

GAUSS 17TM

Language Reference



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Table of Contents

33 Command Reference Introduction	33-1
33.1 Documentation Conventions	33-2
33.2 Command Components	33-3
33.3 Using This Manual	33-5
33.4 Global Control Variables	33-6
33.4.1 Changing the Default Values	33-6
33.4.2 The Procedure <code>gausset</code>	33-7
34 Commands by Category	34-1
34.1 Mathematical Functions	34-4
34.1.1 Scientific Functions	34-4
34.1.2 Differentiation and Integration	34-6
34.1.3 Linear Algebra	34-8
34.1.4 Eigenvalues	34-13
34.1.5 Polynomial Operations	34-13
34.1.6 Fourier Transforms	34-14
34.1.7 Random Numbers	34-14
34.1.8 Fuzzy Conditional Functions	34-17

34.1.9 Statistical Functions	34-19
34.1.10 Optimization and Solution	34-23
34.1.11 Statistical Distributions	34-25
34.1.12 Series and Sequence Functions	34-29
34.1.13 Precision Control	34-30
34.2 Finance Functions	34-30
34.3 Matrix Manipulation	34-32
34.3.1 Creating Vectors and Matrices	34-32
34.3.2 Loading and Storing Matrices	34-33
34.3.3 Size, Ranking, and Range	34-34
34.3.4 Miscellaneous Matrix Manipulation	34-35
34.4 Sparse Matrix Handling	34-38
34.5 N-Dimensional Array Handling	34-40
34.5.1 Creating Arrays	34-40
34.5.2 Size, Ranking and Range	34-40
34.5.3 Setting and Retrieving Data in an Array	34-41
34.5.4 Miscellaneous Array Functions	34-42
34.6 Structures	34-42
34.7 Data File Reading/Writing	34-44
34.7.1 Spreadsheets	34-44
34.7.2 CSV Files	34-45

34.7.3 Text Files	34-45
34.7.4 HDF 5 Files	34-46
34.7.5 Database	34-46
34.8 Compiler Control	34-61
34.9 Multi-Threading	34-63
34.10 Program Control	34-64
34.10.1 Execution Control	34-64
34.10.2 Branching	34-64
34.10.3 Looping	34-65
34.10.4 Subroutines	34-66
34.10.5 Procedures, Keywords, and Functions	34-66
34.10.6 Libraries	34-67
34.10.7 Compiling	34-67
34.10.8 Miscellaneous Program Control	34-68
34.11 OS Functions and File Management	34-69
34.12 Workspace Management	34-70
34.13 Error Handling and Debugging	34-70
34.14 String Handling	34-71
34.15 Time and Date Functions	34-75
34.16 Console I/O	34-77
34.17 Output Functions	34-78

34.17.1 Text Output	34-78
34.17.2 DOS Compatibility Windows	34-79
34.18 GAUSS Graphics	34-80
34.18.1 Graph Types	34-80
34.18.2 Adding Data to Existing Graphs	34-81
34.18.3 Plot Control	34-82
34.18.4 Annotation Control	34-84
34.19 PQG Graphics	34-85
34.19.1 Graph Types	34-85
34.19.2 Axes Control and Scaling	34-86
34.19.3 Text, Labels, Titles, and Fonts	34-87
34.19.4 Main Curve Lines and Symbols	34-87
34.19.5 Extra Lines and Symbols	34-88
34.19.6 Graphic Panel, Page, and Plot Control	34-88
34.19.7 Output Options	34-89
34.19.8 Miscellaneous	34-90
35 Command Reference	35-1
a	35-52
abs	35-52
acf	35-53
aconcat	35-58

aeye	35-61
amax	35-63
amean	35-65
AmericanBinomCall	35-67
AmericanBinomCall_Greeks	35-69
AmericanBinomCall_ImpVol	35-71
AmericanBinomPut	35-73
AmericanBinomPut_Greeks	35-75
AmericanBinomPut_ImpVol	35-77
AmericanBSCall	35-79
AmericanBSCall_Greeks	35-80
AmericanBSCall_ImpVol	35-82
AmericanBSPut	35-84
AmericanBSPut_Greeks	35-85
AmericanBSPut_ImpVol	35-87
amin	35-89
amult	35-92
annotationGetDefaults	35-94
annotationSetBkd	35-95
annotationSetFont	35-97
annotationSetLineColor	35-99

annotationSetLineStyle	35-101
annotationSetLineThickness	35-103
annualTradingDays	35-104
arccos	35-106
arcsin	35-107
areshape	35-108
arrayalloc	35-110
arrayindex	35-112
arrayinit	35-114
arraytomat	35-115
asciiload	35-116
asclabel	35-118
astd	35-120
astds	35-122
asum	35-124
atan	35-126
atan2	35-128
atranspose	35-130
axmargin	35-132
b	35-135
balance	35-135

band	35-136
bandchol	35-138
bandcholsol	35-140
bandltsol	35-141
bandrv	35-143
bandsolpd	35-145
bar	35-146
base10	35-148
begwind	35-149
besselj	35-150
bessely	35-151
beta	35-153
box	35-154
boxcox	35-156
break	35-157
c	35-159
call	35-159
cdfBeta	35-160
cdfBetaInv	35-162
cdfBinomial	35-164
cdfBinomialInv	35-165

cdfBvn	35-167
cdfBvn2	35-169
cdfBvn2e	35-171
cdfCauchy	35-173
cdfCauchyInv	35-174
cdfChic	35-175
cdfChii	35-177
cdfChinc	35-179
cdfChincInv	35-180
cdfEmpirical	35-182
cdfExp	35-185
cdfExpInv	35-186
cdfFc	35-186
cdfFnc	35-189
cdfFncInv	35-190
cdfGam	35-192
cdfGenPareto	35-194
cdfHyperGeo	35-196
cdfLaplace	35-198
cdfLaplaceInv	35-199
cdfLogistic	35-199

cdfLogisticInv	35-201
cdfMvn	35-201
cdfMvnce	35-205
cdfMvne	35-210
cdfMvn2e	35-215
cdfMvtce	35-221
cdfMvte	35-227
cdfMvt2e	35-233
cdfN, cdfNc	35-239
cdfNegBinomial	35-243
cdfNegBinomialInv	35-245
cdfN2	35-246
cdfNi	35-248
cdfPoisson	35-250
cdfPoissonInv	35-251
cdfRayleigh	35-252
cdfRayleighInv	35-254
cdfTc	35-255
cdfTci	35-257
cdfTnc	35-258
cdfTvn	35-261

cdfWeibull	35-263
cdfWeibullInv	35-265
cdir	35-266
ceil	35-267
changeDir	35-269
chdir	35-269
chiBarSquare	35-270
chol	35-272
choldn	35-274
cholsol	35-276
cholup	35-278
chrs	35-279
clear	35-281
clearg	35-282
close	35-283
closeall	35-286
cls	35-289
code	35-290
code (dataloop)	35-293
cols	35-295
colsf	35-296

combinate	35-297
combined	35-299
comlog	35-302
compile	35-303
complex	35-305
con	35-306
cond	35-309
conj	35-310
cons	35-311
ConScore	35-312
continue	35-316
contour	35-318
conv	35-319
convertsatostr	35-320
convertstrtosa	35-321
corrm,corrvc,corrxx	35-322
corrms,corrxxs	35-324
cos	35-327
cosh	35-328
counts	35-329
countwts	35-331

create	35-333
crossprd	35-340
crout	35-341
croutp	35-343
csrcol,csrlin	35-345
csvReadM	35-346
csvReadSA	35-352
csvWriteM	35-356
cumprodc	35-360
cumsumc	35-362
curve	35-363
cvtos	35-365
cvtosa	35-366
d	35-368
datacreate	35-368
datacreatecomplex	35-370
datalist	35-373
dataload	35-374
dataloop (dataloop)	35-376
dataopen	35-377
datasave	35-381

date	35-382
datestr	35-383
datestring	35-384
datestrymd	35-385
dayinyr	35-386
dayofweek	35-387
dbAddDatabase	35-389
dbClose	35-391
dbCommit	35-391
dbCreateQuery	35-392
dbExecQuery	35-394
dbGetConnectOptions	35-395
dbGetDatabaseName	35-396
dbGetDriverName	35-396
dbGetDrivers();	35-397
dbGetHostName	35-398
dbGetLastErrorNum	35-399
dbGetLastErrorText	35-399
dbGetNumericalPrecPolicy	35-400
dbGetPassword	35-401
dbGetPort	35-401

dbGetPrimaryIndex	35-402
dbGetTableHeaders	35-403
dbGetTables	35-404
dbGetUserName	35-404
dbHasFeature	35-405
dbIsDriverAvailable	35-407
dbIsOpen	35-408
dbIsOpenError	35-408
dbIsValid	35-409
dbNumericalPrecPolicy	35-410
dbOpen	35-410
dbQueryBindValue	35-412
dbQueryClear	35-413
dbQueryCols	35-413
dbQueryExecPrepared	35-414
dbQueryFetchAllM	35-415
dbQueryFetchAllSA	35-417
dbQueryFetchOneM	35-418
dbQueryFetchOneSA	35-420
dbQueryFinish	35-421
dbQueryGetBoundValue	35-422

dbQueryGetBoundValues	35-423
dbQueryGetField	35-424
dbQueryGetLastErrorNum	35-426
dbQueryGetLastErrorText	35-426
dbQueryGetLastInsertID	35-427
dbQueryGetLastQuery	35-428
dbQueryGetNumRowsAffected	35-429
dbQueryGetPosition	35-430
dbQueryIsActive	35-431
dbQueryIsForwardOnly	35-432
dbQueryIsNull	35-433
dbQueryIsSelect	35-434
dbQueryIsValid	35-435
dbQueryPrepare	35-436
dbQueryRows	35-438
dbQuerySeek	35-439
dbQuerySeekFirst	35-440
dbQuerySeekLast	35-442
dbQuerySeekNext	35-443
dbQuerySeekPrevious	35-445
dbQuerySetForwardOnly	35-446

dbQuerySetNumericalPrecisionPolicy	35-447
dbRemoveDatabase	35-448
dbRollback	35-448
dbSetConnectOptions	35-449
dbSetDatabaseName	35-452
dbSetHostName	35-453
dbSetPassword	35-454
dbSetPort	35-455
dbSetUserName	35-456
dbTransaction	35-456
debug	35-458
declare	35-458
delete	35-464
delete (dataloop)	35-466
deleteFile	35-467
delif	35-468
delrows	35-473
denseToSp	35-475
denseToSpRE	35-476
denToZero	35-478
design	35-480

det	35-482
detl	35-484
dfft	35-485
dffti	35-486
diag	35-487
diagrv	35-490
digamma	35-492
dlibrary	35-492
dllexport	35-495
do while,do until	35-497
dos	35-500
doswin	35-503
DOSWinCloseall	35-503
DOSWinOpen	35-504
dot	35-507
dotfeq,dotfge, dotfgt,dotfle,dotflt,dotfne	35-509
dotfeqmt,dotfgemt,dotfgtmt,dotflemt,dotflmt,dotfnemt	35-511
draw	35-513
drop (dataloop)	35-515
dsCreate	35-515
dstat	35-516

dstatmt	35-523
dstatmtControlCreate	35-531
dtdate	35-532
dtday	35-534
dttime	35-535
dttodtv	35-536
dttostr	35-537
dttoutc	35-540
dtvnormal	35-542
dtvtodt	35-543
dtvtoutc	35-545
dummy	35-546
dummybr	35-548
dummydn	35-550
e	35-553
ed	35-553
edit	35-554
erfInv,erfCInv	35-555
eig	35-556
eigh	35-558
eighv	35-560

eigv	35-561
elapsedTradingDays	35-563
end	35-565
endp	35-566
endwind	35-567
envget	35-568
eof	35-569
eqSolve	35-571
eqSolveMt	35-576
eqSolveMtControlCreate	35-584
eqSolveMtOutCreate	35-585
eqSolveSet	35-587
erf,erfc	35-587
erfcplx,erfcplx	35-589
error	35-590
errorlog	35-593
errorlogat	35-593
etdays	35-594
ethsec	35-595
etstr	35-597
EuropeanBinomCall	35-598

EuropeanBinomCall_Greeks	35-600
EuropeanBinomCall_ImpVol	35-602
EuropeanBinomPut	35-604
EuropeanBinomPut_Greeks	35-605
EuropeanBinomPut_ImpVol	35-608
EuropeanBSCall	35-609
EuropeanBSCall_Greeks	35-611
EuropeanBSCall_ImpVol	35-613
EuropeanBSPut	35-614
EuropeanBSPut_Greeks	35-616
EuropeanBSPut_ImpVol	35-618
exctsmpl	35-619
exec	35-621
execbg	35-622
exp	35-623
extern (dataloop)	35-625
external	35-626
eye	35-627
f	35-629
fcheckerr	35-629
fclearerr	35-630

feq, fge, fgt, fle, flt, fne	35-631
feqmt, fgemt, fgtmt, flemt, fltmt, fnemt	35-633
fflush	35-635
fft	35-635
ffti	35-636
fftm	35-637
fftn	35-640
fftn	35-643
fgets	35-645
fgetsa	35-646
fgetsat	35-647
fgetst	35-648
fileinfo	35-649
filesa	35-651
floor	35-652
fmod	35-654
fn	35-656
fonts	35-657
fopen	35-658
for	35-660
format	35-663

formatcv	35-673
formatnv	35-674
fputs	35-675
fputst	35-678
fseek	35-681
fstrerror	35-683
ftell	35-684
ftocv	35-685
ftos	35-686
ftostrC	35-690
g	35-693
gamma	35-693
gammacplx	35-694
gammaii	35-695
gausset	35-696
gdaAppend	35-697
gdaCreate	35-699
gdaDStat	35-700
gdaDStatMat	35-704
gdaGetIndex	35-709
gdaGetName	35-710

gdaGetNames	35-711
gdaGetOrders	35-712
gdaGetType	35-714
gdaGetTypes	35-715
gdaGetVarInfo	35-717
gdaIsCplx	35-719
gdaLoad	35-720
gdaPack	35-724
gdaRead	35-726
gdaReadByIndex	35-727
gdaReadSome	35-728
gdaReadSparse	35-731
gdaReadStruct	35-732
gdaReportVarInfo	35-734
gdaSave	35-735
gdaUpdate	35-738
gdaUpdateAndPack	35-740
gdaVars	35-742
gdaWrite	35-743
gdaWrite32	35-744
gdaWriteSome	35-746

getarray	35-749
getdims	35-750
getf	35-751
getGAUSSHome	35-754
getmatrix	35-755
getmatrix4D	35-757
getname	35-759
getnamef	35-760
getNextTradingDay	35-762
getNextWeekDay	35-763
getnr	35-764
getnrmt	35-765
getorders	35-766
getpath	35-767
getPreviousTradingDay	35-769
getPreviousWeekDay	35-770
getRow	35-770
getscalar3D	35-772
getscalar4D	35-773
getTrRow	35-775
getwind	35-776

glm	35-777
gosub	35-807
goto	35-810
gradMT	35-811
gradMTm	35-813
gradMTT	35-815
gradMTTm	35-817
gradp, gradcplx	35-818
graphprt	35-820
graphset	35-824
h	35-825
h5create	35-825
h5open	35-828
h5read	35-831
h5readAttribute	35-838
h5write	35-840
h5writeAttribute	35-842
hasimag	35-845
header	35-846
headermt	35-847
hess	35-849

hessMT	35-850
hessMTg	35-852
hessMTgw	35-854
hessMTm	35-855
hessMTmw	35-857
hessMTT	35-859
hessMTTg	35-861
hessMTTgw	35-863
hessMTTm	35-864
hessMTw	35-866
hessp, hesscplx	35-868
hist	35-870
histf	35-872
histp	35-873
hsec	35-874
i	35-876
if, else, elseif	35-876
imag	35-877
#include	35-878
indcv	35-880
indexcat	35-882

indices	35-883
indices2	35-885
indicesf	35-886
indicesfn	35-888
indnv	35-889
indsav	35-891
integrate1d	35-892
intgrat2	35-897
intgrat3	35-899
inthp1	35-902
inthp2	35-906
inthp3	35-910
inthp4	35-914
inthpControlCreate	35-918
intquad1	35-919
intquad2	35-922
intquad3	35-926
intrleav	35-930
intrleavsa	35-932
intrsect	35-933
intrsectsa	35-936

intsimp	35-938
inv, invpd	35-939
invswp	35-941
iscplx	35-942
iscplx f	35-943
isden	35-944
isinfnanmiss	35-945
ismiss	35-946
itos	35-948
k	35-950
keep (dataloop)	35-950
key	35-951
keyav	35-952
keyw	35-953
keyword	35-953
l	35-956
lag (dataloop)	35-956
lag1	35-957
lagn	35-958
lapeighb	35-960
lapeighi	35-962

lapeighvb	35-964
lapeighvi	35-966
lapgeig	35-968
lapgeigh	35-969
lapgeighv	35-971
lapgeigv	35-973
lapgsvdest	35-974
lapgsvds	35-977
lapgsvdst	35-979
lapgschur	35-982
lapsvdcusv	35-987
lapsvds	35-989
lapsvdusv	35-990
ldl	35-992
ldlp	35-994
ldlsol	35-996
let	35-997
lib	35-1001
library	35-1005
#lineson, #linesoff	35-1009
linsolve	35-1010

listwise (dataloop)	35-1012
ln	35-1012
lncdfbvn	35-1013
lncdfbvn2	35-1014
lncdfmvn	35-1016
lncdfn	35-1017
lncdfn2	35-1018
lncdfnc	35-1020
lnfact	35-1021
lngammacplx	35-1023
lnpdfmvn	35-1024
lnpdfmvt	35-1025
lnpdfn	35-1026
lnpdft	35-1027
load, loadf, loadk, loadm, loadp, loads	35-1028
loadarray	35-1033
loadd	35-1035
loadstruct	35-1039
loadwind	35-1040
local	35-1041
locate	35-1041

loess	35-1042
loessmt	35-1044
loessmtControlCreate	35-1046
log	35-1047
loglog	35-1048
logx	35-1049
logy	35-1050
loopnextindex	35-1051
lower	35-1054
lowmat, lowmat1	35-1056
ltrisol	35-1057
lu	35-1058
lusol	35-1060
m	35-1061
machEpsilon	35-1061
make (dataloop)	35-1061
makevars	35-1062
makewind	35-1064
margin	35-1066
matalloc	35-1067
matinit	35-1068

mattoarray	35-1069
maxc	35-1070
maxindc	35-1071
maxv	35-1073
maxvec	35-1075
maxbytes	35-1076
mbesseli	35-1077
meanc	35-1080
median	35-1081
mergeby	35-1083
mergevar	35-1084
minc	35-1086
minindc	35-1087
minv	35-1089
miss, missrv	35-1090
missex	35-1093
moment	35-1095
momentd	35-1097
movingave	35-1101
movingaveExpwgt	35-1102
movingaveWgt	35-1103

msym	35-1104
n	35-1107
new	35-1107
nextindex	35-1108
nextn, nextnevn	35-1110
nextwind	35-1112
ntos	35-1112
null	35-1115
null1	35-1117
numCombinations	35-1118
o	35-1120
ols	35-1120
olsmt	35-1129
olsmtControlCreate	35-1140
olsqr	35-1141
olsqr2	35-1142
olsqrmt	35-1143
ones	35-1145
open	35-1146
optn, optnevn	35-1152
orth	35-1154

output	35-1156
outtyp (dataloop)	35-1159
outwidth	35-1160
p	35-1162
pacf	35-1162
packedToSp	35-1171
packr	35-1173
parse	35-1175
pause	35-1179
pdfBinomial	35-1179
pdfCauchy	35-1181
pdfexp	35-1182
pdfGenPareto	35-1183
pdfHyperGeo	35-1185
pdfLaplace	35-1187
pdflogistic	35-1188
pdfn	35-1189
pdfPoisson	35-1190
pdfRayleigh	35-1192
pdfWeibull	35-1193
pdfWishartInv	35-1194

pi	35-1196
pinv	35-1197
pinvmt	35-1199
plotAddArea	35-1200
plotAddArrow	35-1204
plotAddBar	35-1206
plotAddBox	35-1207
plotAddErrorBar	35-1208
plotAddHist	35-1216
plotAddHistF	35-1217
plotAddHistP	35-1218
plotAddPolar	35-1219
plotAddScatter	35-1220
plotAddShape	35-1221
plotAddSurface	35-1223
plotAddTextbox	35-1228
plotAddTS	35-1231
plotAddXY	35-1232
plotArea	35-1233
plotBar	35-1234
plotBox	35-1236

plotCDFEmpirical	35-1238
plotClearLayout	35-1241
plotContour	35-1243
plotCustomLayout	35-1246
plotGetDefaults	35-1247
plotHist	35-1249
plotHistF	35-1250
plotHistP	35-1250
plotLayout	35-1251
plotLogLog	35-1253
plotLogX	35-1253
plotLogY	35-1254
plotOpenWindow	35-1255
plotPolar	35-1256
plotSave	35-1257
plotScatter	35-1260
plotSetAxesPen	35-1261
plotSetBar	35-1262
plotSetBkdColor	35-1265
plotSetColorMap	35-1267
plotSetContourLabels	35-1270

plotSetFill	35-1273
plotSetGrid	35-1275
plotSetLegend	35-1276
plotSetLegendFont	35-1278
plotSetLineColor	35-1280
plotSetLineStyle	35-1281
plotSetLineSymbol	35-1282
plotSetLineThickness	35-1285
plotSetNewWindow	35-1286
plotSetTextInterpreter	35-1287
plotSetTitle	35-1292
plotSetWhichY Axis	35-1295
plotSetXLabel	35-1296
plotSetXRange	35-1299
plotSetXTicCount	35-1300
plotSetXTicInterval	35-1303
plotSetXTicLabel	35-1306
plotSetYLabel	35-1311
plotSetYRange	35-1314
plotSetYTicCount	35-1315
plotSetZLabel	35-1319

plotSetZLevels	35-1321
plotSurface	35-1325
plotTS	35-1327
plotXY	35-1331
polar	35-1332
polychar	35-1333
polyeval	35-1334
polygamma	35-1336
polyint	35-1339
polymake	35-1340
polymat	35-1342
polymroot	35-1343
polymult	35-1345
polyroot	35-1347
pop	35-1348
powerM	35-1349
pqgwin	35-1351
previousindex	35-1352
princomp	35-1353
print	35-1355
printdos	35-1366

printfm	35-1367
printfmt	35-1370
proc	35-1373
prodc	35-1375
psi	35-1376
putarray	35-1377
putf	35-1379
putvals	35-1380
pvCreate	35-1382
pvGetIndex	35-1383
pvGetParNames	35-1383
pvGetParVector	35-1385
pvLength	35-1387
pvList	35-1388
pvPack	35-1389
pvPacki	35-1390
pvPackm	35-1392
pvPackmi	35-1394
pvPacks	35-1396
pvPacksi	35-1398
pvPacksm	35-1400

pvPacksmi	35-1402
pvPutParVector	35-1405
pvTest	35-1407
pvUnpack	35-1407
q	35-1409
QNewton	35-1409
QNewtonmt	35-1412
QNewtonmtControlCreate	35-1418
QNewtonmtOutCreate	35-1419
QNewtonSet	35-1420
QProg	35-1420
QProgmt	35-1422
QProgmtInCreate	35-1425
qqr	35-1426
qqre	35-1428
qqrep	35-1431
qr	35-1433
qre	35-1435
qrep	35-1438
qrsol	35-1440
qrtsol	35-1441

qtyr	35-1443
qtyre	35-1446
qtyrep	35-1449
quantile	35-1451
quantiled	35-1454
qyr	35-1458
qyre	35-1460
qyrep	35-1462
qz	35-1464
r	35-1470
rank	35-1470
rankindx	35-1471
readr	35-1472
real	35-1474
reclassify	35-1475
reclassifyCuts	35-1478
recode	35-1483
recode (dataloop)	35-1486
recserar	35-1488
recsercp	35-1490
recsercc	35-1492

renamefile	35-1494
rerun	35-1495
rescale	35-1495
reshape	35-1501
retp	35-1503
return	35-1503
rev	35-1504
rfft	35-1506
rffti	35-1507
rfftip	35-1507
rfftn	35-1509
rfftnp	35-1511
rfftp	35-1513
rndBernoulli	35-1514
rndBeta	35-1516
rndCauchy	35-1519
rndChiSquare	35-1520
rndcon, rndmult, rndseed	35-1522
rndCreateState	35-1524
rndExp	35-1527
rndgam	35-1529

rndGamma	35-1530
rndGeo	35-1532
rndGumbel	35-1534
rndHyperGeo	35-1535
rndi	35-1538
rndKMbeta	35-1541
rndKMgam	35-1543
rndKMi	35-1544
rndKMn	35-1546
rndKMnb	35-1548
rndKMp	35-1550
rndKMu	35-1551
rndKMvm	35-1553
rndLaplace	35-1555
rndLCbeta	35-1557
rndLCgam	35-1559
rndLCi	35-1560
rndLCn	35-1563
rndLCnb	35-1565
rndLCp	35-1567
rndLCu	35-1569

rndLCvm	35-1572
rndLogNorm	35-1574
rndMVn	35-1575
rndMVt	35-1577
rndn	35-1579
rndnb	35-1581
rndNegBinomial	35-1583
rndp	35-1585
rndPoisson	35-1587
rndStateSkip	35-1588
rndu	35-1590
rndvm	35-1593
rndWeibull	35-1594
rndWishart	35-1595
rndWishartInv	35-1597
rotater	35-1598
round	35-1600
rows	35-1601
rowsf	35-1602
rref	35-1603
run	35-1605

s	35-1608
sampleData	35-1608
satostrC	35-1610
save	35-1613
saveall	35-1616
saved	35-1617
savestruct	35-1619
savewind	35-1620
scale	35-1621
scale3d	35-1622
scalerr	35-1623
scalinfnanmiss	35-1626
scalmiss	35-1627
schtoc	35-1628
schur	35-1630
screen	35-1633
searchsourcepath	35-1634
seekr	35-1635
select (dataloop)	35-1636
selif	35-1637
seqa, seqm	35-1639

setarray	35-1640
setdif	35-1642
setdifsa	35-1645
setvars	35-1646
setvwrmode	35-1647
setwind	35-1648
shell	35-1649
shiftr	35-1651
show	35-1653
sin	35-1656
singleindex	35-1657
sinh	35-1659
sleep	35-1660
solpd	35-1661
sortc, sortcc	35-1664
sortd	35-1666
sorthc, sorthcc	35-1667
sortind, sortindc	35-1669
sortmc	35-1670
sortr, sortrc	35-1672
spBiconjGradSol	35-1674

spChol	35-1677
spConjGradSol	35-1678
spCreate	35-1681
spDenseSubmat	35-1682
spDiagRvMat	35-1684
spEigv	35-1687
spEye	35-1690
spGetNZE	35-1691
spline	35-1693
spLDL	35-1694
spLU	35-1695
spNumNZE	35-1697
spOnes	35-1698
spreadSheetReadM	35-1700
spreadSheetReadSA	35-1702
spreadSheetWrite	35-1704
spScale	35-1707
spSubmat	35-1709
spToDense	35-1711
spTrTDense	35-1712
spTScalar	35-1713

spZeros	35-1714
sqpSolve	35-1716
sqpSolveMT	35-1723
sqpSolveMTControlCreate	35-1731
sqpSolveMTlagrangeCreate	35-1732
sqpSolveMToutCreate	35-1733
sqpSolveSet	35-1734
sqrt	35-1735
stdc	35-1736
stdsc	35-1738
stocv	35-1739
stof	35-1740
stop	35-1741
strcombine	35-1742
strindx	35-1744
strlen	35-1745
strput	35-1746
strrindx	35-1748
strjoin	35-1749
strsect	35-1751
strsplit	35-1753

strsplitPad	35-1757
strtodt	35-1758
strtof	35-1760
strtofcp1x	35-1762
strtrim	35-1763
strtriml	35-1765
strtrimr	35-1766
strtrunc	35-1767
strtrunc1	35-1768
strtruncpad	35-1769
strtruncr	35-1770
submat	35-1770
subscat	35-1772
substute	35-1776
subvec	35-1779
sumc	35-1781
sumr	35-1783
surface	35-1785
svd	35-1788
svd1	35-1790
svd2	35-1792

svdcusv	35-1795
svds	35-1798
svdusv	35-1801
sylvester	35-1804
sysstate	35-1806
system	35-1828
t	35-1830
tab	35-1830
tan	35-1831
tanh	35-1832
tempname	35-1833
ThreadBegin	35-1834
ThreadEnd	35-1835
threadfor, threadendfor	35-1836
ThreadJoin	35-1839
ThreadStat	35-1840
time	35-1841
timedt	35-1841
timestr	35-1842
timeutc	35-1843
title	35-1844

tkf2eps	35-1846
tkf2ps	35-1847
tocart	35-1848
todaydt	35-1848
toeplitz	35-1849
token	35-1850
topolar	35-1853
trace	35-1853
trap	35-1855
trapchk	35-1859
trigamma	35-1861
trimr	35-1862
trunc	35-1863
type	35-1865
typecv	35-1867
typef	35-1869
u	35-1871
union	35-1871
unionsa	35-1872
uniqindx	35-1873
uniqindxsa	35-1875

unique	35-1876
uniquesa	35-1882
upmat, upmat1	35-1883
upper	35-1885
use	35-1886
utctodt	35-1888
utctodtv	35-1889
utrisol	35-1891
v	35-1892
vals	35-1892
varCovM, varCovX	35-1894
varCovMS, varCovXS	35-1896
varget	35-1898
vargetl	35-1900
varmall	35-1902
varmares	35-1903
varput	35-1905
varputl	35-1906
vartypef	35-1908
vcm, vcx	35-1909
vcms, vcxs	35-1910

vec, vecr	35-1911
vech	35-1913
vector (dataloop)	35-1914
vget	35-1915
view	35-1916
viewxyz	35-1917
vlist	35-1918
vnamecv	35-1919
volume	35-1920
vput	35-1921
vread	35-1922
vtypecv	35-1923
w	35-1924
wait, waitc	35-1924
walkindex	35-1924
window	35-1926
writer	35-1927
x	35-1930
xlabel	35-1930
xlsGetSheetCount	35-1931
xlsGetSheetSize	35-1932

xlsGetSheetTypes	35-1935
xlsMakeRange	35-1937
xlsReadM	35-1939
xlsReadSA	35-1944
xlsWrite	35-1948
xlsWriteM	35-1952
xlsWriteSA	35-1955
xpnd	35-1958
xtics	35-1960
xy	35-1961
xyz	35-1962
y	35-1964
ylabel	35-1964
ytics	35-1965
z	35-1967
zeros	35-1967
zeta	35-1968
zlabel	35-1969
ztics	35-1970
36 Miscellaneous Topics	36-1
36.1 Change Log	36-1

36.2 Common Errors	36-8
36.3 Error Messages	36-13
36.4 GAUSS Graphics Colors	36-39
36.5 Obsolete Commands	36-42
36.6 Quick Reference	36-46
36.6.1 Basic functions	36-46
36.6.2 Matrix creation	36-47
36.6.3 Matrix manipulation	36-49
36.6.4 Operators	36-51
36.6.5 String creation	36-54
36.6.6 String array manipulation	36-54
36.6.7 Loading and saving data	36-55
36.7 Reserved Words	36-57

33 Command Reference Introduction

The GAUSS LANGUAGE REFERENCE describes each of the commands, procedures and functions available in the **GAUSS**TM programming language. These functions can be divided into four categories:

- Mathematical, statistical and scientific functions.
- Data handling routines, including data matrix manipulation and description routines, and file I/O.
- Programming statements, including branching, looping, display features, error checking, and shell commands.
- Graphics functions.

The first category contains those functions to be expected in a high level mathematical language: trigonometric functions and other transcendental functions, distribution functions, random number generators, numerical differentiation and integration routines, Fourier transforms, Bessel functions and polynomial evaluation routines. And, as a matrix programming language, **GAUSS** includes a variety of routines that perform standard matrix operations. Among these are routines to calculate determinants, matrix inverses, decompositions, eigenvalues and eigenvectors, and condition numbers.

Data handling routines include functions which return dimensions of matrices, and information about elements of data matrices, including functions to locate values lying in specific ranges or with certain values. Also under data handling routines fall all those functions that create, save, open and read from and write to **GAUSS** data sets and

Command Reference Introduction

GAUSS Data Archives. A variety of sorting routines which will operate on both numeric and character data are also available.

Programming statements are all of the commands that make it possible to write complex programs in **GAUSS**. These include conditional and unconditional branching, looping, file I/O, error handling, and system-related commands to execute OS shells and access directory and environment information.

The graphics functions of **GAUSS Publication Quality Graphics** (PQG) are a set of routines built on the graphics functions in GraphiC by Scientific Endeavors Corporation. **GAUSS** PQG consists of a set of main graphing procedures and several additional procedures and global variables for customizing the output.

33.1 Documentation Conventions	33-2
33.2 Command Components	33-3
33.3 Using This Manual	33-5
33.4 Global Control Variables	33-6
33.4.1 Changing the Default Values	33-6
33.4.2 The Procedure gausset	33-7

33.1 Documentation Conventions

The following table describes how text formatting is used to identify **GAUSS** programming elements:

Text Style	Use	Example
regular text	narrative	"... text formatting is used ..."
bold text	emphasis	"...not supported under UNIX."

<i>italics</i>	variables	"... If <i>vnames</i> is a string or has fewer elements than <i>x</i> has columns, it will be ..."
monospace	code example	<code>if scalerr (cm) ; cm = inv (x) ; endif;</code>
monospace	filename, path, etc.	"...is located in the examples subdirectory..."
monospace bold	reference to a GAUSS command or other programming element within a narrative paragraph	"...as explained under plotScatter... "
Bold Text	reference to section of the manual	"...see Operator Precedence , Section 1.1 ..."

33.2 Command Components

The following list describes each of the components used in the **COMMAND REFERENCE**, CHAPTER 35 .

Purpose

Describes what the command or function does.

Library

Lists the library that needs to be activated to access the function.

Command Reference Introduction

Include

Lists files that need to be included to use the function.

Format

Illustrates the syntax of the command or function.

Input

Describes the input parameters of the function.

Global Input

Describes the global variables that are referenced by the function.

Output

Describes the return values of the function.

Global Output

Describes the global variables that are updated by the function.

Portability

Describes differences under various operating systems.

Remarks

Explanatory material pertinent to the command.

Example

Sample code using the command or function.

Source

The source file in which the function is defined, if applicable.

Globals

Global variables that are accessed by the command.

See Also

Other related commands.

Technical Notes

Technical discussion and reference source citations.

References

Reference material citations.

33.3 Using This Manual

Users who are new to **GAUSS** should make sure they have familiarized themselves with **LANGUAGE FUNDAMENTALS**, Chapter [9](#), before proceeding here. That chapter contains the basics of **GAUSS** programming.

In all, there are over 800 routines described in this **GAUSS LANGUAGE REFERENCE**. We suggest that new **GAUSS** users skim through **COMMANDS BY CATEGORY**, CHAPTER 34 , and then browse through **COMMAND REFERENCE**, CHAPTER 35 , the main part of this manual. Here, users can familiarize themselves with the kinds of tasks that **GAUSS** can handle easily.

Chapter 34 gives a categorical listing of all functions in this **GAUSS LANGUAGE REFERENCE** and a short discussion of the functions in each category. Complete syntax, description of input and output arguments, and general remarks regarding each function are given in Chapter 35 .

Command Reference Introduction

If a function is an "extrinsic" (that is, part of the **Run-Time Library**), its source code can be found on the `src` subdirectory. The name of the file containing the source code is given in Chapter 35 under the discussion of that function.

33.4 Global Control Variables

Several **GAUSS** functions use global variables to control various aspects of their performance. The files `gauss.ext`, `gauss.dec` and `gauss.lcg` contain the `external` statements, `declare` statements, and library references to these globals. All globals used by the **GAUSS Run-Time Library** begin with an underscore `'_'`.

Default values for these common globals can be found in the file `gauss.dec`, located on the `src` subdirectory. The default values can be changed by editing this file.

33.4.1 Changing the Default Values33-6

33.4.2 The Procedure `gausset`33-7

33.4.1 Changing the Default Values

To permanently change the default setting of a common global, two files need to be edited: `gauss.dec` and `gauss.src`.

To change the value of the common global `__output` from 1 to 0, for example, edit the file `gauss.dec` and change the statement

```
declare matrix __output = 1;
```

so it reads:

```
declare matrix __output = 0;
```

Also, edit the procedure **`gausset`**, located in the file `gauss.src`, and modify the statement

```
__output = 1;
```

similarly.

33.4.2 The Procedure **gausset**

The global variables affect your program, even if you have not set them directly in a particular command file. If you have changed them in a previous run, they will retain their changed values until you exit **GAUSS** or execute the [new](#) command.

The procedure **gausset** will reset the **Run-Time Library** globals to their default values.

```
gausset;
```

If your program changes the values of these globals, you can use **gausset** to reset them whenever necessary. **gausset** resets the globals as a whole; you can write your own routine to reset specific ones.

34 Commands by Category

34.1 Mathematical Functions	34-4
34.1.1 Scientific Functions	34-4
34.1.2 Differentiation and Integration	34-6
34.1.3 Linear Algebra	34-8
34.1.4 Eigenvalues	34-13
34.1.5 Polynomial Operations	34-13
34.1.6 Fourier Transforms	34-14
34.1.7 Random Numbers	34-14
34.1.8 Fuzzy Conditional Functions	34-17
34.1.9 Statistical Functions	34-19
34.1.10 Optimization and Solution	34-23
34.1.11 Statistical Distributions	34-25
34.1.12 Series and Sequence Functions	34-29
34.1.13 Precision Control	34-30

34.2 Finance Functions	34-30
34.3 Matrix Manipulation	34-32
34.3.1 Creating Vectors and Matrices	34-32
34.3.2 Loading and Storing Matrices	34-33
34.3.3 Size, Ranking, and Range	34-34
34.3.4 Miscellaneous Matrix Manipulation	34-35
34.4 Sparse Matrix Handling	34-38
34.5 N-Dimensional Array Handling	34-40
34.5.1 Creating Arrays	34-40
34.5.2 Size, Ranking and Range	34-40
34.5.3 Setting and Retrieving Data in an Array	34-41
34.5.4 Miscellaneous Array Functions	34-42
34.6 Structures	34-42
34.7 Data File Reading/Writing	34-44
34.7.1 Spreadsheets	34-44
34.7.2 CSV Files	34-45
34.7.3 Text Files	34-45
34.7.4 HDF 5 Files	34-46
34.7.5 Database	34-46
34.8 Compiler Control	34-61
34.9 Multi-Threading	34-63

34.10 Program Control	34-64
34.10.1 Execution Control	34-64
34.10.2 Branching	34-64
34.10.3 Looping	34-65
34.10.4 Subroutines	34-66
34.10.5 Procedures, Keywords, and Functions	34-66
34.10.6 Libraries	34-67
34.10.7 Compiling	34-67
34.10.8 Miscellaneous Program Control	34-68
34.11 OS Functions and File Management	34-69
34.12 Workspace Management	34-70
34.13 Error Handling and Debugging	34-70
34.14 String Handling	34-71
34.15 Time and Date Functions	34-75
34.16 Console I/O	34-77
34.17 Output Functions	34-78
34.17.1 Text Output	34-78
34.17.2 DOS Compatibility Windows	34-79
34.18 GAUSS Graphics	34-80
34.18.1 Graph Types	34-80
34.18.2 Adding Data to Existing Graphs	34-81

34.18.3 Plot Control	34-82
34.18.4 Annotation Control	34-84
34.19 PQG Graphics	34-85
34.19.1 Graph Types	34-85
34.19.2 Axes Control and Scaling	34-86
34.19.3 Text, Labels, Titles, and Fonts	34-87
34.19.4 Main Curve Lines and Symbols	34-87
34.19.5 Extra Lines and Symbols	34-88
34.19.6 Graphic Panel, Page, and Plot Control	34-88
34.19.7 Output Options	34-89
34.19.8 Miscellaneous	34-90

34.1 Mathematical Functions

34.1.1 Scientific Functions

abs	Returns absolute value of argument.
arccos	Computes inverse cosine.
arcsin	Computes inverse sine.
atan	Computes inverse tangent.
atan2	Computes angle given a point x,y .
besselj	Computes Bessel function, first kind.
bessely	Computes Bessel function, second kind.
beta	Computes the complete Beta function, also called the Euler integral.

boxcox	Computes the Box-Cox function.
cos	Computes cosine.
cosh	Computes hyperbolic cosine.
curve	Computes a one-dimensional smoothing curve.
digamma	Computes the digamma function.
exp	Computes the exponential function of x .
fmod	Computes the floating-point remainder of x/y .
gamma	Computes gamma function value.
gammacplx	Computes gamma function for complex inputs.
gammai	Compute the inverse incomplete gamma function.
ln	Computes the natural log of each element.
lnfact	Computes natural log of factorial function.
lngammacplx	Computes the natural log of the gamma function for complex inputs.
log	Computes the log of each element.
mbesseli	Computes modified and exponentially scaled modified Bessels of the first kind of the n th order.
nextn, nextnevn	Returns allowable matrix dimensions for computing FFT's.
optn, optnevn	Returns optimal matrix dimensions for computing FFT's.
pi	Returns π .
polar	Graphs data using polar coordinates.
polygamma	Computes the polygamma function of order n .
psi	Computes the psi (or digamma) function.
sin	Computes sine.
sinh	Computes the hyperbolic sine.
spline	Computes a two-dimensional interpolatory spline.

sqrt	Computes the square root of each element.
tan	Computes tangent.
tanh	Computes hyperbolic tangent.
tocart	Converts from polar to Cartesian coordinates.
topolar	Converts from Cartesian to polar coordinates.
trigamma	Computes trigamma function.
zeta	Computes the Rieman zeta function.

All trigonometric functions take or return values in radian units.

34.1.2 Differentiation and Integration

gradMT	Computes numerical gradient.
gradMTm	Computes numerical gradient with mask.
gradMTT	Computes numerical gradient using available threads.
gradMTTm	Computes numerical gradient with mask using available threads.
gradp, gradcplx	Computes first derivative of a function; gradcplx allows for complex arguments.
hessMT	Computes numerical Hessian.
hessMTg	Computes numerical Hessian using gradient procedure.
hessMTgw	Computes numerical Hessian using gradient procedure with weights.
hessMTm	Computes numerical Hessian with mask.
hessMTmw	Computes numerical Hessian with mask and weights.
hessMTT	Computes numerical Hessian using

	available threads.
hessMTTg	Computes numerical Hessian using gradient procedure with available threads.
hessMTTgw	Computes numerical Hessian using gradient procedure with weights and using available threads.
hessMTTm	Computes numerical Hessian with mask and available threads.
hessMTw	Computes numerical Hessian with weights.
hessp, hesscplx	Computes second derivative of a function; hesscplx allows for complex arguments.
integrate1d	Integrates a user-defined function over a user-defined range, using adaptive quadrature.
intgrat2	Integrates a 2-dimensional function over an user-defined region.
intgrat3	Integrates a 3-dimensional function over an user-defined region.
inthp1	Integrates an user-defined function over an infinite interval.
inthp2	Integrates an user-defined function over the $[a, +\infty)$ interval.
inthp3	Integrates an user-defined function over the $[a, +\infty)$ interval that is oscillatory.
inthp4	Integrates an user-defined function over the $[a, b]$ interval.
inthpControlCreate	Creates default inthpControl structure.
intquad1	Integrates a 1-dimensional function.

intquad2	Integrates a 2-dimensional function over an user-defined rectangular region.
intquad3	Integrates a 3-dimensional function over an user-defined rectangular region.
intsimp	Integrates by Simpson's method.

gradp and **hessp** use a finite difference approximation to compute the first and second derivatives. Use **gradp** to calculate a Jacobian.

intquad1, **intquad2**, and **intquad3** use Gaussian quadrature to calculate the integral of the user-defined function over a rectangular region.

To calculate an integral over a region defined by functions of x and y , use **intgrat2** and **intgrat3**.

In most cases, **integrate1d** will provide the greatest accuracy and fastest performance for 1 dimensional integration. In some cases **intquad1** will be faster, but with a loss of accuracy. **intsimp** is in between the other two with regards to speed and accuracy.

34.1.3 Linear Algebra

balance	Balances a matrix.
band	Extracts bands from a symmetric banded matrix.
bandchol	Computes the Cholesky decomposition of a positive definite banded matrix.
bandcholsol	Solves the system of equations $Ax = b$ for x , given the lower triangle of the Cholesky decomposition of a positive definite banded matrix A .
bandltsol	Solves the system of equations $Ax = b$ for x , where A is a lower triangular banded matrix
bandrv	Creates a symmetric banded matrix, given its compact form.

bandsolpd	Solves the system of equations $Ax = b$ for x , where A is a positive definite banded matrix.
chol	Computes Cholesky decomposition, $X = Y^T Y$.
choldn	Performs Cholesky downdate on an upper triangular matrix.
cholsol	Solves a system of equations given the Cholesky factorization of a matrix.
cholup	Performs Cholesky update on an upper triangular matrix.
cond	Computes condition number of a matrix.
crout	Computes Crout decomposition, $X = LU$ (real matrices only).
croutp	Computes Crout decomposition with row pivoting (real matrices only).
det	Computes determinant of square matrix.
detl	Computes determinant of decomposed matrix.
hess	Computes upper Hessenberg form of a matrix (real matrices only).
inv	Inverts a matrix.
invpd	Inverts a positive definite matrix.
invswp	Computes a generalized sweep inverse.
lapeighb	Computes eigenvalues only of a real symmetric or complex Hermitian matrix selected by bounds.
lapeighi	Computes eigenvalues only of a real symmetric or complex Hermitian matrix selected by index.
lapeighvb	Computes eigenvalues and eigenvectors of a real symmetric or complex Hermitian matrix selected by bounds.
lapeighvi	Computes selected eigenvalues and eigenvectors of a

		Hermitian matrix.
lapgeig		Computes generalized eigenvalues for a pair of real or complex general matrices.
lapgeigh		Computes generalized eigenvalues for a pair of real symmetric or Hermitian matrices.
lapgeighv		Computes generalized eigenvalues and eigenvectors for a pair of real symmetric or Hermitian matrices.
lapgeigv		Computes generalized eigenvalues, left eigenvectors, and right eigenvectors for a pair of real or complex general matrices.
lapgschur		Computes the generalized Schur form of a pair of real or complex general matrices.
lapgsvdct		Computes the generalized singular value decomposition of a pair of real or complex general matrices.
lapgsvds		Computes the generalized singular value decomposition of a pair of real or complex general matrices.
lapgsvdst		Computes the generalized singular value decomposition of a pair of real or complex general matrices.
lapsvdcusv		Computes the singular value decomposition a real or complex rectangular matrix, returns compact u and v .
lapsvds		Computes the singular values of a real or complex rectangular matrix.
lapsvdusv		Computes the singular value decomposition a real or complex rectangular matrix.
ldl		Computes the L and D factors of the LDL factorization of a real symmetric matrix.
ldlp		Computes LDL decomposition with row pivoting of a

	symmetric matrix.
ldlsol	Computes Solves the system of equations $LDL^T x = b$ using a matrix factorized by ldlp .
lu	Computes LU decomposition with row pivoting (real and complex matrices).
lusol	Computes Solves the system of equations $LUx = b$.
null	Computes orthonormal basis for right null space.
null1	Computes orthonormal basis for right null space.
orth	Computes orthonormal basis for column space x .
pinv	Generalized pseudo-inverse: Moore-Penrose.
pinvmt	Generalized pseudo-inverse: Moore-Penrose.
powerM	Computes the power n of a matrix A , as the matrix product of n copies of A .
qqr	QR decomposition: returns Q_I and R .
qqre	QR decomposition: returns Q_I , R and a permutation vector, E .
qqrep	QR decomposition with pivot control: returns Q_I , R , and E .
qr	QR decomposition: returns R .
qre	QR decomposition: returns R and E .
qrep	QR decomposition with pivot control: returns R and E .
qrsol	Solves a system of equations $R'x = b$ given an upper triangular matrix, typically the R matrix from a QR decomposition.
qrtsol	Solves a system of equations $R'x = b$ given an upper triangular matrix, typically the R matrix from a QR decomposition.
qtyr	QR decomposition: returns $Q'Y$ and R .
qtyre	QR decomposition: returns $Q'Y$, R and E .

qtyrep	QR decomposition with pivot control: returns $Q'Y$, R and E .
qyr	QR decomposition: returns QY and R .
qyre	QR decomposition: returns QY , R and E .
qyrep	QR decomposition with pivot control: returns QY , R and E .
qz	Compute the complex QZ, or generalized Schur, form of a pair of real or complex general matrices with an option to sort the eigenvalues.
rank	Computes rank of a matrix.
rref	Computes reduced row echelon form of a matrix.
schtoc	Reduces any 2x2 blocks on the diagonal of the real Schur form of a matrix returned from schur . The transformation matrix is also updated.
schur	Computes real or complex Schur decomposition of a matrix.
solpd	Solves a system of positive definite linear equations.
svd	Computes the singular values of a matrix.
svd1	Computes singular value decomposition, $X = USV'$.
svd2	Computes svd1 with compact U .
svdcusv	Computes the singular value decomposition of a matrix so that: $x = u * s * v'$ (compact u).
svds	Computes the singular values of a matrix.
svdusv	Computes the singular value decomposition of a matrix so that: $x = u * s * v'$.
sylvester	Computes the solution to the Sylvester matrix equation, $AX + XB = C$.

The decomposition routines are **chol** for Cholesky decomposition, **crout** and **croutp** for Crout decomposition, **qqr-qyrep** for QR decomposition, and **svd-svdusv** for singular value decomposition.

null, **null1**, and **orth** calculate orthonormal bases.

inv, **invpd**, **solpd**, **cholsol**, **qrsol** and the "/" operator can all be used to solve linear systems of equations.

rank and **rref** will find the rank and reduced row echelon form of a matrix.

det, **det1** and **cond** will calculate the determinant and condition number of a matrix.

34.1.4 Eigenvalues

eig	Computes eigenvalues of general matrix.
eigh	Computes eigenvalues of complex Hermitian or real symmetric matrix.
eighv	Computes eigenvalues and eigenvectors of complex Hermitian or real symmetric matrix.
eigv	Computes eigenvalues and eigenvectors of general matrix.

There are four eigenvalue-eigenvector routines. Two calculate eigenvalues only, and two calculate eigenvalues and eigenvectors. The three types of matrices handled by these routines are:

General:	eig, eigv
Symmetric or Hermitian:	eigh, eighv

34.1.5 Polynomial Operations

polychar	Computes characteristic polynomial of a square matrix.
polyeval	Evaluates polynomial with given coefficients.
polyint	Calculates Nth order polynomial interpolation given known point pairs.

Commands by Category

polymake	Computes polynomial coefficients from roots.
polymat	Returns sequence powers of a matrix.
polymult	Multiplies two polynomials together.
polyroot	Computes roots of polynomial from coefficients.

See also **recserrc**, **recsercp**, and **conv**.

34.1.6 Fourier Transforms

dfft	Computes discrete 1-D FFT.
dffti	Computes inverse discrete 1-D FFT.
fft	Computes 1- or 2-D FFT.
ffti	Computes inverse 1- or 2-D FFT.
fftm	Computes multi-dimensional FFT.
fftn	Computes multi-dimensional FFT.
fftn	Computes inverse multi-dimensional FFT.
fftn	Computes 1- or 2-D FFT using prime factor algorithm.
rfft	Computes real 1- or 2-D FFT.
rffti	Computes inverse real 1- or 2-D FFT.
rfftip	Computes inverse real 1- or 2-D FFT from packed format FFT.
rfftn	Computes real 1- or 2-D FFT using prime factor algorithm.
rfftnp	Computes real 1- or 2-D FFT using prime factor algorithm, returns packed format FFT.
rfftp	Computes real 1- or 2-D FFT, returns packed format FFT.

34.1.7 Random Numbers

rndBernoulli	Computes random numbers with Bernoulli distribution.
rndBeta	Computes random numbers with beta distribution.

<code>rndCauchy</code>	Computes Cauchy distributed random numbers with a choice of underlying random number generator.
<code>rndChiSquare</code>	Creates pseudo-random numbers with a chi-squared distribution, with an optional non-centrality parameter and a choice of underlying random number generator.
<code>rndCreateState</code>	Creates a new random number stream for a specified generator type from a seed value.
<code>rndExp</code>	Computes exponentially distributed random numbers with a choice of underlying random number generator.
<code>rndGamma</code>	Computes gamma pseudo-random numbers with a choice of underlying random number generator.
<code>rndGeo</code>	Computes geometric pseudo-random numbers with a choice of underlying random number generator.
<code>rndGumbel</code>	Computes Gumbel distributed random numbers with a choice of underlying random number generator.
<code>rndHyperGeo</code>	Computes pseudo-random numbers following a hypergeometric distribution with a choice of underlying random number generator.
<code>rndi</code>	Returns random integers, $0 \leq y < 2^{32}$.
<code>rndKMbeta</code>	Returns uniformly distributed random integers over a user specified range.
<code>rndKMgam</code>	Computes gamma pseudo-random numbers.
<code>rndKMi</code>	Returns random integers, $0 \leq y < 2^{32}$.
<code>rndKMn</code>	Computes standard normal pseudo-random numbers.

rndKMnb	Computes negative binomial pseudo-random numbers.
rndKMp	Computes Poisson pseudo-random numbers.
rndKMu	Computes uniform pseudo-random numbers.
rndKMvm	Computes von Mises pseudo-random numbers.
rndLaplace	Computes Laplacian pseudo-random numbers with the choice of underlying random number generator.
rndLogNorm	Computes lognormal pseudo-random numbers with the choice of underlying random number generator.
rndMVn	Computes multivariate normal random numbers given a covariance matrix.
rndMVt	Computes multivariate Student-t random numbers given a covariance matrix.
rndn	Computes normally distributed pseudo-random numbers with a choice of underlying random number generator.
rndNegBinomial	Computes negative binomial pseudo-random numbers with a choice of underlying random number generator.
rndPoisson	Computes Poisson pseudo-random numbers with a choice of underlying random number generator.
rndseed	Changes seed of the LC random number generator.
rndStateSkip	To advance a state vector by a specified number of values.
rndu	Computes uniform random numbers with a choice of underlying random number generator.

<code>rndWeibull</code>	Computes Weibull pseudo-random numbers with the choice of underlying random number generator.
<code>rndWishart</code>	Computes Wishart pseudo-random matrices with the choice of underlying random number generator.
<code>rndWishartInv</code>	Computes inverse Wishart pseudo-random matrices with the choice of underlying random number generator.

The random number generator can be seeded. Set the seed using `rndseed`. For example:

```
rndseed 44435667;
x = rndu(1,1);
```

34.1.8 Fuzzy Conditional Functions

<code>dot</code>	Returns a scalar dot product of the columns of two matrices.
<code>dotfeq</code>	Fuzzy . ==
<code>dotfeqmt</code>	Fuzzy . ==
<code>dotfge</code>	Fuzzy . >=

dotfgemt	Fuzzy .>
dotfgt	Fuzzy .>
dotfgtmt	Fuzzy .>
dotfle	Fuzzy .<=
dotflemt	Fuzzy .<=
dotflt	Fuzzy .<
dotfltmt	Fuzzy .<
dotfne	Fuzzy ./=
dotfnemt	Fuzzy ./=
feq	Fuzzy ==
feqmt	Fuzzy ==
fge	Fuzzy >=
fgemt	Fuzzy >=
fgt	Fuzzy >
fgtmt	Fuzzy >
file	Fuzzy <=
flemt	Fuzzy <=
flt	Fuzzy <

fltmt	Fuzzy <
fne	Fuzzy /=
fnemt	Fuzzy /=

The **mt** commands use an *fcmtol* argument to control the tolerance used for comparison.

The non-**mt** commands use the global variable `_fcmtol` to control the tolerance used for comparison. By default, this is 1e-15. The default can be changed by editing the file `fcompare.dec`.

34.1.9 Statistical Functions

acf	Computes sample autocorrelations.
astd	Computes the standard deviation of the elements across one dimension of an N-dimensional array.
astds	Computes the 'sample' standard deviation of the elements across one dimension of an N-dimensional array.
chiBarSquare	Computes probability of chi-bar-square statistic.
combine	Computes combinations of <i>n</i> things taken <i>k</i> at a time.
combined	Writes combinations of <i>n</i> things taken <i>k</i> at a time to a GAUSS data set.
conScore	Computes constrained score statistic and its probability.
conv	Computes convolution of two vectors.
corrmm	Computes correlation matrix of a moment matrix.
corrms	Computes sample correlation matrix of a

corrvc	moment matrix. Computes correlation matrix from a variance- covariance matrix.
corrxx	Computes correlation matrix.
corrxxs	Computes sample correlation matrix.
crossprd	Computes cross product.
design	Creates a design matrix of 0's and 1's.
dstatmt	Computes descriptive statistics of a data set or matrix.
dot	Computes a scalar dot product of the columns of two matrices.
dstatmtControlCreate	Creates default dstatmtControl structure.
gdaDStat	Computes descriptive statistics on multiple Nx1 variables in a GDA.
gdaDStatMat	Computes descriptive statistics on a selection of columns in a variable in a GDA.
glm	Computes generalized linear regression of a matrix.
loess	Computes coefficients of locally weighted regression.
loessmt	Computes coefficients of locally weighted regression.
loessmtControlCreate	Creates default loessmtControl structure.
meanc	Computes mean value of each column of a matrix.
median	Computes medians of the columns of a

	matrix.
moment	Computes moment matrix ($x'x$) with special handling of missing values.
momentd	Computes moment matrix from a data set.
movingave	Computes moving average of a series.
movingaveExpwgt	Computes exponentially weighted moving average of a series.
movingaveWgt	Computes weighted moving average of a series.
numCombinations	Computes number of combinations of n things taken k at a time.
ols	Computes least squares regression of data set or matrix.
olsmt	Computes least squares regression of data set or matrix.
olsmtControlCreate	Creates default olsmtControl structure.
olsqr	Computes OLS coefficients using QR decomposition.
olsqr2	Computes OLS coefficients, residuals, and predicted values using QR decomposition.
olsqrmt	Computes OLS coefficients using QR decomposition.
pacf	Computes sample partial autocorrelations.
princomp	Computes principal components of a data matrix.
quantile	Computes quantiles from data in a

		matrix, given specified probabilities.
	quantiled	Computes quantiles from data in a data set, given specified probabilities.
	rndvm	Computes von Mises pseudo-random numbers.
	stdc	Computes standard deviation of the columns of a matrix.
	stdsc	Computes the 'sample' standard deviation of the elements in each column of a matrix.
	toeplitz	Computes Toeplitz matrix from column vector.
	varCovM	Computes the population variance-covariance matrix from a moment matrix.
	varCovMS	Computes a sample variance-covariance matrix from a moment matrix.
	varCovX	Computes the population variance-covariance matrix from a data matrix.
	varCovXS	Computes a sample variance-covariance matrix from a data matrix.
	varmall	Computes the log-likelihood of a Vector ARMA model.
	varmares	Computes the residuals of a Vector ARMA model.
	vcm	Computes a variance-covariance matrix from a moment matrix.
	vcms	Computes a sample variance-covariance matrix from a moment matrix.

vcx	Computes a variance-covariance matrix from a data matrix.
vcxs	Computes a sample variance-covariance matrix from a data matrix.

Advanced statistics and optimization routines are available in the **GAUSS** Applications programs. (Contact Aptech Systems for more information.)

34.1.10 Optimization and Solution

eqSolve	Solves a system of nonlinear equations.
eqSolvemt	Solves a system of nonlinear equations.
eqSolvemtControlCreate	Creates default eqSolvemtControl structure.
eqSolvemtOutCreate	Creates default eqSolvemtOut structure.
eqSolveSet	Sets global input used by eqSolve to default values.
ldlsol	Solves $LDL^T x = b$ using a matrix factorized by ldlp .
linsolve	Solves $Ax = b$ using the inverse function.
ltrisol	Computes the solution of $Lx = b$ where L is a lower triangular matrix.
lusol	Computes the solution of $LUx = b$ where L is a lower triangular matrix and U is an upper triangular matrix.

QNewton	Optimizes a function using the BFGS descent algorithm.
QNewtonmt	Minimizes an arbitrary function.
QNewtonmtControlCreate	Creates default QNewtonmtControl structure.
QNewtonmtOutCreate	Creates default QNewtonmtOut structure.
QProg	Solves the quadratic programming problem.
QProgmt	Solves the quadratic programming problem.
QProgmtInCreate	Creates an instance of a structure of type QProgmtInCreate with the maxit member set to a default value.
sqpSolve	Solves the nonlinear programming problem using a sequential quadratic programming method.
sqpSolveMT	Solves the nonlinear programming problem using a sequential quadratic programming method.
sqpSolveMTControlCreate	Creates an instance of a structure of type sqpSolveMTcontrol set to default values.
sqpSolveMTlagrangeCreate	Creates an instance of a structure of type sqpSolveMTlagrange set to default values.
sqpSolveMToutCreate	Creates an instance of a structure of type sqpSolveMTout set to

	default values.
sqpSolveSet	Resets global variables used by sqpSolve to default values.
utrisol	Computes the solution of $Ux = b$ where U is an upper triangular matrix.

34.1.11 Statistical Distributions

cdfBeta	Computes integral of beta function.
cdfBetaInv	Computes the quantile or inverse of the beta cumulative distribution function.
cdfBinomial	Computes the binomial cumulative distribution function.
cdfBinomialInv	Computes the binomial quantile or inverse cumulative distribution function.
cdfBvn	Computes lower tail of bivariate Normal cdf.
cdfBvn2	Returns cdfbvn of a bounded rectangle.
cdfBvn2e	Returns cdfbvn of a bounded rectangle.
cdfCauchy	Computes the cumulative distribution function for the Cauchy distribution.
cdfCauchyinv	Computes the Cauchy inverse cumulative distribution function.
cdfChic	Computes complement of cdf of χ^2 .
cdfChii	Computes χ^2 abscissae values given probability and degrees of freedom.
cdfChinc	Computes integral of noncentral χ^2 .
cdfEmpirical	Computes the cumulative distribution function for the empirical distribution.

cdfExp	Computes the cumulative distribution function for the exponential distribution.
cdfExpInv	Computes the exponential inverse cumulative distribution function.
cdfFc	Computes complement of cdf of F .
cdfFnc	Computes integral of noncentral F .
cdfFncInv	Computes the quantile or inverse of noncentral F cumulative distribution function.
cdfGam	Computes integral of incomplete Γ function.
cdfGenPareto	Computes the cumulative distribution function for the Generalized Pareto distribution.
cdfHyperGeo	Computes the cumulative distribution function of the hypergeometric distribution.
cdfLaplace	Computes the cumulative distribution function for the Laplace distribution.
cdfLaplaceInv	Computes the Laplace inverse cumulative distribution function.
cdfMvn	Computes multivariate Normal cdf.
cdfMvnce	Computes the complement of the multivariate Normal cumulative distribution function with error management
cdfMvne	Computes multivariate Normal cumulative distribution function with error management
cdfMvn2e	Computes the multivariate Normal cumulative distribution function with error management over the range $[a,b]$
cdfMvtce	Computes complement of multivariate Student's t cumulative distribution function

	with error management
cdfMvte	Computes multivariate Student's t cumulative distribution function with error management
cdfMvt2e	Computes multivariate Student's t cumulative distribution function with error management over [a,b]
cdfN	Computes integral of Normal distribution: lower tail, or cdf.
cdfN2	Computes interval of Normal cdf.
cdfNc	Computes complement of cdf of Normal distribution (upper tail).
cdfNegBinomial	Computes the cumulative distribution function for the negative binomial distribution.
cdfNegBinomialInv	Computes the quantile or inverse negative binomial cumulative distribution function.
cdfNi	Computes the inverse of the cdf of the Normal distribution.
cdfPoisson	Computes the Poisson cumulative distribution function.
cdfPoissonInv	Computes the quantile or inverse Poisson cumulative distribution function.
cdfRayleigh	Computes the Rayleigh cumulative distribution function.
cdfRayleighInv	Computes the Rayleigh inverse cumulative distribution function.
cdfTc	Computes complement of cdf of t -distribution.
cdfTci	Computes the inverse of the complement of

		the Student's t cdf.
cdfTnc		Computes integral of noncentral t -distribution.
cdfTvn		Computes lower tail of trivariate Normal cdf.
cdfWeibull		Computes the cumulative distribution function for the Weibull distribution.
cdfWeibullInv		Computes the Weibull inverse cumulative distribution function.
erf		Computes Gaussian error function.
erfc		Computes complement of Gaussian error function.
erfccplx		Computes complement of Gaussian error function for complex inputs.
erfcplx		Computes Gaussian error function for complex inputs.
lncdfbvn		Computes natural log of bivariate Normal cdf.
lncdfbvn2		Returns log of cdfbvn of a bounded rectangle.
lncdfmvn		Computes natural log of multivariate Normal cdf.
lncdfn		Computes natural log of Normal cdf.
lncdfn2		Computes natural log of interval of Normal cdf.
lncdfnc		Computes natural log of complement of Normal cdf.
lnpdfmvn		Computes multivariate Normal log-probabilities.
lnpdfmvt		Computes multivariate Student's t log-

	probabilities.
lnpdfn	Computes Normal log-probabilities.
lnpdft	Computes Student's t log-probabilities.
pdfBinomial	Computes the probability mass function for the binomial distribution.
pdfCauchy	Computes the probability density function for the Cauchy distribution.
pdfexp	Computes the probability density function for the exponential distribution.
pdfGenPareto	Computes the probability density function for the Generalized Pareto distribution.
pdfHyperGeo	Computes the probability mass function for the hypergeometric distribution.
pdfLaplace	Computes the probability density function for the Laplace distribution.
pdflogistic	Computes the probability density function for the logistic distribution.
pdfn	Computes standard Normal probability density function.
pdfPoisson	Computes the probability mass function for the Poisson distribution.
pdfRayleigh	Computes the probability density function of the Rayleigh distribution.
pdfWeibull	Computes the probability density function of a Weibull random variable.
pdfWishartInv	Computes the probability density function of a inverse Wishart distribution.

34.1.12 Series and Sequence Functions

recserar	Computes autoregressive recursive series.
recsercp	Computes recursive series involving products.
recserrc	Computes recursive series involving division.
sega	Creates an additive sequence.
seqm	Creates a multiplicative sequence.

34.1.13 Precision Control

base10	Converts number to $x.xxx$ and a power of 10.
ceil	Rounds up towards $+\infty$.
floor	Rounds down towards $-\infty$.
machEpsilon	Returns the smallest number such that $1 + \epsilon > 1$.
round	Rounds to the nearest integer.
trunc	Converts numbers to integers by truncating the fractional portion.

round, **trunc**, **ceil** and **floor** convert floating point numbers into integers. The internal representation for the converted integer is double precision (64 bits).

Each matrix element in memory requires 8 bytes of memory.

34.2 Finance Functions

AmericanBinomCall	American binomial method Call.
AmericanBinomCall_Greeks	American binomial method call Delta, Gamma, Theta, Vega, and Rho.
AmericanBinomCall_ImpVol	Implied volatilities for American binomial method calls.
AmericanBinomPut	American binomial method Put.
AmericanBinomPut_Greeks	American binomial method put Delta, Gamma, Theta, Vega, and Rho.

AmericanBinomPut_ImpVol	Implied volatilities for American binomial method puts.
AmericanBSCall	American Black and Scholes Call.
AmericanBSCall_Greeks	American Black and Scholes call Delta, Gamma, Omega, Theta, and Vega.
AmericanBSCall_ImpVol	Implied volatilities for American Black and Scholes calls.
AmericanBSPut	American Black and Scholes Put.
AmericanBSPut_Greeks	American Black and Scholes put Delta, Gamma, Omega, Theta, and Vega.
AmericanBSPut_ImpVol	Implied volatilities for American Black and Scholes puts.
annualTradingDays	Computes number of trading days in a given year.
elapsedTradingDays	Computes number of trading days between two dates inclusively.
EuropeanBinomCall	European binomial method call.
EuropeanBinomCall_Greeks	European binomial method call Delta, Gamma, Theta, Vega and Rho.
EuropeanBinomCall_ImpVol	Implied volatilities for European binomial method calls.
EuropeanBinomPut	European binomial method Put.
EuropeanBinomPut_Greeks	European binomial method put Delta, Gamma, Theta, Vega, and Rho.
EuropeanBinomPut_ImpVol	Implied volatilities for European binomial method puts.
EuropeanBSCall	European Black and Scholes Call.

EuropeanBSCall_Greeks	European Black and Scholes call Delta, Gamma, Omega, Theta, and Vega.
EuropeanBSCall_ImpVol	Implied volatilities for European Black and Scholes calls.
EuropeanBSPut	European Black and Scholes Put.
EuropeanBSPut_Greeks	European Black and Scholes put Delta, Gamma, Omega, Theta, and Vega.
EuropeanBSPut_ImpVol	Implied volatilities for European Black and Scholes puts.
getNextTradingDay	Returns the next trading day.
getNextWeekDay	Returns the next day that is not on a weekend.
getPreviousTradingDay	Returns the previous trading day.
getPreviousWeekDay	Returns the previous day that is not on a weekend.

34.3 Matrix Manipulation

34.3.1 Creating Vectors and Matrices

eye	Creates identity matrix.
matalloc	Allocates a matrix with unspecified contents.
matinit	Allocates a matrix with specified fill value.
ones	Creates a matrix of ones.
zeros	Creates a matrix of zeros.

Use **zeros**, **ones**, or **matinit** to create a constant vector or matrix.

Matrices can also be loaded from an ASCII file, from a **GAUSS** matrix file, or from a **GAUSS** data set. (See **DATA IMPORT/EXPORT**, CHAPTER 1, for more information.)

34.3.2 Loading and Storing Matrices

csvReadM	Loads data from an CSV, or other delimited text file, into a GAUSS matrix.
dataload	Loads matrices, N-dimensional arrays, strings and string arrays from a disk file.
datasave	Saves matrices, N-dimensional arrays, strings and string arrays to a disk file.
load, loadm	Loads a matrix from a matrix (FMT) file.
loadd	Loads matrix from data set (DAT file).
loadf	Loads function from disk file.
loadk	Loads keyword from disk file.
datasave	Saves matrices, N-dimensional arrays, strings and string arrays to a disk file.
load, loadm	Loads a matrix from a matrix (FMT) file.
loadd	Loads matrix from data set (DAT file).
loadf	Loads function from disk file.
loadk	Loads keyword from disk file.
save	Saves symbol to disk file.
saved	Saves matrix to data set.

xlsReadM

Loads data from an XLS or XLSX file into a **GAUSS** matrix.

34.3.3 Size, Ranking, and Range

Commands by
Category

cols

Returns number of columns in a matrix.

colsf

Returns number of columns in an open data set.

counts

Returns number of elements of a vector falling in specified ranges.

countwts

Returns weighted count of elements of a vector falling in specified ranges.

cumprodc

Computes cumulative products of each column of a matrix.

cumsumc

Computes cumulative sums of each column of a matrix.

indexcat

Returns indices of elements falling within a specified range.

maxc

Returns largest element in each column of a matrix.

maxindc

Returns row number of largest element in each column of a matrix.

minc

Returns smallest element in each column of a matrix.

minindc

Returns row number of smallest element in each column of a matrix.

prodc

Computes the product of each column of a matrix.

rankindx

Returns rank index of Nx1 vector.

	(Rank order of elements in vector).
rows	Returns number of rows in a matrix.
rowsf	Returns number of rows in an open data set.
sumc	Computes the sum of each column of a matrix.
sumr	Computes the sum of each row of a matrix.

These functions are used to find the minimum, maximum and frequency counts of elements in matrices.

Use **rows** and **cols** to find the number of rows or columns in a matrix. Use **rowsf** and **colsf** to find the numbers of rows or columns in an open **GAUSS** data set.

34.3.4 Miscellaneous Matrix Manipulation

complex	Creates a complex matrix from two real matrices.
delif	Deletes rows from a matrix using a logical expression.
delrows	Deletes rows from a matrix; the second argument contains the indices of the rows to be deleted.
diag	Extracts the diagonal of a matrix.
diagrv	Puts a column vector into the diagonal of a matrix.
exctsmpl	Creates a random subsample of a data set, with replacement.
imag	Returns the imaginary part of a complex matrix.

indcv	Checks one character vector against another and returns the indices of the elements of the first vector in the second vector.
indnv	Checks one numeric vector against another and returns the indices of the elements of the first vector in the second vector.
intrsect	Returns the intersection of two vectors.
lowmat	Returns the main diagonal and lower triangle.
lowmat1	Returns a main diagonal of 1's and the lower triangle.
putvals	Inserts values into a matrix or N-dimensional array.
real	Returns the real part of a complex matrix.
reshape	Reshapes a matrix to new dimensions.
rev	Reverses the order of rows of a matrix.
rotater	Rotates the rows of a matrix, wrapping elements as necessary.
selif	Selects rows from a matrix using a logical expression.
setdif	Returns elements of one vector that are not in another.
shiftr	Shifts rows of a matrix, filling in holes with a specified value.

submat	Extracts a submatrix from a matrix.
subvec	Extracts an Nx1 vector of elements from an NxK matrix.
trimr	Trims rows from top or bottom of a matrix.
union	Returns the union of two vectors.
upmat	Returns the main diagonal and upper triangle.
upmat1	Returns a main diagonal of 1's and the upper triangle.
vec	Stacks columns of a matrix to form a single column.
vech	Reshapes the lower triangular portion of a symmetric matrix into a column vector.
vecr	Stacks rows of a matrix to form a single column.
vget	Extracts a matrix or string from a data buffer constructed with vput .
vlist	Lists the contents of a data buffer constructed with vput .
vnamecv	Returns the names of the elements of a data buffer constructed with vput .
vput	Inserts a matrix or string into a data buffer.
vread	Reads a string or matrix from a data buffer constructed with vput .
vtypecv	Returns the types of the elements of a data buffer constructed with vput .

xpnd Expands a column vector into a symmetric matrix.

vech and **xpnd** are complementary functions. **vech** provides an efficient way to store a symmetric matrix; **xpnd** expands the stored vector back to its original symmetric matrix.

delif and **selif** are complementary functions. **delif** deletes rows of a matrix based on a logical comparison; **selif** selects rows based on a logical comparison.

lowmat, **lowmat1**, **upmat**, and **upmat1** extract triangular portions of a matrix.

To delete rows which contain missing values from a matrix in memory, see **packr**.

34.4 Sparse Matrix Handling

denseToSp Converts a dense matrix to a sparse matrix.

denseToSpRE Converts a dense matrix to a sparse matrix using a relative epsilon.

packedToSp Creates a sparse matrix from a packed matrix of non-zero values and row and column indices.

spBiconjGradSol Solves the system of linear equations $Ax=b$ using the biconjugate gradient method.

spChol Computes the LL' decomposition of a sparse matrix.

spConjGradSol Solves the system of linear equations $Ax=b$ for symmetric matrices using the conjugate gradient method.

spCreate Creates a sparse matrix from vectors of non-zero values, row indices, and

	column indices.
spDenseSubmat	Returns a dense submatrix of a sparse matrix.
spDiagRvMat	Inserts submatrices along the diagonal of a sparse matrix.
spEigv	Computes a specified number of eigenvalues and eigenvectors of a square, sparse matrix.
spEye	Creates a sparse identity matrix.
spGetNZE	Returns the non-zero values in a sparse matrix, as well as their corresponding row and column indices.
spGetNumNZE	Returns the number of non-zero elements in a sparse matrix.
spLDL	Computes the LDL decomposition of a symmetric sparse matrix.
spLU	Computes the LU decomposition of a sparse matrix with partial pivoting.
spOnes	Generates a sparse matrix containing only ones and zeros
spSubmat	Returns a sparse submatrix of sparse matrix.
spToDense	Converts a sparse matrix to a dense matrix.
spTrTDense	Multiplies a sparse matrix transposed by a dense matrix.
spTScalar	Multiplies a sparse matrix by a scalar.
spZeros	Creates a sparse matrix containing no

non-zero values.

34.5 N-Dimensional Array Handling

34.5.1 Creating Arrays

aconcat	Concatenates conformable matrices and arrays in a user-specified dimension.
aeye	Creates an N-dimensional array in which the planes described by the two trailing dimensions of the array are equal to the identity.
areshape	Reshapes a scalar, matrix, or array into an array of user-specified size.
arrayalloc	Creates an N-dimensional array with unspecified contents.
arrayinit	Creates an N-dimensional array with a specified fill value.
mattoarray	Converts a matrix to a type array.

34.5.2 Size, Ranking and Range

amax	Moves across one dimension of an N-dimensional array and finds the largest element.
amin	Moves across one dimension of an N-dimensional array and finds the smallest element.
asum	Computes the sum across one dimension of an N-dimensional array.

getdims	Gets the number of dimensions in an array.
getorders	Gets the vector of orders corresponding to an array.

34.5.3 Setting and Retrieving Data in an Array

aconcat	Concatenates conformable matrices and arrays in a user-specified dimension.
areshape	Reshapes a scalar, matrix, or array into an array of user-specified size.
arraytomat	Changes an array to type matrix.
getarray	Gets a contiguous subarray from an N-dimensional array.
getmatrix	Gets a contiguous matrix from an N-dimensional array.
getmatrix4D	Gets a contiguous matrix from a 4-dimensional array.
getscalar3D	Gets a scalar from a 3-dimensional array.
getscalar4D	Gets a scalar from a 4-dimensional array.
putarray	Puts a contiguous subarray into an N-dimensional array and returns the resulting array.
setarray	Sets a contiguous subarray of an N-dimensional array.

34.5.4 Miscellaneous Array Functions

Commands by
Category

<code>amean</code>	Computes the mean across one dimension of an N-dimensional array.
<code>amult</code>	Performs matrix multiplication on the planes described by the two trailing dimensions of N-dimensional arrays.
<code>arrayindex</code>	Saves a matrix of structures to a file on the disk.
<code>atranspose</code>	Transposes an N-dimensional array.
<code>loopnextindex</code>	Increments an index vector to the next logical index and jumps to the specified label if the index did not wrap to the beginning.
<code>nextindex</code>	Returns the index of the next element or subarray in an array.
<code>previousindex</code>	Returns the index of the previous element or subarray in an array.
<code>singleindex</code>	Converts a vector of indices for an N-dimensional array to a scalar vector index.
<code>walkindex</code>	Walks the index of an array forward or backward through a specified dimension.

34.6 Structures

<code>dsCreate</code>	Creates an instance of a structure of type DS set to default values.
<code>loadstruct</code>	Loads a structure into memory from a

	file on the disk.
pvCreate	Returns an initialized an instance of structure of type PV .
pvGetIndex	Gets row indices of a matrix in a parameter vector.
pvGetParNames	Generates names for parameter vector stored in structure of type PV .
pvGetParVector	Retrieves parameter vector from structure of type PV .
pvLength	Returns the length of a parameter vector.
pvList	Retrieves names of packed matrices in structure of type PV .
pvPack	Packs general matrix into a structure of type PV with matrix name.
pvPacki	Packs general matrix or array into a PV instance with name and index.
pvPackm	Packs general matrix into a structure of type PV with a mask and matrix name.
pvPackmi	Packs general matrix or array into a PV instance with a mask, name, and index.
pvPacks	Packs symmetric matrix into a structure of type PV .
pvPacksi	Packs symmetric matrix into a PV instance with matrix name and index.
pvPacksm	Packs symmetric matrix into a structure of type PV with a mask.

pvPacksmi	Packs symmetric matrix into a PV instance with a mask, matrix name, and index.
pvPutParVector	Inserts parameter vector into structure of type PV .
pvTest	Tests an instance of structure of type PV to determine if it is a proper structure of type PV .
pvUnpack	Unpacks matrices stored in a structure of type PV .
savestruct	Saves a matrix of structures to a file on the disk.

34.7 Data File Reading/Writing

34.7.1 Spreadsheets

spreadSheetReadM	Reads numeric data from an Excel file into a GAUSS matrix.
spreadSheetReadSA	Reads text or string data from an Excel file into a GAUSS string array.
spreadSheetWrite	Writes numeric or string data to an Excel file.
xlsGetSheetCount	Gets the number of sheets in an Excel spreadsheet.
xlsGetSheetSize	Gets the size (rows and columns) of a specified sheet in an Excel spreadsheet.
xlsGetSheetTypes	Gets the cell format types of a row in an Excel spreadsheet.

xlsMakeRange	Builds an Excel range string from a row/column pair.
xlsReadM	Reads from an Excel spreadsheet, into a GAUSS matrix.
xlsReadSA	Reads from an Excel spreadsheet, into a GAUSS string array or string.
xlsWrite	Writes a GAUSS matrix, string, or string array to an Excel spreadsheet.
xlsWriteM	Writes a GAUSS matrix to an Excel spreadsheet.
xlsWriteSA	Writes a GAUSS string or string array to an Excel spreadsheet.

34.7.2 CSV Files

csvReadM	Reads numeric data from a CSV file into a matrix.
csvReadSA	Reads text or string data from a CSV file into a GAUSS string array.

34.7.3 Text Files

fcheckerr	Gets the error status of a file.
fclearerr	Gets the error status of a file, then clears it.
fflush	Flushes a file's output buffer.
fgets	Reads a line of text from a file.
fgetsa	Reads lines of text from a file into a string array.
fgetsat	Reads lines of text from a file into a string array.

fgetst	Reads a line of text from a file.
fopen	Opens a file.
fputs	Writes strings to a file.
fputst	Writes strings to a file.
fseek	Positions the file pointer in a file.
fstrerror	Returns an error message explaining the cause of the most recent file I/O error.
ftell	Gets the position of the file pointer in a file.

34.7.4 HDF 5 Files

h5create	Create a HDF5 dataset (.h5).
h5open	Open a HDF5 file.
h5read	Reads data from a HDF5 file (.h5) into a GAUSS matrix.
h5readAttribute	Read attributes from a HDF5 file into GAUSS.
h5write	Write a GAUSS matrix to a HDF5 file.
h5writeAttribute	Write a GAUSS string array as attributes into a HDF5 file

34.7.5 Database

This section summarizes all procedures within the GAUSS database module. A general usage description will be found in **Databases with GAUSS**, Section 1 .

Database Setup

dbAddDatabase	Adds a database to the list of database
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	connections using the driver type or a connection URL.
dbGetDrivers	Returns a list of available database drivers.
dbIsDriverAvailable	Returns 1 if a specified database driver is available.
dbRemoveDatabase	Removes a database connection from the list of open database connections. Frees all related resources.

Database Properties

dbGetConnectOptions	Returns the connection options string used for a database connection.
dbGetDatabaseName	Returns the name of the database.
dbGetDriverName	Returns the name of the connection's database driver.
dbGetHostName	Returns the database connection's host name.
dbGetPassword	Returns a connection's password.
dbGetNumericalPrecPolicy	Returns the default numerical precision policy for a specified database connection.
dbGetPort	Returns the database connection's port number if it has been set.
dbIsOpen	Reports whether a specified database connection is open.
dbIsValid	Reports whether a specified database connection has a valid driver.

dbSetConnectOptions	Sets database-specific options.
dbSetDatabaseName	Sets the connection's database name to name.
dbSetHostName	Sets the specified database connection's host name.
dbSetNumericalPrecPolicy	Sets the default numerical precision policy used by queries created on this database connection.
dbSetPassword	Sets the database connection's password.
dbSetPort	Sets the specified database connection's port number.

Database Information

dbGetPrimaryIndex	Returns the primary index for the specified table.
dbGetTableHeaders	Returns a string array populated with the names of all the fields in a specified table (or view).
dbGetTables	Returns the database's tables, system tables and views.
dbHasFeature	Returns a 1 if the database supports the specified feature.

Database Errors

dbGetLastErrorNum	Returns numerical information about the last error that occurred on the database.
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dbGetLastErrorText	Returns text information about the last error that occurred on the database.
dbIsOpenError	Reports whether an error occurred while attempting to open the database connection.
dbQueryGetLastErrorNum	Returns numerical error information about the last error that occurred (if any) with the last executed query.
dbQueryGetLastErrorText	Returns text error information about the last error that occurred (if any) with the last executed query.

Database Connect

dbClose	Closes a database connection and destroys any remaining queries.
dbOpen	Opens a specified database connection using the current connection values.

Database Transaction

dbCommit	Commits a transaction to the database if the driver supports transactions and a dbTransaction() has been started.
dbCreateQuery	Process an SQL statement and prepare a query.
dbExecQuery	Executes an SQL statement and creates a query.
dbRollback	Rolls back a transaction on the database.

dbTransaction Begins a transaction on the database.

Query Building

dbQueryBindValue Set the placeholder placeholder to be bound to value val in the prepared statement.

dbQueryGetBoundValue Returns the value for a placeholder in a query.

dbQueryGetBoundValues Returns an Nx2 string array containing the placeholders and their corresponding values in a query.

dbQueryExecPrepared Executes a previously created and prepared query.

dbQueryPrepare Prepares a SQL query for execution.

Query Manipulation

dbQueryClear Clears the result set and releases any resources held by the query. Sets the query state to inactive.

dbQueryFinish Instructs the database driver that no more data will be fetched from this query until it is re-executed.

Query Information

dbQueryCols Returns the number of fields in the record.

dbQueryGetLastInsertID Returns the object ID of the most recent inserted row if supported by the database.

dbQueryGetLastQuery	Returns the text of the current query being used.
dbQueryGetNumRowsAffected	Reports the number of rows affected by the result's SQL statement.
dbQueryIsActive	Returns 1 if the query is active.
dbQueryIsForwardOnly	Reports whether you can only scroll forward through a result set.
dbQueryIsNull	Reports whether the current field pointed at by an active query positioned on a valid record is NULL.
dbQueryIsSelect	Reports whether the specified query is a SELECT statement.
dbQueryIsValid	Reports whether the specified query is positioned on a valid record.
dbQueryRows	Returns the size of the result (number of rows returned), or -1 if the size cannot be determined or if the database does not support reporting information about query sizes.
dbQuerySetForwardOnly	Sets forward only mode to forward. If forward is true, only dbQuerySeekNext() and dbQuerySeek() with positive values, are allowed for navigating the results.

Query Iteration

<code>dbQueryGetPosition</code>	Returns the current internal position of the query.
<code>dbQuerySeek</code>	Retrieves the record at a specified position, if available, and positions the query on the retrieved record.
<code>dbQuerySeekFirst</code>	Retrieves the first record in the result, if available, and positions the query on the retrieved record.
<code>dbQuerySeekLast</code>	Retrieves the last record in the result, if available, and positions the query on the retrieved record.
<code>dbQuerySeekNext</code>	Retrieves the next record in the result, if available, and positions the query on the retrieved record.
<code>dbQuerySeekPrevious</code>	Retrieves the previous record in the result, if available, and positions the query on the retrieved record.

Query Data Retrieval

<code>dbQueryFetchAllM</code>	Returns the result set for the current query as a matrix.
<code>dbQueryFetchAllSA</code>	Returns the result set for the current query as a string array.
<code>dbQueryFetchOneM</code>	Returns a single row as an Nx1 matrix where N is the column count of the SELECT statement.
<code>dbQueryFetchOneSA</code>	Returns a single row as a string vector containing the field information for

dbQueryGetField	the current query. Returns the value of a specified field in the current record.
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GAUSS Data Archives

gdaAppend	Appends data to a variable in a GDA.
gdaCreate	Creates a GDA.
gdaDStat	Computes descriptive statistics on multiple Nx1 variables in a GDA.
gdaDStatMat	Computes descriptive statistics on a selection of columns in a variable in a GDA.
gdaGetIndex	Gets the index of a variable in a GDA.
gdaGetName	Gets the name of a variable in a GDA.
gdaGetNames	Gets the names of all the variables in a GDA.
gdaGetOrders	Gets the orders of a variable in a GDA.
gdaGetType	Gets the type of a variable in a GDA.
gdaGetTypes	Gets the types of all the variables in a GDA.
gdaGetVarInfo	Gets information about all of the variables in a GDA.
gdaIsCplx	Checks to see if a variable in a GDA is complex.
gdaLoad	Loads variables in a GDA into the

gdaPack	workspace. Packs the data in a GDA, removing all empty bytes.
gdaRead	Gets a variable from a GDA.
gdaReadByIndex	Gets a variable from a GDA, given a variable index.
gdaReadSome	Reads part of a variable from a GDA.
gdaReadSparse	Gets a sparse matrix from a GAUSS Data Archive.
gdaReadStruct	Gets a structure from a GAUSS Data Archive.
gdaReportVarInfo	Gets information about all of the variables in a GAUSS Data Archive and returns it in a string array formatted for printing.
gdaSave	Writes variables in a workspace to a GDA.
gdaUpdate	Updates a variable in a GDA.
gdaUpdateAndPack	Updates a variable in a GDA, leaving no empty bytes if the updated variable is smaller or larger than the variable it is replacing.
gdaWrite	Writes a variable to a GDA.
gdaWrite32	Writes a variable to a GDA using 32-bit system file write commands.
gdaWriteSome	Overwrites part of a variable in a GDA.

These functions all operate on **GAUSS** Data Archives (GDA's). For more information, see **GAUSS Data Archives**, Section 1.0.1.

Data Sets

close	Closes an open data set (.dat file).
closeall	Closes all open data sets.
create	Creates and opens a data set.
datacreate	Creates a v96 real data set.
datacreatecomplex	Creates a v96 complex data set.
datalist	Lists selected variables from a data set.
dataopen	Opens a data set.
eof	Tests for end of file.
getnr	Computes number of rows to read per iteration for a program that reads data from a disk file in a loop.
getnrmt	Computes number of rows to read per iteration for a program that reads data from a disk file in a loop.
iscplx	Returns whether a data set is real or complex.
load	Loads a small data set.
open	Opens an existing data set.
readr	Reads rows from open data set.
saved	Creates small data sets.
seekr	Moves pointer to specified location in open data set.
tempname	Creates a temporary file with a unique name.
typef	Returns the element size (2, 4 or 8 bytes) of data in open data set.

writer Writes matrix to an open data set.

These functions all operate on **GAUSS** data sets (`.dat` files). For more information, see **DATA IMPORT/EXPORT**, CHAPTER 1.

To create a **GAUSS** data set from a matrix in memory, use **saved**. To create a data set from an existing one, use **create**. To create a data set from a large ASCII file, use the `ATOG` utility (see **ATOG**, CHAPTER 1.)

Data sets can be opened, read from, and written to using **open**, **readr**, **seekr** and **writer**. Test for the end of a file using **eof**, and close the data set using **close** or **closeall**.

The data in data sets may be specified as character or numeric. (See **DATA IMPORT/EXPORT**, CHAPTER 1.) See also **create** and **vartypef**.

typef returns the element size of the data in an open data set.

Data Set Variable Names

getname	Returns column vector of variable names in a data set.
getnamef	Returns string array of variable names in a data set.
indices	Retrieves column numbers and names from a data set.
indices2	Similar to indices , but matches columns with names for dependent and independent variables.
indicesf	Retrieves column numbers and names from a data set.
indicesfn	Retrieves column numbers and names from a data set.
makevars	Decomposes matrix to create column vectors.

setvars	Creates globals using the names in a data set.
vartypef	Returns column vector of variable types (numeric/character) in a data set.

Use **getnamef** to retrieve the variable names associated with the columns of a **GAUSS** data set and **vartypef** to retrieve the variable types. Use **makevars** and **setvars** to create global vectors from those names. Use **indices** and **indices2** to match names with column numbers in a data set.

Data Coding

code	Codes the data in a vector by applying a logical set of rules to assign each data value to a category.
code (dataloop)	Creates new variables with different values based on a set of logical expressions.
dataloop (dataloop)	Specifies the beginning of a data loop.
delete (dataloop)	Removes specific rows in a data loop based on a logical expression.
drop (dataloop)	Specifies columns to be dropped from the output data set in a data loop.
dummy	Creates a dummy matrix, expanding values in vector to rows with ones in columns corresponding to true categories and zeros elsewhere.
dummybr	Similar to dummy .
dummydn	Similar to dummy .
extern (dataloop)	Allows access to matrices or strings in memory from inside a data loop.

isinfnanmiss	Returns true if the argument contains an infinity, NaN, or missing value.
scalmiss	Returns 1 if matrix has any missing values, 0 otherwise.
keep (dataloop)	Specifies columns (variables) to be saved to the output data set in a data loop.
lag (dataloop)	Lags variables a specified number of periods.
lag1	Lags a matrix by one time period for time series analysis.
lagN	Lags a matrix a specified number of time periods for time series analysis.
listwise (dataloop)	Controls listwise deletion of missing values.
make (dataloop)	Specifies the creation of a new variable within a data loop.
miss	Changes specified values to missing value code.
missex	Changes elements to missing value using logical expression.
missrv	Changes missing value codes to specified values.
msym	Sets symbol to be interpreted as missing value.
outtyp (dataloop)	Specifies the precision of the output data set.
packr	Delete rows with missing values.
reclassify	Replaces specified values of a matrix,

	array or string array
reclassifyCuts	Replaces values of a matrix or array within specified ranges
recode	Changes the values of an existing vector from a vector of new values. Used in data transformations.
recode (dataloop)	Changes the value of a variable with different values based on a set of logical expressions.
scalinfnanmiss	Returns true if the argument is a scalar infinity, NaN, or missing value.
scalmiss	Tests whether a scalar is the missing value code.
select (dataloop)	Selects specific rows (observations) in a data loop based on a logical expression.
subscat	Simpler version of recode , but uses ascending bins instead of logical conditions.
substute	Similar to recode , but operates on matrices.
vector (dataloop)	Specifies the creation of a new variable within a data loop.

code, **recode**, and **subscat** allow the user to code data variables and operate on vectors in memory. **substute** operates on matrices, and **dummy**, **dummybr** and **dummydn** create matrices.

missex, **missrv** and **miss** should be used to recode missing values.

Sorting and Merging

intrleav

Produces one large sorted data file from two smaller sorted files having the same keys.

intrleavsa

Interleaves the rows of two string arrays that have been sorted on a common column.

mergeby

Produces one large sorted data file from two smaller sorted files having a single key column in common.

mergevar

Accepts a list of names of global matrices, and concatenates the corresponding matrices horizontally to form a single matrix.

sortc

Quick-sorts rows of matrix based on numeric key.

sortcc

Quick-sorts rows of matrix based on character key.

sortd

Sorts data set on a key column.

sorthc

Heap-sorts rows of matrix based on numeric key.

sorthcc

Heap-sorts rows of matrix based on character key.

sortind

Returns a sorted index of a numeric vector.

sortindc

Returns a sorted index of a character vector.

sortmc

Sorts rows of matrix on the basis of multiple columns.

sortr	Sorts rows of a matrix of numeric data.
sortrc	Sorts rows of a matrix of character data.
uniqindx	Returns a sorted unique index of a vector.
uniqindxsa	Computes the sorted index of a string vector, omitting duplicate elements.
unique	Removes duplicate elements of a vector.
uniquesa	Removes duplicate elements from a string vector.

sortc, **sorthc**, and **sortind** operate on numeric data only. **sortcc**, **sorthcc**, and **sortindc** operate on character data only.

sortd, **sortmc**, **unique**, and **uniqindx** operate on both numeric and character data.

Use **sortd** to sort the rows of a data set on the basis of a key column.

Both **intrleav** and **mergeby** operate on data sets.

34.8 Compiler Control

#define	Defines a case-insensitive text-replacement or flag variable.
#definesc	Defines a case-sensitive text-replacement or flag variable.
#else	Alternates clause for #if-#else-#endif code block.
#endif	End of #if-#else-#endif code block.

Commands by Category

Commands by Category

<code>#ifdef</code>	Compiles code block if a variable has been <code>#define</code> 'd.
<code>#iflight</code>	Compiles code block if running GAUSS Light.
<code>#ifndef</code>	Compiles code block if a variable has not been <code>#define</code> 'd.
<code>#ifmac</code>	Compiles code block if running on Mac.
<code>#ifos2win</code>	Compiles code block if running Windows.
<code>#ifunix</code>	Compiles code block if running UNIX.
<code>#include</code>	Includes code from another file in program.
<code>#linesoff</code>	Compiles program without line number and file name records.
<code>#lineson</code>	Compiles program with line number and file name records.
<code>#srcfile</code>	Inserts source file name record at this point (currently used when doing data loop translation).
<code>#srcline</code>	Inserts source file line number record at this point (currently used when doing data loop translation).
<code>#undef</code>	Undefines a text-replacement or flag variable.

These commands are compiler directives. That is, they do not generate **GAUSS** program instructions; rather, they are instructions that tell **GAUSS** how to process a program during compilation. They determine what the final compiled form of a

program will be. They are not executable statements and have no effect at run-time. (See **COMPILER DIRECTIVES**, CHAPTER 1.1, for more information.)

34.9 Multi-Threading

<code>threadBegin</code>	Marks beginning of a block of code to be executed as a thread.
<code>threadEnd</code>	Marks end of a block of code to be executed as a thread.
<code>threadEndFor</code>	Marks end of a parallel <code>for</code> loop.
<code>threadFor</code>	Marks the start of a parallel <code>for</code> loop.
<code>threadJoin</code>	Completes definition of a set of threads, waits for their work.
<code>threadStat</code>	Marks a single statement to be executed as a thread.

Together, `threadBegin/threadEnd` and `threadStat` define a set of threads that will execute simultaneously. `threadJoin` completes the definition of that set. `threadJoin` waits for the threads in the set to finish their calculations, the results of which are then available for further use.

```
threadBegin; // Thread 1
  y = x'x;
  z = y'y;
threadEnd;
threadBegin; // Thread 2
  q = r'r;
  r = q'q;
threadEnd;
threadStat n = m'm; // Thread 3
threadStat p = o'o; // Thread 4
threadJoin; // waits for Threads 1-4 to finish
```

```
b = z + r + n'p; // Using the results
```

34.10 Program Control

34.10.1 Execution Control

<code>call</code>	Calls function and discards return values.
<code>end</code>	Terminates a program and closes all files.
<code>pause</code>	Pauses for the specified time.
<code>run</code>	Runs a program in a text file.
<code>sleep</code>	Sleeps for the specified time.
<code>stop</code>	Stops a program and leaves files open.
<code>system</code>	Quits and returns to the OS.

Both `stop` and `end` will terminate the execution of a program; `end` will close all open files, and `stop` will leave those files open. Neither `stop` nor `end` is required in a **GAUSS** program.

34.10.2 Branching

<code>goto</code>	Unconditional branching.
<code>if...endif</code>	Conditional branching.
<code>pop</code>	Retrieves <code>goto</code> arguments.

```
if iter > itlim;
    goto errout("Iteration limit exceeded");
elseif iter =\,= 1;
    j = setup(x,y);
else;
```



```

        j = iterate (x,y);
    endif;

    .
    .
    .
errout:

    pop errmsg;
    print errmsg;
end;

```

34.10.3 Looping

`break`

Jumps out the bottom of a `do` or `for` loop.

`continue`

Jumps to the top of a `do` or `for` loop.

`do while...enddo`

Executes a series of statements in a loop as long as a given expression is TRUE (or FALSE).

`do until...enddo`

Loops if FALSE.

`for...endfor`

Loops with integer counter.

```

iter = 0;
do while dif > tol;
    { x,x0 } = eval (x,x0);
    dif = abs (x-x0);
    iter = iter + 1;
    if iter > maxits;
        break;
    endif;
    if not prtiter;
        continue;
    endif;
    format /rdn 1,0;

```

Commands by Category

```
print "Iteration: " iter;;
format /re 16,8;
print ", Error: " maxc(dif);
endo;

for i (1, cols(x), 1);
  for j (1, rows(x), 1);
    x[i,j] = x[i,j] + 1;
  endfor;
endfor;
```

34.10.4 Subroutines

<code>gosub</code>	Branches to subroutine.
<code>pop</code>	Retrieves <code>gosub</code> arguments.
<code>return</code>	Returns from subroutine.

Arguments can be passed to subroutines in the branch to the subroutine label and then popped, in first-in-last-out order, immediately following the subroutine label definition. See `gosub`.

Arguments can then be returned in an analogous fashion through the `return` statement.

34.10.5 Procedures, Keywords, and Functions

<code>endp</code>	Terminates a procedure definition.
<code>fn</code>	Allows user to create one-line functions.
<code>keyword</code>	Begins the definition of a keyword procedure. Keywords are user-defined functions with local or global variables.
<code>local</code>	Declares variables local to a procedure.
<code>proc</code>	Begins definition of multi-line procedure.
<code>retp</code>	Returns from a procedure.

Here is an example of a **GAUSS** procedure:

```

proc (3) = crosprod(x,y);
    local r1, r2, r3;
    r1 = x[2,.].*y[3,.]-x[3,.].*y[2,.];
    r2 = x[3,.].*y[1,.]-x[1,.].*y[3,.];
    r3 = x[1,.].*y[2,.]-x[2,.].*y[1,.];
    retp( r1,r2,r3);
endp;

```

The "(3) =" indicates that the procedure returns three arguments. All local variables, except those listed in the argument list, must appear in the `local` statement. Procedures may reference global variables. There may be more than one `retp` per procedure definition; none is required if the procedure is defined to return 0 arguments. The `endp` is always necessary and must appear at the end of the procedure definition. Procedure definitions cannot be nested. The syntax for using this example function is

```
{ a1,a2,a3 } = crosprod(u,v);
```

See **PROCEDURES AND KEYWORDS**, CHAPTER 1, and **LIBRARIES**, CHAPTER 1, for details.

34.10.6 Libraries

<code>declare</code>	Initializes variables at compile time.
<code>external</code>	External symbol definitions.
<code>lib</code>	Builds or updates a GAUSS library.
<code>library</code>	Sets up list of active libraries.

`call` allows functions to be called when return values are not needed. This is especially useful if a function produces printed output (`dstat`, `ols` for example) as well as return values.

34.10.7 Compiling

<code>compile</code>	Compiles and saves a program to a
----------------------	-----------------------------------

	<code>.gcg</code> file.
<code>#include</code>	Inserts code from another file into a GAUSS program.
<code>loadp</code>	Loads compiled procedure.
<code>save</code>	Saves the compiled image of a procedure to disk.
<code>saveall</code>	Saves the contents of the current workspace to a file.
<code>use</code>	Loads previously compiled code.

GAUSS procedures and programs may be compiled to disk files. By then using this compiled code, the time necessary to compile programs from scratch is eliminated. Use `compile` to compile a command file. All procedures, matrices and strings referenced by that program will be compiled as well.

Stand-alone applications may be created by running compiled code under the **GAUSS** Run-Time Module. Contact Aptech Systems for more information on this product.

To save the compiled images of procedures that do not make any global references, use `save`. This will create an `.fcg` file. To load the compiled procedure into memory, use `loadp`. (This is not recommended because of the restriction on global references and the need to explicitly load the procedure in each program that references it. It is included here to maintain backward compatibility with previous versions.)

34.10.8 Miscellaneous Program Control

<code>gausset</code>	Resets the global control variables declared in <code>gauss.dec</code> .
<code>sysstate</code>	Gets or sets general system parameters.

34.11 OS Functions and File Management

cd	Returns current directory.
changeDir	Changes the working directory in program.
chdir	Changes directory interactively.
deleteFile	Deletes files.
dlibrary	Dynamically links and unlinks shared libraries.
dllcall	Calls functions located in dynamic libraries.
dos	Provides access to the operating system from within GAUSS .
envget	Gets an environment string.
exec	Executes an executable program file.
execbg	Provides access to the operating system from within GAUSS .
fileinfo	Takes a file specification, returns names and information of files that match.
filesa	Takes a file specification, returns names of files that match.
getGAUSSHome	Returns a string with the full path to the GAUSS home directory.
getpath	Returns an expanded filename including the drive and path.
searchsourcepath	Searches the source path and (if specified) the src subdirectory of the GAUSS installation directory for a specified file.

`shell` Shells to OS.

34.12 Workspace Management

<code>clear</code>	Sets matrices equal to 0.
<code>clearg</code>	Sets global symbols to 0.
<code>delete</code>	Deletes specified global symbols.
<code>hasimag</code>	Examines matrix for nonzero imaginary part.
<code>iscplx</code>	Returns whether a matrix is real or complex.
<code>maxbytes</code>	Returns maximum memory to be read from a dataset at a time inside some GAUSS functions.
<code>maxvec</code>	Returns maximum allowed vector size.
<code>new</code>	Clears current workspace.
<code>show</code>	Displays global symbol table.
<code>type</code>	Returns type of argument (matrix or string).
<code>typecv</code>	Returns types of symbols (argument contains the names of the symbols to be checked).

When working with limited workspace, it is a good idea to `clear` large matrices that are no longer needed by your program.

34.13 Error Handling and Debugging

`debug` Executes a program under the source

error	debugger.
errorlog	Creates user-defined error code.
#linesoff	Sends error message to screen and log file.
#lineson	Omits line number and file name records from program.
scalerr	Includes line number and file name records in program.
trace	Tests for a scalar error code.
trap	Traces program execution for debugging.
trapchk	Controls trapping of program errors.
	Examines the trap flag.

To trace the execution of a program, use **trace**.

User-defined error codes may be generated using **error**.

34.14 String Handling

chrs	Converts ASCII values to a string.
convertsatotr	Converts a 1x1 string array to a string.
convertstrtosa	Converts a string to a 1x1 string array.
cvtos	Converts a character vector to a string.
cvtosa	Converts an NxK character vector to an NxK string array.
ftocv	Converts an NxK matrix to a

ftos	character matrix. Converts a floating point scalar to a string.
ftostrC	Converts a matrix to a string array using a C language format specification.
getf	Loads ASCII or binary file into string.
indsav	Checks one string array against another and returns
intrsect	Returns the intersection of two string vectors (or matrices), with duplicates removed.
loads	Loads a string file (.fst file).
lower	Converts a string to lowercase.
parse	Parses a string, returning a character vector of tokens.
putf	Writes a string to disk file.
stocv	Converts a string to a character vector.
stof	Converts a string to floating point numbers.
strcombine	Converts an NxM string array to an Nx1 string vector by combining each element in a column separated by a user-defined delimiter string.
strindx	Finds starting location of one string in another string.
strjoin	Converts an NxM string array to an Nx1 string vector by combining each

	element in a column separated by a user-defined delimiter string.
strlen	Returns length of a string.
strput	Lays a substring over a string.
strrindx	Finds starting location of one string in another string, searching from the end to the start of the string.
strsect	Extracts a substring of a string.
strsplit	Splits an Nx1 string vector into an NxK string array of the individual tokens.
strsplitPad	Splits an Nx1 string vector into an NxK string array of the individual tokens. Pads on the right with null strings.
strtof	Converts a string array to a numeric matrix.
strtofcplx	Converts a string array to a complex numeric matrix.
strtrim	Strips all whitespace characters from the left and right side of each element in a string array.
strtriml	Strips all whitespace characters from the left side of each element in a string array.
strtrimr	Strips all whitespace characters from the right side of each element in a string array.
strtrunc	Truncates all elements of a string array to not longer than the specified

strtrunc1	number of characters. Truncates the left side of all elements of a string array by a user-specified number of characters.
strtruncpad	Truncates all elements of a string array to the specified number of characters, adding spaces on the end as needed to achieve the exact length.
strtruncr	Truncates the right side of all elements of a string array by a user-specified number of characters.
token	Extracts the leading token from a string.
upper	Changes a string to uppercase.
vals	Converts a string to ASCII values.
varget	Accesses the global variable named by a string.
varget1	Accesses the local variable named by a string.
varput	Assigns a global variable named by a string.
varput1	Assigns a local variable named by a string.
 strlen , strindx , strrindx , and strsect can be used together to parse strings.	
Use ftos to print to a string.	

To create a list of generic variable names (X1,X2,X3,X4,... for example), use `ftocv`.

34.15 Time and Date Functions

<code>date</code>	Returns current system date.
<code>datestr</code>	Formats date as " mm/dd/yy ".
<code>datestring</code>	Formats date as " mm/dd/yyyy ".
<code>datestrymd</code>	Formats date as " yyyymmdd ".
<code>dayinyr</code>	Returns day number of a date.
<code>dayofweek</code>	Returns day of week.
<code>dtdate</code>	Creates a matrix in DT scalar format.
<code>dtday</code>	Creates a matrix in DT scalar format containing only the year, month, and day. Time of day information is zeroed out.
<code>dttime</code>	Creates a matrix in DT scalar format containing only the hour, minute, and second. The date information is zeroed out.
<code>dttodtv</code>	Converts DT scalar format to DTV vector format.
<code>dttostr</code>	Converts a matrix containing dates in DT scalar format to a string array.
<code>dttoutc</code>	Converts DT scalar format to UTC scalar format.
<code>dtvnormal</code>	Normalizes a date and time (DTV) vector.
<code>dtvtodt</code>	Converts DTV vector format to DT scalar format.

dtvtoutc	Converts DTV vector format to UTC scalar format.
etdays	Difference between two times in days.
ethsec	Difference between two times in hundredths of a second.
etstr	Converts elapsed time to string.
hsec	Returns elapsed time since midnight in hundredths of a second.
strtodt	Converts a string array of dates to a matrix in DT scalar format.
time	Returns current system time.
timedt	Returns system date and time in DT scalar format.
timestr	Formats time as " hh:mm:ss ".
timeutc	Returns the number of seconds since January 1, 1970 Greenwich Mean Time.
todaydt	Returns system date in DT scalar format. The time returned is always midnight (00:00:00), the beginning of the returned day.
utctodt	Converts UTC scalar format to DT scalar format.
utctodtv	Converts UTC scalar format to DTV vector format.

Use **hsec** to time segments of code. For example,

```
et = hsec;
x = y*y;
et = hsec - et;
```

will time the **GAUSS** multiplication operator.

34.16 Console I/O

con	Requests console input, creates matrix.
cons	Requests console input, creates string.
key	Gets the next key from the keyboard buffer. If buffer is empty, returns a 0.
keyav	Checks if keystroke is available.
keyw	Gets the next key from the keyboard buffer. If buffer is empty, waits for a key.
wait	Waits for a keystroke.
waitc	Flushes buffer, then waits for a keystroke.

key can be used to trap most keystrokes. For example, the following loop will trap the ALT-H key combination:

```
kk = 0;
do until kk == 1035;
    kk = key;
endo;
```

Other key combinations, function keys and cursor key movement can also be trapped. See **key**.

cons and **con** can be used to request information from the console. **keyw**, **wait**, and **waitc** will wait for a keystroke.

34.17 Output Functions

34.17.1 Text Output

<code>cls</code>	Clears the window.
<code>comlog</code>	Controls interactive command logging.
<code>csrcol</code>	Gets column position of cursor on window.
<code>csrlin</code>	Gets row position of cursor on window.
<code>ed</code>	Accesses an alternate editor.
<code>edit</code>	Edits a file with the GAUSS editor.
<code>format</code>	Defines format of matrix printing.
<code>formatcv</code>	Sets the character data format used by printfmt .
<code>formatnv</code>	Sets the numeric data format used by printfmt .
<code>header</code>	Prints a header for a report.
<code>headermt</code>	Prints a header for a report.
<code>locate</code>	Positions the cursor on the window.
<code>output</code>	Redirects <code>print</code> statements to auxiliary output.
<code>outwidth</code>	Sets line width of auxiliary output.
<code>print</code>	Prints to window.
<code>printdos</code>	Prints a string for special handling by the OS.

printfm	Prints matrices using a different format for each column.
printfmt	Prints character, numeric, or mixed matrix using a default format controlled by the functions formatcv and formatnv .
satostrC	Copies from one string array to another using a C language format specifier string for each element.
screen [on off]	Directs/suppresses print statements to window.
tab	Positions the cursor on the current line.

The results of all printing can be sent to an output file using **output**. This file can then be printed or ported as an ASCII file to other software.

To produce boxes, etc. using characters from the extended ASCII set, use **chrs**.

34.17.2 DOS Compatibility Windows

doswin	Opens the DOS compatibility window with default settings.
DOSWinCloseall	Closes the DOS compatibility window.
DOSWinOpen	Opens the DOS compatibility window and gives it the specified title and attributes.

34.18 GAUSS Graphics

This section summarizes all procedures available within the **GAUSS** graphics system. A general usage description will be found in **GAUSS GRAPHICS, CHAPTER 1**.

34.18.1 Graph Types

Commands by
Category

plotArea	Creates a cumulative area plot.
plotBar	Creates a bar plot.
plotBox	Creates a box plot.
plotCDFEmpirical	Creates a cumulative distribution function (cdf) of the empirical distribution plot.
plotContour	Creates a contour plot.
plotHist	Calculates and creates a frequency histogram plot.
plotHistF	Creates a histogram plot from a vector of frequencies.
plotHistP	Calculates and creates a percentage frequency histogram plot.
plotLogLog	Creates a 2-dimensional line plot with logarithmic scaling of the both the X and Y axes.
plotLogX	Creates a 2-dimensional line plot with logarithmic scaling of the X axis.
plotLogY	Creates a 2-dimensional line plot with logarithmic scaling of the Y axis.
plotPolar	Creates a polar plot.
plotScatter	Creates a 2-dimensional scatter plot.
plotSurface	Creates a 3-dimensional surface plot.

plotTS	Creates a graph of time series data.
plotXY	Creates a 2-dimensional line plot.

34.18.2 Adding Data to Existing Graphs

plotAddArea	Adds a cumulative area plot to an existing 2-D graph.
plotAddBar	Adds a bar or a set of bars to an existing 2-D graph.
plotAddErrorBar	Adds an error bar or a set of bars to an existing 2-D graph.
plotAddBox	Adds a box plot to an existing 2-D graph.
plotAddHist	Adds a histogram to an existing 2-D graph.
plotAddHistF	Adds a frequency histogram to an existing 2-D graph.
plotAddHistP	Adds a percent frequency histogram to an existing 2-D graph.
plotAddPolar	Adds a graph using polar coordinates to an existing polar graph.
plotAddScatter	Adds a set of points to an existing 2-D graph.
plotAddSurface	Adds a surface plot to an existing 3-D plot.
plotAddTS	Adds a curve of time series data to an existing time series plot.
plotAddXY	Adds an XY plot to an existing 2-D graph.

34.18.3 Plot Control

plotClearLayout	Clears any previously set plot layouts.
plotCustomLayout	Plots a graph of user-specified size at a user-specified location.
plotGetDefaults	Gets default settings for graph types.
plotLayout	Divides a plot into a grid of subplots and assigns the cell location in which to draw the next created graph.
plotOpenWindow	Opens a new, empty graph window to be used by the next drawn graph.
plotSave	Saves the last created graph to a user specified file type.
plotSetAxesPen	Sets the color for the axes line.
plotSetBar	Sets the fill style and format of bars in a histogram or bar graph.
plotSetBkdColor	Sets background color of a graph.
plotSetColorMap	Sets the color maps for a surface or contour plot.
plotSetContourLabels	Sets the contour labels for a contour plot.
plotSetFill	Settings for the background grid of a plot.
plotSetGrid	Controls the settings for the background grid of a plot.
plotSetLegend	Adds a legend to a graph.
plotSetLegendFont	Set the legend font for a graph.
plotSetLineColor	Sets line colors for a graph.

plotSetLineStyle	Sets line styles for a graph.
plotSetLineSymbol	Sets line symbols displayed on the plotted points of a graph.
plotSetLineThickness	Sets line thickness for a graph.
plotSetNewWindow	Sets whether or not graph should be drawn in the same window or a new window.
plotSetTextInterpreter	Controls the text interpreter settings for a graph.
plotSetTitle	Controls the settings for the title for a graph.
plotSetWhichYAxis	Assigns curves to the right or left Y-axis.
plotSetXLabel	Controls the settings for the X-axis label on a graph.
plotSetXRange	Sets the range for the X-axis.
plotSetXTicCount	Controls the number of major tics on the X-axis of a 2-D plot.
plotSetXTicInterval	Controls the interval between X-axis tic labels and also allows the user to specify the first tic to be labeled for 2-D time series graphs.
plotSetXTicLabel	Controls the formatting and angle of X-axis tic labels for 2-D time series graphs.
plotSetYLabel	Controls the settings for the Y-axis label on a graph.
plotSetYRange	Sets the range for the y-axis.
plotSetYTicCount	Controls the number of major tics on

plotSetZLabel

the y-axis of a 2-D plot.

Controls the settings for the Z-axis label on a graph.

plotSetZLevels

Controls the heights at which lines are drawn on a contour plot.

34.18.4 Annotation Control

annotationGetDefaults

Fills in an instance of a plotAnnotation structure with default values.

annotationSetBkd

Sets the background color and transparency level for a textbox, rectangle or ellipse.

annotationSetFont

Sets the font properties of a plotAnnotation structure for controlling text boxes added to a graph.

annotationSetLineColor

Sets the line color for textbox, rectangle or ellipse borders as well as the color for lines and arrows.

annotationSetLineStyle

Sets the line style for textbox, rectangle or ellipse borders as well as the style for lines and arrows.

annotationSetLineThickness

Sets the line thickness for textbox, rectangle or

ellipse borders as well as the color for lines and arrows.

34.19 PQG Graphics

This section summarizes all procedures and global variables available within the PUBLICATION QUALITY GRAPHICS (PQG) System. A general usage description will be found in **PQG GRAPHICS COLORS**. Note that PUBLICATION QUALITY GRAPHICS (PQG) graphic functions are included as legacy code and have been replaced with new plot functions.

34.19.1 Graph Types

bar	Generates bar graph.
box	Graphs data using the box graph percentile method.
contour	Graphs contour data.
draw	Supplies additional graphic elements to graphs.
hist	Computes and graphs frequency histogram.
histf	Graphs a histogram given a vector of frequency counts.
histp	Graphs a percent frequency histogram of a vector.
loglog	Graphs X,Y using logarithmic X and Y axes.
logx	Graphs X,Y using logarithmic X axis.
logy	Graphs X,Y using logarithmic Y axis.
surface	Graphs a 3-D surface.

xy	Graphs X,Y using Cartesian coordinate system.
xyz	Graphs X,Y,Z using 3-D Cartesian coordinate system.

34.19.2 Axes Control and Scaling

<i>_paxes</i>	Turns axes on or off.
<i>_pcross</i>	Controls where axes intersect.
<i>_pgrid</i>	Controls major and minor grid lines.
<i>_pticout</i>	Controls direction of tick marks on axes.
<i>_pxpmax</i>	Controls precision of numbers on X axis.
<i>_pxsci</i>	Controls use of scientific notation on X axis.
<i>_pypmax</i>	Controls precision of numbers on Y axis.
<i>_pysci</i>	Controls use of scientific notation on Y axis.
<i>_pzpmax</i>	Controls precision of numbers on Z axis.
<i>_pzsci</i>	Controls use of scientific notation on Z axis.
scale	Scales X,Y axes for 2-D plots.
scale3d	Scales X,Y, and Z axes for 3-D plots.
xtics	Scales X axis and controls tick marks.
ytics	Scales Y axis and controls tick marks.
ztics	Scales Z axis and controls tick marks.

34.19.3 Text, Labels, Titles, and Fonts

<i>_paxht</i>	Controls size of axes labels.
<i>_pdate</i>	Controls date string contents.
<i>_plegctl</i>	Sets location and size of plot legend.
<i>_plegstr</i>	Specifies legend text entries.
<i>_pmsgctl</i>	Controls message position.
<i>_pmsgstr</i>	Specifies message text.
<i>_pnum</i>	Controls axes numeric labels and orientation.
<i>_pnumht</i>	Controls size of axes numeric labels.
<i>_ptitlht</i>	Controls main title size.
asclabel	Defines character labels for tick marks.
fonts	Loads fonts for labels, titles, messages, and legend.
title	Specifies main title for graph.
xlabel	Specifies X axis label.
ylabel	Specifies Y axis label.
zlabel	Specifies Z axis label.

34.19.4 Main Curve Lines and Symbols

<i>_pboxctl</i>	Controls box plotter.
<i>_pboxlim</i>	Outputs percentile matrix from box plotter.
<i>_pcolor</i>	Controls line color for main curves.
<i>_plctrl</i>	Controls main curve and frequency of data symbols.

Commands by Category

<code>_plttype</code>	Controls line style for main curves.
<code>_plwidth</code>	Controls line thickness for main curves.
<code>_pstype</code>	Controls symbol type for main curves.
<code>_psymsiz</code>	Controls symbol size for main curves.
<code>_pzclr</code>	Z level color control for contour and surface .

34.19.5 Extra Lines and Symbols

<code>_parrow</code>	Creates arrows.
<code>_parrow3</code>	Creates arrows for 3-D graphs.
<code>_perrbar</code>	Plots error bars.
<code>_pline</code>	Plots extra lines and circles.
<code>_pline3d</code>	Plots extra lines for 3-D graphs.
<code>_psym</code>	Plots extra symbols.
<code>_psym3d</code>	Plots extra symbols for 3-D graphs.

34.19.6 Graphic Panel, Page, and Plot Control

<code>_pageshf</code>	Shifts the graph for printer output.
<code>_pagesiz</code>	Controls size of graph for printer output.
<code>_plotshf</code>	Controls plot area position.
<code>_plotsiz</code>	Controls plot area size.
<code>_protate</code>	Rotates the graph 90 degrees.
axmargin	Controls axes margins and plot size.
begwind	Graphic panel initialization procedure.
endwind	Ends graphic panel manipulation; displays graphs.

getwind	Gets current graphic panel number.
loadwind	Loads a graphic panel configuration from a file.
makewind	Creates graphic panel with specified size and position.
margin	Controls graph margins.
nextwind	Sets to next available graphic panel number.
savewind	Saves graphic panel configuration to a file.
setwind	Sets to specified graphic panel number.
window	Creates tiled graphic panels of equal size.

axmargin is preferred to the older *_plotsiz* and *_plotshf* globals for establishing an absolute plot size and position.

34.19.7 Output Options

<i>_pscreen</i>	Controls graphics output to window.
<i>_psilent</i>	Controls final beep.
<i>_ptek</i>	Controls creation and name of <code>graphics.tkf</code> file.
<i>_pzoom</i>	Specifies zoom parameters.
graphprt	Generates print, conversion file.
pqgwin	Sets the graphics viewer mode.
setvwrmode	Sets the graphics viewer mode.
tkf2eps	Converts <code>.tkf</code> file to Encapsulated PostScript file.

tkf2ps

Converts `.tkf` file to PostScript file.

34.19.8 Miscellaneous

Commands by
Category

_pbox

Draws a border around graphic panel/window.

_pcrop

Controls cropping of graphics data outside axes area.

_pframe

Draws a frame around 2-D, 3-D plots.

_pmcolor

Controls colors to be used for axes, title, x and y labels, date, box, and background.

graphset

Resets all PQG globals to default values.

rerun

Displays most recently created graph.

view

Sets 3-D observer position in workbox units.

viewxyz

Sets 3-D observer position in plot coordinates.

volume

Sets length, width, and height ratios of 3-D workbox.

35 Command Reference

a	35-52
abs	35-52
acf	35-53
aconcat	35-58
aeye	35-61
amax	35-63
amean	35-65
AmericanBinomCall	35-67
AmericanBinomCall_Greeks	35-69
AmericanBinomCall_ImpVol	35-71
AmericanBinomPut	35-73
AmericanBinomPut_Greeks	35-75
AmericanBinomPut_ImpVol	35-77
AmericanBSCall	35-79
AmericanBSCall_Greeks	35-80

AmericanBSCall_ImpVol	35-82
AmericanBSPut	35-84
AmericanBSPut_Greeks	35-85
AmericanBSPut_ImpVol	35-87
amin	35-89
amult	35-92
annotationGetDefaults	35-94
annotationSetBkd	35-95
annotationSetFont	35-97
annotationSetLineColor	35-99
annotationSetLineStyle	35-101
annotationSetLineThickness	35-103
annualTradingDays	35-104
arccos	35-106
arcsin	35-107
areshape	35-108
arrayalloc	35-110
arrayindex	35-112
arrayinit	35-114
arraytomat	35-115
asciiload	35-116

asclabel	35-118
astd	35-120
astds	35-122
asum	35-124
atan	35-126
atan2	35-128
atranspose	35-130
axmargin	35-132
b	35-135
balance	35-135
band	35-136
bandchol	35-138
bandcholsol	35-140
bandltsol	35-141
bandrv	35-143
bandsolpd	35-145
bar	35-146
base10	35-148
begwind	35-149
besselj	35-150
bessely	35-151

beta	35-153
box	35-154
boxcox	35-156
break	35-157
c	35-159
call	35-159
cdfBeta	35-160
cdfBetaInv	35-162
cdfBinomial	35-164
cdfBinomialInv	35-165
cdfBvn	35-167
cdfBvn2	35-169
cdfBvn2e	35-171
cdfCauchy	35-173
cdfCauchyInv	35-174
cdfChic	35-175
cdfChii	35-177
cdfChinc	35-179
cdfChincInv	35-180
cdfEmpirical	35-182
cdfExp	35-185

cdfExpInv	35-186
cdfFc	35-186
cdfFnc	35-189
cdfFncInv	35-190
cdfGam	35-192
cdfGenPareto	35-194
cdfHyperGeo	35-196
cdfLaplace	35-198
cdfLaplaceInv	35-199
cdfLogistic	35-199
cdfLogisticInv	35-201
cdfMvn	35-201
cdfMvnce	35-205
cdfMvne	35-210
cdfMvn2e	35-215
cdfMvtce	35-221
cdfMvte	35-227
cdfMvt2e	35-233
cdfN, cdfNc	35-239
cdfNegBinomial	35-243
cdfNegBinomialInv	35-245

cdfN2	35-246
cdfNi	35-248
cdfPoisson	35-250
cdfPoissonInv	35-251
cdfRayleigh	35-252
cdfRayleighInv	35-254
cdfTc	35-255
cdfTci	35-257
cdfTnc	35-258
cdfTvn	35-261
cdfWeibull	35-263
cdfWeibullInv	35-265
cdir	35-266
ceil	35-267
changeDir	35-269
chdir	35-269
chiBarSquare	35-270
chol	35-272
choldn	35-274
cholsol	35-276
cholup	35-278

chrs	35-279
clear	35-281
clearg	35-282
close	35-283
closeall	35-286
cls	35-289
code	35-290
code (dataloop)	35-293
cols	35-295
colsf	35-296
combinate	35-297
combined	35-299
comlog	35-302
compile	35-303
complex	35-305
con	35-306
cond	35-309
conj	35-310
cons	35-311
ConScore	35-312
continue	35-316

contour	35-318
conv	35-319
convertsatostr	35-320
convertstrtosa	35-321
corrmm,corrvc,corrxx	35-322
corrms,corrxs	35-324
cos	35-327
cosh	35-328
counts	35-329
countwts	35-331
create	35-333
crossprd	35-340
crout	35-341
croutp	35-343
csrcol,csrlin	35-345
csvReadM	35-346
csvReadSA	35-352
csvWriteM	35-356
cumprodc	35-360
cumsumc	35-362
curve	35-363

cvtos	35-365
cvtos	35-366
d	35-368
datacreate	35-368
datacreatecomplex	35-370
datalist	35-373
dataload	35-374
dataloop (dataloop)	35-376
dataopen	35-377
datasave	35-381
date	35-382
datestr	35-383
datestring	35-384
datestrymd	35-385
dayinyr	35-386
dayofweek	35-387
dbAddDatabase	35-389
dbClose	35-391
dbCommit	35-391
dbCreateQuery	35-392
dbExecQuery	35-394

dbGetConnectOptions	35-395
dbGetDatabaseName	35-396
dbGetDriverName	35-396
dbGetDrivers();	35-397
dbGetHostName	35-398
dbGetLastErrorNum	35-399
dbGetLastErrorText	35-399
dbGetNumericalPrecPolicy	35-400
dbGetPassword	35-401
dbGetPort	35-401
dbGetPrimaryIndex	35-402
dbGetTableHeaders	35-403
dbGetTables	35-404
dbGetUserName	35-404
dbHasFeature	35-405
dbIsDriverAvailable	35-407
dbIsOpen	35-408
dbIsOpenError	35-408
dbIsValid	35-409
dbNumericalPrecPolicy	35-410
dbOpen	35-410

dbQueryBindValue	35-412
dbQueryClear	35-413
dbQueryCols	35-413
dbQueryExecPrepared	35-414
dbQueryFetchAllM	35-415
dbQueryFetchAllSA	35-417
dbQueryFetchOneM	35-418
dbQueryFetchOneSA	35-420
dbQueryFinish	35-421
dbQueryGetBoundValue	35-422
dbQueryGetBoundValues	35-423
dbQueryGetField	35-424
dbQueryGetLastErrorNum	35-426
dbQueryGetLastErrorText	35-426
dbQueryGetLastInsertID	35-427
dbQueryGetLastQuery	35-428
dbQueryGetNumRowsAffected	35-429
dbQueryGetPosition	35-430
dbQueryIsActive	35-431
dbQueryIsForwardOnly	35-432
dbQueryIsNull	35-433

dbQueryIsSelect	35-434
dbQueryIsValid	35-435
dbQueryPrepare	35-436
dbQueryRows	35-438
dbQuerySeek	35-439
dbQuerySeekFirst	35-440
dbQuerySeekLast	35-442
dbQuerySeekNext	35-443
dbQuerySeekPrevious	35-445
dbQuerySetForwardOnly	35-446
dbQuerySetNumericalPrecisionPolicy	35-447
dbRemoveDatabase	35-448
dbRollback	35-448
dbSetConnectOptions	35-449
dbSetDatabaseName	35-452
dbSetHostName	35-453
dbSetPassword	35-454
dbSetPort	35-455
dbSetUserName	35-456
dbTransaction	35-456
debug	35-458

declare	35-458
delete	35-464
delete (dataloop)	35-466
deleteFile	35-467
delif	35-468
delrows	35-473
denseToSp	35-475
denseToSpRE	35-476
denToZero	35-478
design	35-480
det	35-482
detl	35-484
dfft	35-485
dffti	35-486
diag	35-487
diagrv	35-490
digamma	35-492
dlibrary	35-492
dllcall	35-495
do while,do until	35-497
dos	35-500

doswin	35-503
DOSWinCloseall	35-503
DOSWinOpen	35-504
dot	35-507
dotfeq,dotfge, dotfgt,dotfle,dotflt,dotfne	35-509
dotfeqmt,dotfgemt,dotfgtmt,dotflemt,dotfltmt,dotfnemt	35-511
draw	35-513
drop (dataloop)	35-515
dsCreate	35-515
dstat	35-516
dstatmt	35-523
dstatmtControlCreate	35-531
dtdate	35-532
dtday	35-534
dttime	35-535
dttodtv	35-536
dttostr	35-537
dttoutc	35-540
dtvnormal	35-542
dtvtodt	35-543
dtvtoutc	35-545

dummy	35-546
dummybr	35-548
dummydn	35-550
e	35-553
ed	35-553
edit	35-554
erfInv,erfCInv	35-555
eig	35-556
eigh	35-558
eighv	35-560
eigv	35-561
elapsedTradingDays	35-563
end	35-565
endp	35-566
endwind	35-567
envget	35-568
eof	35-569
eqSolve	35-571
eqSolvemt	35-576
eqSolvemtControlCreate	35-584
eqSolvemtOutCreate	35-585

eqSolveSet	35-587
erf,erfc	35-587
erfcplx,erfccplx	35-589
error	35-590
errorlog	35-593
errorlogat	35-593
etdays	35-594
ethsec	35-595
etstr	35-597
EuropeanBinomCall	35-598
EuropeanBinomCall_Greeks	35-600
EuropeanBinomCall_ImpVol	35-602
EuropeanBinomPut	35-604
EuropeanBinomPut_Greeks	35-605
EuropeanBinomPut_ImpVol	35-608
EuropeanBSCall	35-609
EuropeanBSCall_Greeks	35-611
EuropeanBSCall_ImpVol	35-613
EuropeanBSPut	35-614
EuropeanBSPut_Greeks	35-616
EuropeanBSPut_ImpVol	35-618

exctsmpl	35-619
exec	35-621
execbg	35-622
exp	35-623
extern (dataloop)	35-625
external	35-626
eye	35-627
f	35-629
fcheckerr	35-629
fclearerr	35-630
feq,fge,fgt,fle,flt,fne	35-631
feqmt,fgemt,fgtmt,flemt,fltmt,fnemt	35-633
fflush	35-635
fft	35-635
ffti	35-636
fftm	35-637
fftmi	35-640
fftn	35-643
fgets	35-645
fgetsa	35-646
fgetsat	35-647

fgetst	35-648
fileinfo	35-649
filesa	35-651
floor	35-652
fmod	35-654
fn	35-656
fonts	35-657
fopen	35-658
for	35-660
format	35-663
formatcv	35-673
formatnv	35-674
fputs	35-675
fputst	35-678
fseek	35-681
fstrerror	35-683
ftell	35-684
ftocv	35-685
ftos	35-686
ftostrC	35-690
g	35-693

gamma	35-693
gammacplx	35-694
gammaii	35-695
gausset	35-696
gdaAppend	35-697
gdaCreate	35-699
gdaDStat	35-700
gdaDStatMat	35-704
gdaGetIndex	35-709
gdaGetName	35-710
gdaGetNames	35-711
gdaGetOrders	35-712
gdaGetType	35-714
gdaGetTypes	35-715
gdaGetVarInfo	35-717
gdaIsCplx	35-719
gdaLoad	35-720
gdaPack	35-724
gdaRead	35-726
gdaReadByIndex	35-727
gdaReadSome	35-728

gdaReadSparse	35-731
gdaReadStruct	35-732
gdaReportVarInfo	35-734
gdaSave	35-735
gdaUpdate	35-738
gdaUpdateAndPack	35-740
gdaVars	35-742
gdaWrite	35-743
gdaWrite32	35-744
gdaWriteSome	35-746
getarray	35-749
getdims	35-750
getf	35-751
getGAUSSHome	35-754
getmatrix	35-755
getmatrix4D	35-757
getname	35-759
getnamef	35-760
getNextTradingDay	35-762
getNextWeekDay	35-763
getnr	35-764

getnrmt	35-765
getorders	35-766
getpath	35-767
getPreviousTradingDay	35-769
getPreviousWeekDay	35-770
getRow	35-770
getscalar3D	35-772
getscalar4D	35-773
getTrRow	35-775
getwind	35-776
glm	35-777
gosub	35-807
goto	35-810
gradMT	35-811
gradMTm	35-813
gradMTT	35-815
gradMTTm	35-817
gradp, gradcplx	35-818
graphprt	35-820
graphset	35-824
h	35-825

h5create	35-825
h5open	35-828
h5read	35-831
h5readAttribute	35-838
h5write	35-840
h5writeAttribute	35-842
hasimag	35-845
header	35-846
headermt	35-847
hess	35-849
hessMT	35-850
hessMTg	35-852
hessMTgw	35-854
hessMTm	35-855
hessMTmw	35-857
hessMTT	35-859
hessMTTg	35-861
hessMTTgw	35-863
hessMTTm	35-864
hessMTw	35-866
hessp, hesscplx	35-868

hist	35-870
histf	35-872
histp	35-873
hsec	35-874
i	35-876
if, else, elseif	35-876
imag	35-877
#include	35-878
indcv	35-880
indexcat	35-882
indices	35-883
indices2	35-885
indicesf	35-886
indicesfn	35-888
indnv	35-889
indsav	35-891
integrate1d	35-892
intgrat2	35-897
intgrat3	35-899
inthp1	35-902
inthp2	35-906

inthp3	35-910
inthp4	35-914
inthpControlCreate	35-918
intquad1	35-919
intquad2	35-922
intquad3	35-926
intrleav	35-930
intrleavsa	35-932
intrsect	35-933
intrsectsa	35-936
intsimp	35-938
inv, invpd	35-939
invswp	35-941
iscplx	35-942
iscplxsf	35-943
isden	35-944
isinfnanmiss	35-945
ismiss	35-946
itos	35-948
k	35-950
keep (dataloop)	35-950

key	35-951
keyav	35-952
keyw	35-953
keyword	35-953
l	35-956
lag (dataloop)	35-956
lag1	35-957
lagn	35-958
lapeighb	35-960
lapeighi	35-962
lapeighvb	35-964
lapeighvi	35-966
lapgeig	35-968
lapgeigh	35-969
lapgeighv	35-971
lapgeigv	35-973
lapgsvdst	35-974
lapgsvds	35-977
lapgsvdst	35-979
lapgschur	35-982
lapsvdcusv	35-987

lapsvds	35-989
lapsvdusv	35-990
ldl	35-992
ldlp	35-994
ldlsol	35-996
let	35-997
lib	35-1001
library	35-1005
#lineson, #linesoff	35-1009
linsolve	35-1010
listwise (dataloop)	35-1012
ln	35-1012
lncdfbvn	35-1013
lncdfbvn2	35-1014
lncdfmvn	35-1016
lncdfn	35-1017
lncdfn2	35-1018
lncdfnc	35-1020
lnfact	35-1021
lngammacplx	35-1023
lnpdfmvn	35-1024

lnpdfmvt	35-1025
lnpdfn	35-1026
lnpdft	35-1027
load, loadf, loadk, loadm, loadp, loads	35-1028
loadarray	35-1033
loadd	35-1035
loadstruct	35-1039
loadwind	35-1040
local	35-1041
locate	35-1041
loess	35-1042
loessmt	35-1044
loessmtControlCreate	35-1046
log	35-1047
loglog	35-1048
logx	35-1049
logy	35-1050
loopnextindex	35-1051
lower	35-1054
lowmat, lowmat1	35-1056
ltrisol	35-1057

lu	35-1058
lusol	35-1060
m	35-1061
machEpsilon	35-1061
make (dataloop)	35-1061
makevars	35-1062
makewind	35-1064
margin	35-1066
matalloc	35-1067
matinit	35-1068
mattoarray	35-1069
maxc	35-1070
maxindc	35-1071
maxv	35-1073
maxvec	35-1075
maxbytes	35-1076
mbesseli	35-1077
meanc	35-1080
median	35-1081
mergeby	35-1083
mergevar	35-1084

minc	35-1086
minindc	35-1087
minv	35-1089
miss, missrv	35-1090
missex	35-1093
moment	35-1095
momentd	35-1097
movingave	35-1101
movingaveExpwgt	35-1102
movingaveWgt	35-1103
msym	35-1104
n	35-1107
new	35-1107
nextindex	35-1108
nextn, nextnevn	35-1110
nextwind	35-1112
ntos	35-1112
null	35-1115
null1	35-1117
numCombinations	35-1118
o	35-1120

ols	35-1120
olsmt	35-1129
olsmtControlCreate	35-1140
olsqr	35-1141
olsqr2	35-1142
olsqrmt	35-1143
ones	35-1145
open	35-1146
optn, optnevn	35-1152
orth	35-1154
output	35-1156
outtyp (dataloop)	35-1159
outwidth	35-1160
p	35-1162
pacf	35-1162
packedToSp	35-1171
packr	35-1173
parse	35-1175
pause	35-1179
pdfBinomial	35-1179
pdfCauchy	35-1181

pdfexp	35-1182
pdfGenPareto	35-1183
pdfHyperGeo	35-1185
pdfLaplace	35-1187
pdflogistic	35-1188
pdfn	35-1189
pdfPoisson	35-1190
pdfRayleigh	35-1192
pdfWeibull	35-1193
pdfWishartInv	35-1194
pi	35-1196
pinv	35-1197
pinvmt	35-1199
plotAddArea	35-1200
plotAddArrow	35-1204
plotAddBar	35-1206
plotAddBox	35-1207
plotAddErrorBar	35-1208
plotAddHist	35-1216
plotAddHistF	35-1217
plotAddHistP	35-1218

plotAddPolar	35-1219
plotAddScatter	35-1220
plotAddShape	35-1221
plotAddSurface	35-1223
plotAddTextbox	35-1228
plotAddTS	35-1231
plotAddXY	35-1232
plotArea	35-1233
plotBar	35-1234
plotBox	35-1236
plotCDFEmpirical	35-1238
plotClearLayout	35-1241
plotContour	35-1243
plotCustomLayout	35-1246
plotGetDefaults	35-1247
plotHist	35-1249
plotHistF	35-1250
plotHistP	35-1250
plotLayout	35-1251
plotLogLog	35-1253
plotLogX	35-1253

plotLogY	35-1254
plotOpenWindow	35-1255
plotPolar	35-1256
plotSave	35-1257
plotScatter	35-1260
plotSetAxesPen	35-1261
plotSetBar	35-1262
plotSetBkdColor	35-1265
plotSetColorMap	35-1267
plotSetContourLabels	35-1270
plotSetFill	35-1273
plotSetGrid	35-1275
plotSetLegend	35-1276
plotSetLegendFont	35-1278
plotSetLineColor	35-1280
plotSetLineStyle	35-1281
plotSetLineSymbol	35-1282
plotSetLineThickness	35-1285
plotSetNewWindow	35-1286
plotSetTextInterpreter	35-1287
plotSetTitle	35-1292

plotSetWhichY Axis	35-1295
plotSetXLabel	35-1296
plotSetXRange	35-1299
plotSetXTicCount	35-1300
plotSetXTicInterval	35-1303
plotSetXTicLabel	35-1306
plotSetYLabel	35-1311
plotSetYRange	35-1314
plotSetYTicCount	35-1315
plotSetZLabel	35-1319
plotSetZLevels	35-1321
plotSurface	35-1325
plotTS	35-1327
plotXY	35-1331
polar	35-1332
polychar	35-1333
polyeval	35-1334
polygamma	35-1336
polyint	35-1339
polymake	35-1340
polymat	35-1342

polymroot	35-1343
polymult	35-1345
polyroot	35-1347
pop	35-1348
powerM	35-1349
pqgwin	35-1351
previousindex	35-1352
princomp	35-1353
print	35-1355
printdos	35-1366
printfm	35-1367
printfmt	35-1370
proc	35-1373
prodc	35-1375
psi	35-1376
putarray	35-1377
putf	35-1379
putvals	35-1380
pvCreate	35-1382
pvGetIndex	35-1383
pvGetParNames	35-1383

pvGetParVector	35-1385
pvLength	35-1387
pvList	35-1388
pvPack	35-1389
pvPacki	35-1390
pvPackm	35-1392
pvPackmi	35-1394
pvPacks	35-1396
pvPacksi	35-1398
pvPacksm	35-1400
pvPacksmi	35-1402
pvPutParVector	35-1405
pvTest	35-1407
pvUnpack	35-1407
q	35-1409
QNewton	35-1409
QNewtonmt	35-1412
QNewtonmtControlCreate	35-1418
QNewtonmtOutCreate	35-1419
QNewtonSet	35-1420
QProg	35-1420

QProgmt	35-1422
QProgmtInCreate	35-1425
qqr	35-1426
qqre	35-1428
qqrep	35-1431
qr	35-1433
qre	35-1435
qrep	35-1438
qrsol	35-1440
qrtsol	35-1441
qtyr	35-1443
qtyre	35-1446
qtyrep	35-1449
quantile	35-1451
quantiled	35-1454
qyr	35-1458
qyre	35-1460
qyrep	35-1462
qz	35-1464
r	35-1470
rank	35-1470

rankindx	35-1471
readr	35-1472
real	35-1474
reclassify	35-1475
reclassifyCuts	35-1478
recode	35-1483
recode (dataloop)	35-1486
recserar	35-1488
recsercp	35-1490
recserrc	35-1492
renamefile	35-1494
rerun	35-1495
rescale	35-1495
reshape	35-1501
retp	35-1503
return	35-1503
rev	35-1504
rfft	35-1506
rffti	35-1507
rfftip	35-1507
rfftn	35-1509

rfftnp	35-1511
rfftp	35-1513
rndBernoulli	35-1514
rndBeta	35-1516
rndCauchy	35-1519
rndChiSquare	35-1520
rndcon, rndmult, rndseed	35-1522
rndCreateState	35-1524
rndExp	35-1527
rndgam	35-1529
rndGamma	35-1530
rndGeo	35-1532
rndGumbel	35-1534
rndHyperGeo	35-1535
rndi	35-1538
rndKMbeta	35-1541
rndKMgam	35-1543
rndKMi	35-1544
rndKMn	35-1546
rndKMnb	35-1548
rndKMp	35-1550

rndKMu	35-1551
rndKMvm	35-1553
rndLaplace	35-1555
rndLCbeta	35-1557
rndLCgam	35-1559
rndLCi	35-1560
rndLCn	35-1563
rndLCnb	35-1565
rndLCp	35-1567
rndLCu	35-1569
rndLCvm	35-1572
rndLogNorm	35-1574
rndMVn	35-1575
rndMVt	35-1577
rndn	35-1579
rndnb	35-1581
rndNegBinomial	35-1583
rndp	35-1585
rndPoisson	35-1587
rndStateSkip	35-1588
rndu	35-1590

rndvm	35-1593
rndWeibull	35-1594
rndWishart	35-1595
rndWishartInv	35-1597
rotater	35-1598
round	35-1600
rows	35-1601
rowsf	35-1602
rref	35-1603
run	35-1605
s	35-1608
sampleData	35-1608
satostrC	35-1610
save	35-1613
saveall	35-1616
saved	35-1617
savestruct	35-1619
savewind	35-1620
scale	35-1621
scale3d	35-1622
scalerr	35-1623

scalinfnanmiss	35-1626
scalmiss	35-1627
schtoc	35-1628
schur	35-1630
screen	35-1633
searchsourcepath	35-1634
seekr	35-1635
select (dataloop)	35-1636
selif	35-1637
seqa, seqm	35-1639
setarray	35-1640
setdif	35-1642
setdifsa	35-1645
setvars	35-1646
setvwrmode	35-1647
setwind	35-1648
shell	35-1649
shiftr	35-1651
show	35-1653
sin	35-1656
singleindex	35-1657

sinh	35-1659
sleep	35-1660
solpd	35-1661
sortc, sortcc	35-1664
sortd	35-1666
sorthc, sorthec	35-1667
sortind, sortindc	35-1669
sortmc	35-1670
sortr, sortrc	35-1672
spBiconjGradSol	35-1674
spChol	35-1677
spConjGradSol	35-1678
spCreate	35-1681
spDenseSubmat	35-1682
spDiagRvMat	35-1684
spEigv	35-1687
spEye	35-1690
spGetNZE	35-1691
spline	35-1693
spLDL	35-1694
spLU	35-1695

spNumNZE	35-1697
spOnes	35-1698
spreadSheetReadM	35-1700
spreadSheetReadSA	35-1702
spreadSheetWrite	35-1704
spScale	35-1707
spSubmat	35-1709
spToDense	35-1711
spTrTDense	35-1712
spTScalar	35-1713
spZeros	35-1714
sqpSolve	35-1716
sqpSolveMT	35-1723
sqpSolveMTControlCreate	35-1731
sqpSolveMTlagrangeCreate	35-1732
sqpSolveMToutCreate	35-1733
sqpSolveSet	35-1734
sqrt	35-1735
stdc	35-1736
stdsc	35-1738
stocv	35-1739

stof	35-1740
stop	35-1741
strcombine	35-1742
strindx	35-1744
strlen	35-1745
strput	35-1746
strrindx	35-1748
strjoin	35-1749
strsect	35-1751
strsplit	35-1753
strsplitPad	35-1757
strtodt	35-1758
strtof	35-1760
strtofcp1x	35-1762
strtrim	35-1763
strtriml	35-1765
strtrimr	35-1766
strtrunc	35-1767
strtrunc1	35-1768
strtruncpad	35-1769
strtruncr	35-1770

submat	35-1770
subscat	35-1772
substute	35-1776
subvec	35-1779
sumc	35-1781
sumr	35-1783
surface	35-1785
svd	35-1788
svd1	35-1790
svd2	35-1792
svdcusv	35-1795
svds	35-1798
svdusv	35-1801
sylvester	35-1804
sysstate	35-1806
system	35-1828
t	35-1830
tab	35-1830
tan	35-1831
tanh	35-1832
tempname	35-1833

ThreadBegin	35-1834
ThreadEnd	35-1835
threadfor, threadendfor	35-1836
ThreadJoin	35-1839
ThreadStat	35-1840
time	35-1841
timedt	35-1841
timestr	35-1842
timeutc	35-1843
title	35-1844
tkf2eps	35-1846
tkf2ps	35-1847
tocart	35-1848
todaydt	35-1848
toeplitz	35-1849
token	35-1850
topolar	35-1853
trace	35-1853
trap	35-1855
trapchk	35-1859
trigamma	35-1861

trimr	35-1862
trunc	35-1863
type	35-1865
typecv	35-1867
typef	35-1869
u	35-1871
union	35-1871
unionsa	35-1872
uniqindx	35-1873
uniqindxsa	35-1875
unique	35-1876
uniquesa	35-1882
upmat, upmat1	35-1883
upper	35-1885
use	35-1886
utctodt	35-1888
utctodtv	35-1889
utrisol	35-1891
v	35-1892
vals	35-1892
varCovM, varCovX	35-1894

varCovMS, varCovXS	35-1896
varget	35-1898
vargetl	35-1900
varmall	35-1902
varmares	35-1903
varput	35-1905
varputl	35-1906
vartypef	35-1908
vcm, vcx	35-1909
vcms, vcxs	35-1910
vec, vecr	35-1911
vech	35-1913
vector (dataloop)	35-1914
vget	35-1915
view	35-1916
viewxyz	35-1917
vlist	35-1918
vnamecv	35-1919
volume	35-1920
vput	35-1921
vread	35-1922

vtypecv	35-1923
w	35-1924
wait, waitc	35-1924
walkindex	35-1924
window	35-1926
writer	35-1927
x	35-1930
xlabel	35-1930
xlsGetSheetCount	35-1931
xlsGetSheetSize	35-1932
xlsGetSheetTypes	35-1935
xlsMakeRange	35-1937
xlsReadM	35-1939
xlsReadSA	35-1944
xlsWrite	35-1948
xlsWriteM	35-1952
xlsWriteSA	35-1955
xpnd	35-1958
xtics	35-1960
xy	35-1961
xyz	35-1962

y	35-1964
ylabel	35-1964
ytics	35-1965
z	35-1967
zeros	35-1967
zeta	35-1968
ylabel	35-1969
ztics	35-1970

abs

a

abs

Purpose

Returns the absolute value or complex modulus of x .

Format

$y = \mathbf{abs}(x);$

Input

x NxK matrix or sparse matrix or N-dimensional array.

Output

y NxK matrix or sparse matrix or N-dimensional array
containing absolute values of x .

Example

```
//Set random number generator seed for
//repeatable random numbers
randseed 929212;

x = randn(2,2);
y = abs(x);
```

The code above assigns the variables as follows:

```
x =  -0.23061709    0.054931120
      0.88863202   -0.82246522

y =   0.23061709    0.054931120
      0.88863202    0.82246522
```

In this example, a 2x2 matrix of Normal random numbers is generated and the absolute value of the matrix is computed.

acf

Purpose

Computes sample autocorrelations.

Format

```
rk = acf(y, k, d);
```

Input

<i>y</i>	Nx1 vector, data.
<i>k</i>	scalar, maximum number of autocorrelations to compute.
<i>d</i>	scalar, order of differencing.

acf

Output

rk

Kx1 vector, sample autocorrelations.

Examples

Example 1

```
//Short time-series
x = { 12.92,
      14.28 ,
      13.31 ,
      13.34 ,
      12.71 ,
      13.08 ,
      11.86 ,
      9.000 ,
      8.190 ,
      7.970 ,
      8.350 ,
      8.200 ,
      8.120 ,
      8.390 ,
      8.660 };

//Maximum number of lags
k = 4;

//Order of differencing
d = 1;

//Calculate and print result of autocorrelation function
rk = acf(x, k, d);
print rk;
```

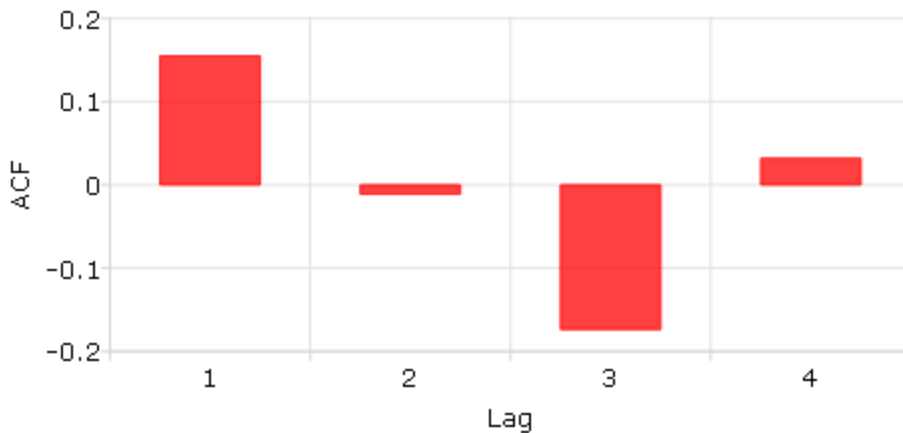

The code above produces the following output:

```
0.15488076  
-0.011078336  
-0.17367167  
0.031921209
```

Plot the ACF results with **plotBar** function. The first input *0* means using sequence numbers to name elements in the *rk*.

```
plotBar(0, rk);
```

You can type the labels for x-axis and y-axis in the graphics window. The plot is shown below:



Example 2

Calculate the autocorrelation function and plot the results for "beef_prices" data.

```
//Get file name with full path
file = getGAUSSHome() $+ "examples/beef_prices.csv";

//Import data set starting with row 2 and column 2
beef = csvReadM(file, 2, 2);

//Max lags
k = 10;

//Order of differencing
d = 0;

//Call acf function
beef_acf = acf(beef, k, d);
```

The following code can give the time plot and sample ACF plot based on the *beef* and *beef_acf*:

```
// Time series plot
//Step 1: Declare a plotControl structure
struct plotControl ctl;
ctl = plotGetDefaults("xy");

//Step 2: the time plot
//Making a 1 by 2 plot, the first plot is the time plot
plotLayout(1,2,1);

//Labels and format setting based on 'beef' matrix
```

```
plotSetYLabel(&ctl, "cents/lb");
plotSetXLabel(&ctl, "Year");
plotSetXTicLabel(&ctl, "YYYY");
plotSetXTicInterval(&ctl, 120, 199501);

//Draw time series plot
plotTS(ctl, 1992, 12, beef);

//Making a 1 by 2 plot, the second plot is the ACF plot
plotLayout(1,2,2);

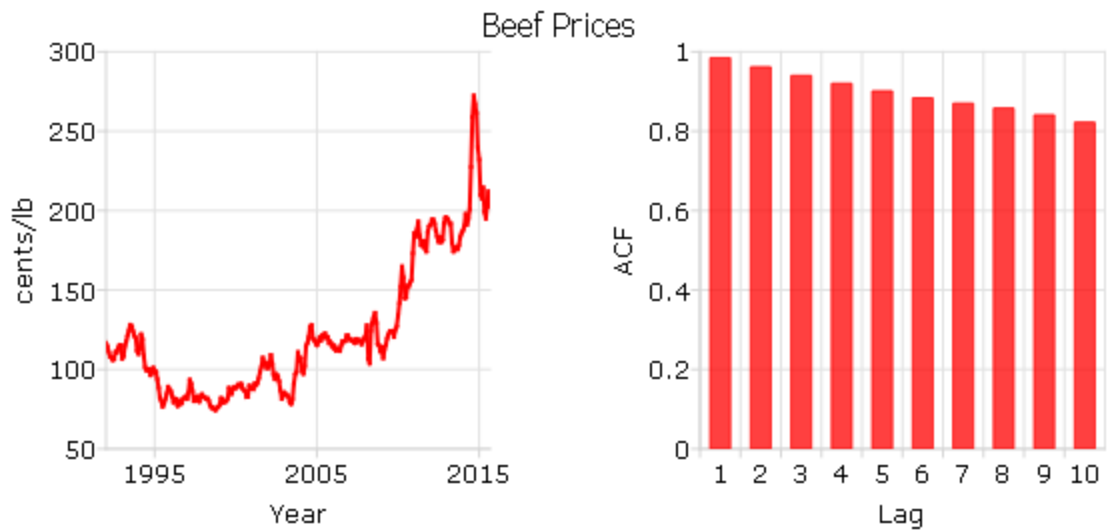
//Step 3: ACF plot
//Fill 'ctl' structure with bar plot defaults
ctl = plotGetDefaults("bar");

//Set labels and format based on 'beef_acf' matrix
plotSetYLabel(&ctl, "ACF");
plotSetXLabel(&ctl, "Lag");
plotSetXTicInterval(&ctl, 1, 5);

// ACF plot with plotBar function
plotBar(ctl, seqa(1,1,k), beef_acf);
```

You can use 'Add Text' to type 'Beef Prices' as the title in the graphics window. The plot is:

aconcat



Source

tsutil.src

See Also

[pacf](#)

aconcat

Purpose

Concatenates conformable matrices and arrays in a user-specified dimension.

Format

```
y = aconcat(a, b, dim);
```

Input

<i>a</i>	matrix or N-dimensional array.
<i>b</i>	matrix or K-dimensional array, conformable with <i>a</i> .
<i>dim</i>	scalar, dimension in which to concatenate.

Output

<i>y</i>	M-dimensional array, the result of the concatenation.
----------	---

Remarks

a and *b* are conformable only if all of their dimensions except *dim* have the same sizes. If *a* or *b* is a matrix, then the size of dimension 1 is the number of columns in the matrix, and the size of dimension 2 is the number of rows in the matrix.

Example

```
//Create a 2x3x4 array with each element set to 0
a = arrayinit(2|3|4,0);

//Create a 3x4 matrix with each element set to 3
b = 3*ones(3,4);
y = aconcat(a,b,3);
```

y will be a 3x3x4 array, where [1,1,1] through [2,3,4] are zeros and [3,1,1] through [3,2,4] are threes.

aconcat

```
//Create an additive sequence from 1-20 and 'reshape' it
//into a 4x5 matrix
a = reshape(seqa(1,1,20),4,5);

b = zeros(4,5);
y = aconcat(a,b,3);
```

y will be a 2x4x5 array, where [1,1,1] through [1,4,5] are sequential integers beginning with 1, and [2,1,1] through [2,4,5] are zeros.

```
//The pipe operator '|' causes vertical concatenation so
//that the statement 2|3|4 creates a 3x1 column vector
//equal to { 2, 3, 4 }
a = arrayinit(2|3|4,0);
b = seqa(1,1,24);

//'Reshape' the vector 'b' into a 2x3x4 dimensional array
b = areshape(b,2|3|4);
y = aconcat(a,b,5);
```

y will be a 2x1x2x3x4 array, where [1,1,1,1,1] through [1,1,2,3,4] are zeros, and [2,1,1,1,1] through [2,1,2,3,4] are sequential integers beginning with 1.

```
a = arrayinit(2|3|4,0);
b = seqa(1,1,6);
b = areshape(b,2|3|1);
y = aconcat(a,b,1);
print "y = " y;
```

y will be a 2x3x5 array:

```
y =

Plane [1,...]
```

```
0.00    0.00    0.00    0.00    1.0
0.00    0.00    0.00    0.00    2.0
0.00    0.00    0.00    0.00    3.0
```

```
Plane [2,...]
```

```
0.00    0.00    0.00    0.00    4.0
0.00    0.00    0.00    0.00    5.0
0.00    0.00    0.00    0.00    6.0
```

See Also

[areshape](#)

aeye

Purpose

Creates an N-dimensional array in which the planes described by the two trailing dimensions of the array are equal to the identity.

Format

```
a = aeye(ord);
```

Input

ord Nx1 vector of orders, the sizes of the dimensions of *a*.

aeeye

Output

<code>a</code>	N-dimensional array, containing 2-dimensional identity arrays.
----------------	--

Remarks

If `ord` contains numbers that are not integers, they will be truncated to integers.

The planes described by the two trailing dimensions of `a` will contain 1's down the diagonal and 0's everywhere else.

Example

`a`

```
v = { 2, 3, 3 };  
a = aeeye (v);
```

`a` will be a 2x3x3 array, such that:

[1,1,1] through [1,3,3] =

```
1 0 0  
0 1 0  
0 0 1
```

[2,1,1] through [2,3,3] =

```
1 0 0  
0 1 0  
0 0 1
```

See Also

[eye](#)

amax

Purpose

Moves across one dimension of an N-dimensional array and finds the largest element.

Format

```
y = amax(x, dim);
```

Input

<i>x</i>	N-dimensional array.
<i>dim</i>	scalar, number of dimension across which to find the maximum value.

Output

<i>y</i>	N-dimensional array.
----------	----------------------

Remarks

The output *y*, will have the same sizes of dimensions as *x*, except that the dimension indicated by *dim* will be collapsed to 1.

Example

```
randseed 9823432;

//Create random normal numbers with a standard deviation
//of 10 and round them to the nearest integer
```

amax

```
x = round(10*randn(24,1));

//Reshape them from a 24x1 vector into 2x3x4 array
x = areshape(x,2|3|4);

// Calculate the max across the second dimension
dim = 2;
y = amax(x,dim);
```

After this calculation:

x[1,1,1] through x[1,3,4] =

-14.000000	4.0000000	6.0000000	-4.0000000
1.0000000	8.0000000	10.000000	9.0000000
-3.0000000	12.000000	5.0000000	-26.000000

x[2,1,1] through x[2,3,4] =

4.0000000	6.0000000	4.0000000	2.0000000
1.0000000	16.000000	9.0000000	-4.0000000
-4.0000000	-8.0000000	-10.000000	8.0000000

y[1,1,1] through y[1,1,4] =

1.0000000	12.000000	10.000000	9.0000000
-----------	-----------	-----------	-----------

y[2,1,1] through y[2,1,4] =

4.0000000	16.000000	9.0000000	8.0000000
-----------	-----------	-----------	-----------

Use the same x array and calculate the max across dimension 1:

```
y2 = amax(x,1);
```

After this calculation, x remains the same, but y2 is:

```
y2[1,1,1] through y2[1,3,1] =
```

```
6.0000000  
10.000000  
12.000000
```

```
y2[2,1,1] through y2[2,3,1] =
```

```
6.0000000  
16.000000  
8.0000000
```

See Also

[amin](#), [maxc](#)

amean

Purpose

Computes the mean across one dimension of an N-dimensional array.

Format

```
y = amean(x, dim);
```

Input

<i>x</i>	N-dimensional array.
<i>dim</i>	scalar, number of dimension to compute the mean across.

amean

Output

y	[N-1]-dimensional array.
-----	--------------------------

Remarks

The output y , will have the same sizes of dimensions as x , except that the dimension indicated by dim will be collapsed to 1.

Example

3

```
//Create an additive sequence from 1-24
x = seqa(1,1,24);

//'Reshape' this 24x1 vector into a 2x3x4 dimensional array
x = areshape(x,2|3|4);

y = amean(x,3);
```

x is a 2x3x4 array, such that:

[1,1,1] through [1,3,4] =

1.0000000	2.0000000	3.0000000	4.0000000
5.0000000	6.0000000	7.0000000	8.0000000
9.0000000	10.000000	11.000000	12.000000

[2,1,1] through [2,3,4] =

13.000000	14.000000	15.000000	16.000000
17.000000	18.000000	19.000000	20.000000
21.000000	22.000000	23.000000	24.000000

y will be a 1x3x4 array, such that:

[1,1,1] through [1,3,4] =

AmericanBinomCall

7.0000000	8.0000000	9.0000000	10.000000
11.000000	12.000000	13.000000	14.000000
15.000000	16.000000	17.000000	18.000000

```
y = amean(x,1);
```

Using the same array x as the above example, this example computes the mean across the first dimension. y will be a 2x3x1 array, such that:

[1,1,1] through [1,3,1] =

```
2.5000000
6.5000000
10.500000
```

[2,1,1] through [2,3,1] =

```
14.500000
18.500000
22.500000
```

See Also

[asum](#)

AmericanBinomCall

Purpose

Prices American call options using binomial method.

Format

```
c = AmericanBinomCall(S0, K, r, div, tau, sigma, N);
```

AmericanBinomCall

Input

S_0	scalar, current price.
K	Mx1 vector, strike prices.
r	scalar, risk free rate.
div	continuous dividend yield.
τ	scalar, elapsed time to exercise in annualized days of trading.
σ	scalar, volatility.
N	number of time segments.

Output

c	Mx1 vector, call premiums.
-----	----------------------------

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach," *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
S0 = 718.46;  
K = { 720, 725, 730 };  
r = .0498;  
sigma = .2493;  
  
t0 = dtday(2001, 1, 30);  
t1 = dtday(2001, 2, 16);  
tau = elapsedTradingDays(t0,t1) /  
      annualTradingDays(2001);
```

```
c = AmericanBinomCall(S0,K,r,0,tau,sigma,60);  
print c;
```

produces the output:

```
17.344044  
15.058486  
12.817427
```

Source

finprocs.src

AmericanBinomCall_Greeks

Purpose

Computes Delta, Gamma, Theta, Vega, and Rho for American call options using binomial method.

Format

```
{ d, g, t, v, rh } = AmericanBinomCall_Greeks(S0, K, r,  
div, tau, sigma, N);
```

Input

S_0	scalar, current price.
K	Mx1 vector, strike prices.
r	scalar, risk free rate.
div	continuous dividend yield.

AmericanBinomCall_Greeks

τ	scalar, elapsed time to exercise in annualized days of trading.
σ	scalar, volatility.
N	number of time segments.

Global Input

_fin_thetaType	scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.
_fin_epsilon	scalar, finite difference stepsize. Default = 1e-8.

Output

d	Mx1 vector, delta.
g	Mx1 vector, gamma.
t	Mx1 vector, theta.
v	Mx1 vector, vega.
ρ	Mx1 vector, rho.

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach," *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
S0 = 305;  
K = 300;
```



```
r = .08;  
sigma = .25;  
tau = .33;  
div = 0;  
  
print AmericanBinomCall_Greeks(S0,K,r,0,tau,sigma,30);
```

produces:

```
0.70631204  
0.00076381912  
-17.400851  
68.703851  
76.691829
```

Source

finprocs.src

See Also

[AmericanBinomCall_ImpVol](#), [AmericanBinomCall](#), [AmericanBinomPut Greeks](#),
[AmericanBSCall Greeks](#)

AmericanBinomCall_ImpVol

Purpose

Computes implied volatilities for American call options using binomial method.

Format

```
sigma = AmericanBinomCall_ImpVol(c, S0, K, r, div, tau,  
N);
```

AmericanBinomCall_ImpVol

Input

c	Mx1 vector, call premiums
S_0	scalar, current price.
K	Mx1 vector, strike prices.
r	scalar, risk free rate.
div	continuous dividend yield.
τ	scalar, elapsed time to exercise in annualized days of trading.
N	number of time segments.

Output

σ	Mx1 vector, volatility.
----------	-------------------------

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach," *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
c = { 13.70, 11.90, 9.10 };
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;
div = 0;

t0 = dtday(2001, 1, 30);
t1 = dtday(2001, 2, 16);
tau = elapsedTradingDays(t0,t1) /
```

```
annualTradingDays(2001);

sigma = AmericanBinomCall_ImpVol(c,S0,K,r,0,tau,30);
print sigma;
```

produces:

```
0.19629517
0.16991943
0.12874756
```

Source

finprocs.src

AmericanBinomPut

Purpose

Prices American put options using binomial method.

Format

```
c = AmericanBinomPut(S0, K, r, div, tau, sigma, N);
```

Input

S_0	scalar, current price.
K	Mx1 vector, strike prices.
r	scalar, risk free rate.
div	continuous dividend yield.
tau	scalar, elapsed time to exercise in annualized days of

AmericanBinomPut

	trading.
σ	scalar, volatility.
N	number of time segments.

Output

c	Mx1 vector, put premiums.
-----	---------------------------

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach," *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;
sigma = .2493;

t0 = dtday(2001, 1, 30);
t1 = dtday(2001, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2001);

c = AmericanBinomPut(S0,K,r,0,tau,sigma,60);
print c;
```

produces:

```
16.986117
19.729923
22.548538
```

Source

finprocs.src

AmericanBinomPut_Greeks

Purpose

Computes Delta, Gamma, Theta, Vega, and Rho for American put options using binomial method.

Format

```
{ d, g, t, v, rh } = AmericanBinomPut_Greeks(S0, K, r,
div, tau, sigma, N);
```

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.
<i>N</i>	number of time segments.

AmericanBinomPut_Greeks

Global Input

<code>_fin_</code>	scalar, if 1, one day look ahead, else, infinitesimal.
<code>thetaType</code>	Default = 0.
<code>_fin_</code>	scalar, finite difference stepsize. Default = 1e-8.
<code>epsilon</code>	

Output

<code>d</code>	Mx1 vector, delta.
<code>g</code>	Mx1 vector, gamma.
<code>t</code>	Mx1 vector, theta.
<code>v</code>	Mx1 vector, vega.
<code>rh</code>	Mx1 vector, rho.

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach," *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
S0 = 305;
K = 300;
r = .08;
div = 0;
sigma = .25;
tau = .33;

print AmericanBinomPut_Greeks (S0,K,r,0,tau,sigma,60);
```

produces

```
-0.38324908  
0.00076381912  
8.1336630  
68.337294  
-27.585043
```

Source

finprocs.src

See Also

[AmericanBinomPut_ImpVol](#), [AmericanBinomPut](#), [AmericanBinomCall Greeks](#),
[AmericanBSPut Greeks](#)

AmericanBinomPut_ImpVol

Purpose

Computes implied volatilities for American put options using binomial method.

Format

```
sigma = AmericanBinomPut_ImpVol(c, S0, K, r, div, tau, N);
```

Input

<i>c</i>	Mx1 vector, put premiums
<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.

AmericanBinomPut_ImpVol

<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>N</i>	number of time segments.

Output

<i>sigma</i>	Mx1 vector, volatility.
--------------	-------------------------

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach," *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
p = { 14.60, 17.10, 20.10 };
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;
div = 0;

t0 = dtday(2001, 1, 30);
t1 = dtday(2001, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2001);

sigma = AmericanBinomPut_ImpVol(p,S0,K,r,0,tau,30);
print sigma;
```

produces:


```
0.12466064
0.16583252
0.21203735
```

Source

finprocs.src

AmericanBSCall

Purpose

Prices American call options using Black, Scholes, and Merton method.

Format

c = **AmericanBSCall**(*S0*, *K*, *r*, *div*, *tau*, *sigma*);

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.

Output

<i>c</i>	Mx1 vector, call premiums.
----------	----------------------------

AmericanBSCall_Greeks

Example

```
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;
sigma = .2493;

t0 = dtday(2001, 1, 30);
t1 = dtday(2001, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2001);

c = AmericanBSCall(S0,K,r,0,tau,sigma);
print c;
```

produces:

```
32.005720
31.083232
30.367548
```

Source

finprocs.src

AmericanBSCall_Greeks

Purpose

Computes Delta, Gamma, Theta, Vega, and Rho for American call options using Black, Scholes, and Merton method.

Format

```
{ d, g, t, v, rh } = AmericanBSCall_Greeks(S0, K, r, div,
tau, sigma);
```

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.

Global Input

<i>_fin_thetaType</i>	scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.
<i>_fin_epsilon</i>	scalar, finite difference stepsize. Default = 1e-8.

Output

<i>d</i>	Mx1 vector, delta.
<i>g</i>	Mx1 vector, gamma.
<i>t</i>	Mx1 vector, theta.
<i>v</i>	Mx1 vector, vega.
<i>rh</i>	Mx1 vector, rho.

Example

```
S0 = 305;
```

AmericanBSCall_ImpVol

```
K = 300;  
r = .08;  
sigma = .25;  
tau = .33;  
print AmericanBSCall_Greeks(S0,K,r,0,tau,sigma);
```

produces:

```
0.40034039  
0.016804021  
-55.731079  
115.36906  
46.374528
```

Source

finprocs.src

See Also

[AmericanBSCall_ImpVol](#), [AmericanBSCall](#), [AmericanBSPut Greeks](#),
[AmericanBinomCall Greeks](#)

AmericanBSCall_ImpVol

Purpose

Computes implied volatilities for American call options using Black, Scholes, and Merton method.

Format

```
sigma = AmericanBSCall_ImpVol(c, S0, K, r, div, tau);
```

Input

<i>c</i>	Mx1 vector, call premiums.
<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.

Output

<i>sigma</i>	Mx1 vector, volatility.
--------------	-------------------------

Example

```
c = { 13.70, 11.90, 9.10 };
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;

t0 = dtday(2001, 1, 30);
t1 = dtday(2001, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2001);

sigma = AmericanBSCall_ImpVol(c,S0,K,r,0,tau);
print sigma;
```

produces:

AmericanBSPut

```
0.10259888
0.088370361
0.066270752
```

Source

finprocs.src

AmericanBSPut

Purpose

Prices American put options using Black, Scholes, and Merton method.

Format

```
c = AmericanBSPut(S0, K, r, div, tau, sigma);
```

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.

Output

<i>c</i>	Mx1 vector, put premiums.
----------	---------------------------

Example

```
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;
sigma = .2493;

t0 = dtday(2001, 1, 30);
t1 = dtday(2001, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2001);

c = AmericanBSPut(S0,K,r,0,tau,sigma);
print c;
```

produces:

```
16.870783
19.536842
22.435487
```

Source

finprocs.src

AmericanBSPut_Greeks

Purpose

Computes Delta, Gamma, Theta, Vega, and Rho for American put options using Black, Scholes, and Merton method.

AmericanBSPut_Greeks

Format

```
{ d, g, t, v, rh } = AmericanBSPut_Greeks(S0, K, r, div, tau, sigma);
```

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.

Global Input

<i>_fin_thetaType</i>	scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.
<i>_fin_epsilon</i>	scalar, finite difference stepsize. Default = 1e-8.

Output

<i>d</i>	Mx1 vector, delta.
<i>g</i>	Mx1 vector, gamma.
<i>t</i>	Mx1 vector, theta.
<i>v</i>	Mx1 vector, vega.
<i>rh</i>	Mx1 vector, rho.

Example

```
S0 = 305;  
K = 300;  
r = .08;  
sigma = .25;  
tau = .33;  
  
print AmericanBSPut_Greeks(S0,K,r,0,tau,sigma);
```

produces:

```
-0.33296721  
0.0091658294  
-17.556118  
77.614237  
-40.575963
```

Source

finprocs.src

See Also

[AmericanBSCall ImpVol](#), [AmericanBSCall Greeks](#), [AmericanBSPut ImpVol](#)

AmericanBSPut_ImpVol

Purpose

Computes implied volatilities for American put options using Black, Scholes, and Merton method.

AmericanBSPut_ImpVol

Format

```
sigma = AmericanBSPut_ImpVol(c, S0, K, r, div, tau);
```

Input

<i>c</i>	Mx1 vector, put premiums.
<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.

Output

<i>sigma</i>	Mx1 vector, volatility.
--------------	-------------------------

Example

```
p = { 14.60, 17.10, 20.10 };
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;

t0 = dtday(2001, 1, 30);
t1 = dtday(2001, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2001);

sigma = AmericanBSPut_ImpVol(p,S0,K,r,0,tau);
print sigma;
```

produces:

```
0.12753662
0.16780029
0.21396729
```

Source

`finprocs.src`

amin

Purpose

Moves across one dimension of an N-dimensional array and finds the smallest element.

Format

```
y = amin(x, dim);
```

Input

<i>x</i>	N-dimensional array.
<i>dim</i>	scalar, number of dimension across which to find the minimum value.

Output

<i>y</i>	N-dimensional array.
----------	----------------------

amin

Remarks

The output *y*, will have the same sizes of dimensions as *x*, except that the dimension indicated by *dim* will be collapsed to 1.

Example

3

```
//Setting the rng seed allows for repeatable
//random numbers
rndseed 8237348;

//Create a 24x1 vector of random normal numbers
//with a standard deviation of 10 and then round
//to the nearest integer value
x = round(10*rndn(24,1));

//Reshape the 24x1 vector into a 2x3x4 dimensional array
//NOTE: The pipe operator '|' is for vertical concatenation
x = areshape(x,2|3|4);

dim = 2;
y = amin(x,dim);
```

x is a 2x3x4 array, such that:

[1,1,1] through [1,3,4] =

1.0000000	-11.000000	9.0000000	-8.0000000
-2.0000000	-10.000000	-6.0000000	-5.0000000
-5.0000000	17.000000	9.0000000	-2.0000000

[2,1,1] through [2,3,4] =

-4.0000000	-2.0000000	7.0000000	-2.0000000
4.0000000	13.000000	-16.000000	11.000000
2.0000000	-1.0000000	12.000000	-16.000000

y will be a 2x1x4 array, such that:

[1,1,1] through [1,1,4] =

-5.0000000	-11.000000	-6.0000000	-8.0000000
------------	------------	------------	------------

[2,1,1] through [2,1,4] =

-4.0000000	-2.0000000	-16.000000	-16.000000
------------	------------	------------	------------

```
y = amin(x,1);
```

Using the same array x as the above example, this example finds the minimum value across the first dimension.

y will be a 2x3x1 array, such that:

[1,1,1] through [1,3,1] =

-11.000000
-10.000000
-5.0000000

[2,1,1] through [2,3,1] =

-4.0000000
-16.000000
-16.000000

See Also

[amax](#), [minc](#)

amult

amult

Purpose

Performs matrix multiplication on the planes described by the two trailing dimensions of N-dimensional arrays.

Format

```
y = amult(a, b);
```

Input

a

a	N-dimensional array.
b	N-dimensional array.

Output

y	N-dimensional array, containing the product of the matrix multiplication of the planes described by the two trailing dimensions of a and b.
---	---

Remarks

All leading dimensions must be strictly conformable, and the two trailing dimensions of each array must be matrix-product conformable.

Example

```
//Create an additive sequence from 1-12 and reshape it into  
//a 2x3x2 dimensional array  
a = areshape(seqa(1,1,12),2|3|2);
```

```

b = areshape(seqa(1,1,16),2|2|4);

//Multiply the two 3x2 matrices in 'a' by the corresponding
//2x4 matrices in 'b'
y = amult(a,b);

```

a is a 2x3x2 array, such that:

[1,1,1] through [1,3,2] =

1.0000000	2.0000000
3.0000000	4.0000000
5.0000000	6.0000000

[2,1,1] through [2,3,2] =

7.0000000	8.0000000
9.0000000	10.000000
11.000000	12.000000

b is a 2x2x4 array, such that:

[1,1,1] through [1,2,4] =

1.0000000	2.0000000	3.0000000	4.0000000
5.0000000	6.0000000	7.0000000	8.0000000

[2,1,1] through [2,2,4] =

9.0000000	10.000000	11.000000	12.000000
13.000000	14.000000	15.000000	16.000000

y will be a 2x3x4 array, such that:

[1,1,1] through [1,3,4] =

annotationGetDefaults

11.000000	14.000000	17.000000	20.000000
23.000000	30.000000	37.000000	44.000000
35.000000	46.000000	57.000000	68.000000

[2,1,1] through [2,3,4] =

167.00000	182.00000	197.00000	212.00000
211.00000	230.00000	249.00000	268.00000
255.00000	278.00000	301.00000	324.00000

annotationGetDefaults

Purpose

Fills in an instance of a **plotAnnotation** structure with default values.

Format

```
myAnnotation = annotationGetDefaults();
```

Output

<i>myAnnotation</i>	An instance of a plotAnnotation structure with all members set to defaults.
---------------------	--

Remarks

plotAnnotation structures are used with the **annotationSet** functions to programmatically control the attributes of the annotations that you add to graphs.

To see a full example of adding an annotation to a graph, see the command reference page for **plotAddLine**, **plotAddShape** or **plotAddTextbox**

Example

```
//Declare 'myAnnotation' to be an instance of a plotAn-
notation structure
struct plotAnnotation myAnnotation;

//Fill in 'myAnnotation' with default values
myAnnotation = annotationGetDefaults() ;
```

See Also

[plotAddShape](#), [plotAddTextbox](#), [annotationSetLineColor](#), [annotationSetBkd](#)

annotationSetBkd

Purpose

Sets the background color and transparency level for a textbox, rectangle or ellipse.

Format

```
annotationSetBkd(&myAnnotation, color, transparency);
```

Input

<i>&myAnnotation</i>	A pointer to an instance of a plotAnnotation structure.
<i>color</i>	String, color name or hex HTML color code.
<i>transparency</i>	Scalar, transparency percentage. Valid range is between 0 and 1.

Examples

Example 1: Using a color string

```
//Declare 'myAnnotation' to be an instance of a plotAn-
notation structure
//and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set background to light gray with 40% opacity
annotationSetBkd(&myAnnotation, "light gray", 0.4);
```

Example 2: Full example using an HTML color code

```
//Create and plot some simple data
x = sega(1, 1, 10);
y = rndu(10, 1);
plotXY(x, y);

//Declare 'myAnnotation' to be an instance of a plotAn-
notation
//structure and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set background to white with 80% opacity
annotationSetBkd(&myAnnotation, "#FFFFFF", 0.8);

//Add rectangle to 'xy' plot from above
//using settings from 'myAnnotation'
plotAddShape(myAnnotation, "rectangle", 1, 0.2, 3, 0.5);
```

See Also

[plotAddTextbox](#), [plotAddShape](#), [annotationGetDefaults](#)

annotationSetFont

Purpose

Sets the font properties of a **plotAnnotation** structure for controlling text boxes added to a graph.

Format

```
annotationSetFont(&myAnnotation, fontname, fontsize, fontcolor);
```

Input

&myAnnotation	A plotAnnotation structure pointer.
fontname	String, the name of the font.
fontsize	Scalar, the size of the font in points.
fontcolor	String, a color or HTML hexadecimal color code.

Remarks

annotationSetFont does not currently support surface plots.

Examples

Example 1: Basic usage

```
//Declare an instance of a 'plotAnnotation' structure
//and fill in with default values
struct plotAnnotation myAnnotation;
```

annotationSetFont

```
myAnnotation = annotationGetDefaults();

annotationSetFont(&myAnnotation, "arial", 14, "black");
```

Example 2: Customized textbox

```
//Create a simple plot on which to add a textbox
x = seqa(pi, 0.1, 50);
plotXY(x, sin(x) + rndu(50, 1));

//Declare instance of plotAnnotation structure
//and fill in with default values
struct plotAnnotation myTextbox;
myTextbox = annotationGetDefaults();

//Set font to dark-gray, 14pt times
annotationSetFont(&myTextbox, "times", 14, "dark gray");

//Create text for textbox
box_text = "Trend change in Q2";

//The top-left corner of the text box
//will start at the point (3.5,1.5)
x_start = 3.5;
y_start = 1.5;

//Add textbox to last draw graph
plotAddTextbox(myTextbox, box_text, x_start, y_start);
```

See Also

[plotAddShape](#), [plotAddTextbox](#), [annotationGetDefaults](#)

annotationSetLineColor

Purpose

Sets the line color for textbox, rectangle or ellipse borders as well as the color for lines and arrows.

Format

```
annotationSetLineColor(&myAnnotation, color);
```

Input

<i>&myAnnotation</i>	A pointer to an instance of a plotAnnotation structure.
<i>color</i>	String, named color or HTML hexadecimal color code. Note that HTML color codes must start with a #.

Examples

Example 1: Basic usage with named color

```
//Declare 'myAnnotation' to be an instance of a plotAn-
notation structure
//and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set line color
annotationSetLineColor (&myAnnotation, "blue");
```

Example 2: Basic usage with HTML color code

annotationSetLineColor

```
//Declare 'myAnnotation' to be an instance of a plotAn-
notation structure
//and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set line color
annotationSetLineColor(&myAnnotation, "#CCCCCC");
```

Example 3: Full example adding a red arrow to a graph

```
//Create and plot some simple data
x = sega(0.1, 0.1, 30);
y = cos(x);
plotXY(x, y);

//Declare 'myAnnotation' to be an instance of a plotAn-
notation
//structure and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set line color for arrow
annotationSetLineColor(&myAnnotation, "red");

//Add arrow to 'xy' plot from above
//using settings from 'myAnnotation'
x_start = 0.15;
y_start = 0.2;
x_end = 1;
y_end = 0.5;
head_size = 15;
plotAddArrow(myAnnotation, x_start, y_start, x_end, y_end,
head_size);
```

See Also

[plotAddTextbox](#), [plotAddArrow](#), [plotAddShape](#), [annotationGetDefaults](#)

annotationSetLineStyle

Purpose

Sets the line style for textbox, rectangle or ellipse borders as well as the style for lines and arrows.

Format

```
annotationSetLineStyle(&myAnnotation, style);
```

Input

<i>&myAnnotation</i>	A pointer to an instance of a plotAnnotation structure.
<i>style</i>	Matrix, line style. Valid options include: <ul style="list-style-type: none"> • 1 - solid • 2 - dash • 3 - dot • 4 - dash-dot • 5 - dash-dot-dot

Examples

Example 1: Basic usage

```
//Declare 'myAnnotation' to be an instance of a
```

annotationSetLineStyle

```
plotAnnotation structure
//and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set line style to 'dot'
line_style = 3;
annotationSetLineStyle(&myAnnotation,line_style);
```

Example 2: Full example creating an ellipse with a dash border

```
//Create and plot some simple data
x = sega(0.1, 0.2, 10);
y = 3 .* sin(x) + randu(10, 1);
plotXY(x, y);

//Declare 'myAnnotation' to be an instance of a plotAn-
notation
//structure and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set line style to dash (for ellipse border in this case)
annotationSetLineStyle(&myAnnotation, 2);

//Add ellipse to 'xy' plot from above
//using settings from 'myAnnotation'
plotAddShape(myAnnotation, "ellipse", 0.4, 1.5, 1, 2.9);
```

See Also

[plotAddTextbox](#), [plotAddArrow](#), [plotAddShape](#), [annotationGetDefaults](#)

annotationSetLineThickness

Purpose

Sets the line thickness for textbox, rectangle or ellipse borders as well as the color for lines and arrows.

Format

```
annotationSetLineThickness(&myAnnotation, thickness);
```

Input

<i>myAnnotation</i>	A pointer to an instance of a plotAnnotation structure.
<i>thickness</i>	Scalar, the thickness of the line in pixels.

Examples

Example 1: Basic usage

```
//Declare 'myAnnotation' to be an instance of a plotAn-
notation structure
//and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set line thickness to 1 pixel
annotationSetLineThickness(&myAnnotation, 1);
```

Example 2: Full example setting thickness to zero to turn off rectangle border

```
//Create and plot some simple data
```

annualTradingDays

```
x = sega(-1.5, 0.1, 31);
y = cos(x);
plotXY(x, y);

//Declare 'myAnnotation' to be an instance of a plotAn-
notation
//structure and fill it in with default values
struct plotAnnotation myAnnotation;
myAnnotation = annotationGetDefaults();

//Set line thickness to 1 pixel (for rectangle border in
this case)
annotationSetLineThickness(&myAnnotation, 1);

//Add rectangle to 'xy' plot from above
//using settings from 'myAnnotation'
x_start = -pi ./ 4;
y_start = 0.07;
x_end = pi ./ 4;
y_end = 0.71;

plotAddShape(myAnnotation, "rectangle", x_start, y_start,
x_end, y_end);
```

See Also

[plotAddTextbox](#), [plotAddArrow](#), [plotAddShape](#), [annotationGetDefaults](#)

annualTradingDays

Purpose

Computes number of trading days in a given year.

Format

```
 $n$  = annualTradingDays( $a$ );
```

Input

a	scalar, year.
-----	---------------

Output

n	number of trading days in year.
-----	---------------------------------

Remarks

A trading day is a weekday that is not a holiday as defined by the New York Stock Exchange from 1888 through 2012. Holidays are defined in `holidays.asc`. You may edit that file to modify or add holidays.

Source

`finutils.src`

Globals

`_fin_annualTradingDays`, `_fin_holidays`

See Also

[eTD](#), [gNTD](#), [gPTD](#), [gNWD](#), [gPWD](#)

arccos

arccos

Purpose

Computes the inverse cosine.

Format

```
y = arccos(x);
```

Input

x	NxK matrix or N-dimensional array.
-----	------------------------------------

Output

y	NxK matrix or N-dimensional array containing the angle in radians whose cosine is x .
-----	---

Remarks

If x is complex or has any elements whose absolute value is greater than 1, complex results are returned.

Example

```
//Format print statements to show 3 digits
//after the decimal point
format /rd 6,3;

x = { -1, -0.5, 0, 0.5, 1 };
y = arccos(x);

print "x = " x;
```

```
print "y = " y;
```

The code above, produces the following output:

```
x =  
  -1.000  
  -0.500  
   0.000  
   0.500  
   1.000  
y =  
  3.142  
  2.094  
  1.571  
  1.047  
  0.000
```

Source

trig.src

arcsin

Purpose

Computes the inverse sine.

Format

```
y = arcsin(x);
```

Input

x	NxK matrix or N-dimensional array.
---	------------------------------------

areshape

Output

y

$N \times K$ matrix or N -dimensional array, the angle in radians whose sine is x .

Remarks

If x is complex or has any elements whose absolute value is greater than 1, complex results are returned.

Example

```
//Set 'x' to be the sequence -1, -0.5, 0, 0.5, 1  
x = sega(-1, 0.5, 5);  
y = arcsin(x);
```

Assigns y to be equal to:

```
-1.5707963  
-0.52359878  
0.00000000  
0.52359878  
1.5707963
```

Source

trig.src

areshape

Purpose

Reshapes a scalar, matrix, or array into an array of user-specified size.

Format

```
y = areshape(x, o);
```

Input

x	scalar, matrix, or N-dimensional array.
o	Mx1 vector of orders, the sizes of the dimensions of the new array.

Output

y	M-dimensional array, created from data in x.
---	--

Remarks

If there are more elements in *x* than in *y*, the remaining elements are discarded. If there are not enough elements in *x* to fill *y*, then when **areshape** runs out of elements, it goes back to the first element of *x* and starts getting additional elements from there.

Example

```
x = 3;  
orders = { 2,3,4 };  
y = areshape(x,orders);
```

y will be a 2x3x4 array of threes.

```
x = reshape(seqa(1,1,90),30,3);  
orders = { 2,3,4,5 };  
y = areshape(x,orders);
```

arrayalloc

y will be a 2x3x4x5 array. Since *y* contains 120 elements and *x* contains only 90, the first 90 elements of *y* will be set to the sequence of integers from 1 to 90 that are contained in *x*, and the last 30 elements of *y* will be set to the sequence of integers from 1 to 30 contained in the first 30 elements of *x*.

```
x = reshape(seqa(1,1,60),20,3);  
orders = { 3,2,4 };  
y = areshape(x,orders);
```

y will be a 3x2x4 array. Since *y* contains 24 elements, and *x* contains 60, the elements of *y* will be set to the sequence of integers from 1 to 24 contained in the first 24 elements of *x*.

See Also

[aconcat](#)

arrayalloc

Purpose

Creates an N-dimensional array with unspecified contents.

Format

```
y = arrayalloc(o, cf);
```

Input

<i>o</i>	Nx1 vector of orders, the sizes of the dimensions of the array.
<i>cf</i>	scalar, 0 to allocate real array, or 1 to allocate complex array.

Output

y N-dimensional array.

Remarks

The contents are unspecified. To create a new array with all elements initialized to a particular scalar value, use **arrayinit**.

arrayalloc is used to allocate an array that will be written to in sections using [setarray](#), or indexed assignments. It is much faster to preallocate an array and fill in sections during a loop rather than adding new sections with concatenation.

Example

```
orders = { 2,3,4 };
y = arrayalloc(orders, 1);
```

y will be a complex 2x3x4 array with unspecified contents.

```
//Tell GAUSS to replace all instances of 'REAL' with a 0
#define REAL 0
orders = { 7, 5, 3 };

//Create a real 7x5x3 dimensional array; before GAUSS
//interprets this statement it will replace 'REAL' with
//a scalar 0
y = arrayalloc(orders, REAL);
```

See Also

[arrayinit](#), [setarray](#)

arrayindex

arrayindex

Purpose

Converts a scalar vector index to a vector of indices for an N-dimensional array.

Format

```
i = arrayindex(scalar_idx, orders);
```

Input

<i>scalar_idx</i>	scalar, index into vector or 1-dimensional array.
<i>orders</i>	Nx1 vector of orders of an N-dimensional array.

Output

<i>i</i>	Nx1 vector of indices, index of corresponding element in N-dimensional array.
----------	---

Remarks

This function and its opposite, **singleindex**, allow you to easily convert between an N-dimensional index and its corresponding location in a 1-dimensional object of the same size.

Example

```
//Set the rng seed for repeatable random numbers  
rndseed 982348;
```

```

orders = { 2,3,4,5 };

//Create 120x1 vector of uniform random numbers
// (2*3*4*5 = 120)
v = randu(prodc(orders),1);

//Reshape the 120x1 random vector into a
//2x3x4x5 dimensional array
a = areshape(v,orders);

vi = 50;
ai = arrayindex(vi,orders);

print "vi = " vi;
print "ai = " ai;
print "v[vi] = " v[vi];
print "getarray(a, ai) = "; getarray(a,ai);

```

The code above, produces the following output:

```

vi =    50.000
ai =
    1.000
    3.000
    2.000
    5.000
v[vi] =    0.047
getarray(a, ai) =    0.047

```

This example allocates a vector of random numbers and creates a 4-dimensional array using the same data. The 50th element of the vector *v* corresponds to the element of array *a* that is indexed with *ai*.

See Also

[singleindex](#)

arrayinit

arrayinit

Purpose

Creates an N-dimensional array with a specified fill value.

Format

```
y = arrayinit(orders, value);
```

Input

<i>orders</i>	Nx1 vector of orders, the sizes of the dimensions of the array.
<i>value</i>	scalar, value to initialize. If <i>value</i> is complex the result will be complex.

Output

<i>y</i>	N-dimensional array with each element equal to the value of <i>value</i> .
----------	--

Example

```
val = 3.14;  
orders = { 2, 100, 9 };  
y = arrayinit(orders, val);
```

y will be a 2x100x9 array with each element equal to 3.14.

See Also

[arrayalloc](#)

arraytomat

Purpose

Converts an array to type matrix.

Format

```
y = arraytomat(a);
```

Input

<i>a</i>	N-dimensional array.
----------	----------------------

Output

<i>y</i>	KxL or 1xL matrix or scalar, where L is the size of the fastest moving dimension of the array and K is the size of the second fastest moving dimension.
----------	---

Remarks

arraytomat will take an array of 1 or 2 dimensions or an N-dimensional array, in which the N-2 slowest moving dimensions each have a size of 1.

Example

```
//Create 25x1 vector containing the sequence 0.5, 1,  
//1.5...12.5  
x = seqa(0.5, 0.5, 25);  
  
//Reshape into a 1x6x4 array, discarding the 25th element
```

asciiload

```
//of 'x'
a = areshape(x, 1|6|4);

//Set 'y' to be a 6x4 variable of type matrix, with the
//same contents as 'a'
y = arraytomat(a);
```

The code above sets *y* equal to:

0.5	1.0	1.5	2.0
2.5	3.0	3.5	4.0
4.5	5.0	5.5	6.0
6.5	7.0	7.5	8.0
8.5	9.0	9.5	10.0
10.5	11.0	11.5	12.0

See Also

[mattoarray](#)

asciiload

Purpose

Loads data from a delimited ASCII text file into an Nx1 vector. NOTE: This function is deprecated. Use **csvReadM** instead.

Format

```
y = asciiload(filename);
```

Input

<i>filename</i>	string, name of data file.
-----------------	----------------------------

Output

y	Nx1 vector.
-----	-------------

Remarks

NOTE: This function is deprecated. Use **csvReadM** instead.

The file extension must be included in the file name.

Numbers in ASCII files must be delimited with spaces, commas, tabs, or newlines.

This command loads as many elements as possible from the file into an Nx1 vector. This allows you to verify if the load was successful by calling **rows(y)** after **asciiload** to see how many elements were actually loaded. You may then **reshape** the Nx1 vector to the desired form. You could, for instance, put the number of rows and columns of the matrix right in the file as the first and second elements and **reshape** the remainder of the vector to the desired form using those values.

Example

To load the file `myfile.asc`, containing the following data:

```
2.805  16.568
-4.871   3.399
17.361 -12.725
```

you may use any of the following commands:

```
//This statement assumes 'myfile.asc' is in the current
//working directory
y = asciiload("myfile.asc");
```

asclabel

```
//This code assumes that 'myfile.asc' is
//located in the C:\gauss17 directory
//Note the double backslashes for path separators
fpath = "C:/gauss/myfile.asc";
y = asciiload(fpath);

path = "C:/gauss/";
fname = "myfile.asc";
//The '$+' operator adds two strings together into one
//string
y = asciiload(path$+fname);
```

All of the above commands will set *y* to be equal to:

```
2.805
16.568
-4.871
3.399
17.361
-12.725
```

See Also

[csvReadM](#), [load](#), [dataload](#)

asclabel

Purpose

To set up character labels for the X and Y axes. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
asclabel(xl, yl);
```

Input

<i>xl</i>	string or Nx1 character vector, labels for the tick marks on the X axis. Set to 0 if no character labels for this axis are desired.
<i>yl</i>	string or Mx1 character vector, labels for the tick marks on the Y axis. Set to 0 if no character labels for this axis are desired.

Example

This illustrates how to label the X axis with the months of the year:

```
library pgraph;  
let lab = JAN FEB MAR APR MAY JUN JUL AUG SEP OCT NOV DEC;  
asclabel(lab,0);
```

This will also work:

```
lab = "JAN FEB MAR APR MAY JUN JUL AUG SEP OCT NOV DEC";  
asclabel(lab,0);
```

If the string format is used, then escape characters may be embedded in the labels. For example, the following produces character labels that are multiples of λ . The font Simgrma must be previously loaded in a **fonts** command.

astd

```
fonts("simplex simgrma");  
lab = "\2010.25\2021 \2010.5\2021 \2010.75\2021 1";  
asclabel(lab,0);
```

Here, the "\2021" produces the "λ" symbol from Simgrma.

Source

pgraph.src

See Also

[xtics](#), [ytics](#), [scale](#), [scale3d](#), [fonts](#)

astd

Purpose

Computes the standard deviation of the elements across one dimension of an N-dimensional array.

Format

```
y = astd(x, dim);
```

Input

<i>x</i>	N-dimensional array.
<i>dim</i>	scalar, number of dimension to sum across.

Output

<i>y</i>	N-dimensional array, standard deviation across specified dimension of <i>x</i> .
----------	--

Remarks

The output y , will have the same sizes of dimensions as x , except that the dimension indicated by `dim` will be collapsed to 1.

For each column, this function essentially computes sample standard deviation, s :

$$s = \sqrt{\frac{1}{n-1} \times \sum_{i=1}^n (X_i - \bar{X})^2}$$

Thus, the divisor is N-1 rather than N, where N is the number of elements being summed. See **astds** for the alternate definition.

Example

```
//Create a 1e6x1 vector of random normal numbers with a
//standard deviation of 25 and reshape it into a
//2e5x3x2 array
rndseed 456;
a = areshape(25*rndn(2e6,1),2e5|3|2);
y = astd(a,3);
```

The code above should produce a 3x2 matrix with all elements close to 25 similar to what we see below.

25.070091	24.994774
24.988263	24.990370
24.956467	24.987882

See Also

[astds](#), [stdc](#)

astds

astds

Purpose

Computes the biased standard deviation of the elements across one dimension of an N-dimensional array.

Format

$y = \text{astds}(x, \text{dim});$

Input

x	N-dimensional array.
dim	scalar, number of dimension to sum across.

Output

y	N-dimensional array, standard deviation across specified dimension of x .
-----	---

Remarks

The output y , will have the same sizes of dimensions as x , except that the dimension indicated by dim will be collapsed to 1.

This function essentially computes:

$$\sigma = \sqrt{\frac{1}{n} \times \sum_{i=1}^n (X_i - \mu)^2}$$

Thus, the divisor is N rather than $N-1$, where N is the number of elements being summed. See **astd** for the alternate definition.

Example

```
a = areshape(25*randn(16,1),4|2|2);
y = astds(a,3);

print "a = " a;
print "y = " y;
```

The code above produces the following output (due to the use of random data in this example your answers will be different):

```
a =

Plane [1,...]

    12.538   -56.786
   -40.283   -58.287

Plane [2,...]

     4.047    -0.325
    17.617    -9.248

Plane [3,...]

    17.908    40.048
     8.916   -37.247

Plane [4,...]

    -0.977    16.058
```

3

asum

```
-38.189    0.984

y =

Plane [1,...]

    7.321    35.659
   26.441    23.333
```

In this example, 16 standard Normal random variables are generated. They are multiplied by 25 and **areshape**'d into a 4x2x2 array, and the standard deviation is computed across the third dimension of the array.

See Also

[astd](#), [stdsc](#)

asum

Purpose

Computes the sum across one dimension of an N-dimensional array.

Format

```
y = asum(x, dim);
```

Input

<i>x</i>	N-dimensional array.
<i>dim</i>	scalar, number of dimension to sum across.

Output

y N-dimensional array.

Remarks

The output y , will have the same sizes of dimensions as x , except that the dimension indicated by dim will be collapsed to 1.

Example

```
x = sega(1,1,24);

dims = { 2, 3, 4 };
x = areshape(x,dims);

y = asum(x,3);
```

x is a 2x3x4 array, such that:

Plane [1, ., .]

1.000	2.000	3.000	4.000
5.000	6.000	7.000	8.000
9.000	10.000	11.000	12.000

Plane [2, ., .]

13.000	14.000	15.000	16.000
17.000	18.000	19.000	20.000
21.000	22.000	23.000	24.000

and y is equal to:

atan

Plane [1, ., .]

14.000	16.000	18.000	20.000
22.000	24.000	26.000	28.000
30.000	32.000	34.000	36.000

`y = asum(x,1);`

Using the same array `x` as the above example, this example computes the sum across the first dimension. `y` will be a 2x3x1 array, such that:

Plane [1, ., .]

10.000
26.000
42.000

Plane [2, ., .]

58.000
74.000
90.000

See Also

[amean](#)

atan

Purpose

Returns the arctangent of its argument.

Format

```
y = atan(x);
```

Input

x	NxK matrix or N-dimensional array.
-----	------------------------------------

Output

y	NxK matrix or N-dimensional array containing the arctangents of x in radians.
-----	---

Remarks

y will be the same size as x , containing the arctangents of the corresponding elements of x .

For real x , the arctangent of x is the angle whose tangent is x . The result is a value in radians in the range $-\pi/2$ to $+\pi/2$. To convert radians to degrees, multiply by $180/\pi$.

For complex x , the arctangent is defined everywhere except i and $-i$. If x is complex, y will be complex.

Example

```
//Create a sequence with 5 elements starting at -pi and
//increasing by pi/2
x = seqa(-pi, pi/2, 5)
y = atan(x);
print "x = " x;
print "y = " y;
```

After the code above:

atan2

```
-3.142    -1.263
-1.571    -1.004
x = 0.000  y = 0.000
 1.571     1.004
 3.142     1.263
```

See Also

[atan2](#), [sin](#), [cos](#), [pi](#), [tan](#)

atan2

Purpose

Computes an angle from an x , y coordinate.

Format

```
z = atan2(y, x);
```

Input

y	$N \times K$ matrix or P-dimensional array where the last two dimensions are $N \times K$, the y coordinate.
x	$L \times M$ matrix or P-dimensional array where the last two dimensions are $L \times M$, $E \times E$ conformable with y , the x coordinate.

Output

z	$\max(N,L)$ by $\max(K,M)$ matrix or P-dimensional array where the last two dimensions are $\max(N,L)$ by $\max(K,M)$.
-----	---

Remarks

Given a point x, y in a Cartesian coordinate system, **atan2** will give the correct angle with respect to the positive X axis. The answer will be in radians from $-\pi$ to $+\pi$.

To convert radians to degrees, multiply by $180/\pi$.

atan2 operates only on the real component of x , even if x is complex.

Example

```
//Create the sequence -pi, -pi/2, 0, pi/2, pi
x = seqa(-pi, pi/2, 5);
y = 1;

zpol = atan2(y,x);
zdeg = zpol*(180/pi);
print "x = " x;
print "zpol = " zpol;
print "zdeg = " zdeg;
```

After the code above:

-3.142	2.833	162.343
-1.571	2.575	147.518
x = 0.000	zpol = 1.571	zdeg = 90.000
1.571	0.567	32.482
3.142	0.308	17.657

See Also

[atan](#), [sin](#), [cos](#), [pi](#), [tan](#), [arcsin](#), [arccos](#)

atranspose

atranspose

Purpose

Transposes an N-dimensional array.

Format

```
y = atranspose(x, nd);
```

Input

<i>x</i>	N-dimensional array.
<i>nd</i>	Nx1 vector of dimension indices, the new order of dimensions.

Output

<i>y</i>	N-dimensional array, transposed according to <i>nd</i> .
----------	--

Remarks

The vector of dimension indices must be a unique vector of integers, 1-N, where 1 corresponds to the first element of the vector of orders.

Example

```
x = sega (1,1,24) ;  
x = areshape (x,2|3|4) ;  
nd = { 2,1,3 } ;  
y = atranspose (x,nd) ;
```

This example transposes the dimensions of `x` that correspond to the first and second elements of the vector of orders. `x` is a 2x3x4 array, such that:

Plane [1,...]

1.000	2.000	3.000	4.000
5.000	6.000	7.000	8.000
9.000	10.000	11.000	12.000

Plane [2,...]

13.000	14.000	15.000	16.000
17.000	18.000	19.000	20.000
21.000	22.000	23.000	24.000

`y` is a 3x2x4 array, such that:

Plane [1,...]

1.000	2.000	3.000	4.000
13.000	14.000	15.000	16.000

Plane [2,...]

5.000	6.000	7.000	8.000
17.000	18.000	19.000	20.000

Plane [3,...]

9.000	10.000	11.000	12.000
21.000	22.000	23.000	24.000

```
nd = { 2,3,1 };  
y = atranspose(x,nd);
```

axmargin

Using the same array `x` as the example above, this example transposes all three dimensions of `x`, returning a 3x4x2 array `y`, such that:

```
Plane [1,...]

    1.000    13.000
    2.000    14.000
    3.000    15.000
    4.000    16.000

Plane [2,...]

    5.000    17.000
    6.000    18.000
    7.000    19.000
    8.000    20.000

Plane [3,...]

    9.000    21.000
   10.000    22.000
   11.000    23.000
   12.000    24.000
```

See Also

[areshape](#)

axmargin

Purpose

Sets absolute margins for the plot axes which control placement and size of plot.
NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
axmargin(l, r, t, b);
```

Input

<i>l</i>	scalar, the left margin in inches.
<i>r</i>	scalar, the right margin in inches.
<i>t</i>	scalar, the top margin in inches.
<i>b</i>	scalar, the bottom margin in inches.

Remarks

axmargin sets an absolute distance from the axes to the edge of the graphic panel. Note that the user is responsible for allowing enough space in the margin if axes labels, numbers and title are used on the graph, since **axmargin** does not size the plot automatically as in the case of **margin**.

All input inch values for this procedure are based on a full size window of 9x6.855 inches. If this procedure is used within a graphic panel, the values will be scaled to window inches automatically.

If both **margin** and **axmargin** are used for a graph, **axmargin** will override any sizes specified by **margin**.

Example

The statement:

axmargin

```
library pgraph;  
axmargin(1,1,.5,.855);
```

will create a plot area of 7 inches horizontally by 5.5 inches vertically, and positioned 1 inch right and .855 up from the lower left corner of the graphic panel/page.

Source

pgraph.src

b

balance

Purpose

Balances a square matrix.

Format

$\{ b, z \} = \text{balance}(x);$

Input

x	KxK matrix or N-dimensional array where the last two dimensions are KxK.
-----	--

Output

b	KxK matrix or N-dimensional array where the last two dimensions are KxK, balanced matrix.
z	KxK matrix or N-dimensional array where the last two dimensions are KxK, diagonal scale matrix.

Remarks

balance returns a balanced matrix b and another matrix z with scale factors in powers of two on its diagonal. b is balanced in the sense that the absolute sums of the magnitudes of elements in corresponding rows and columns are nearly equal.

band

balance is most often used to scale matrices to improve the numerical stability of the calculation of their eigenvalues. It is also useful in the solution of matrix equations.

In particular,

$$b = z^{-1}xz$$

balance uses the BALANC function from EISPACK..

Example

```
let x[3,3] = 100 200 300
             40  50  60
             7   8   9;
{ b,z } = balance(x);

b = 100.0  100.0  37.5
     80.0   50.0  15.0
     56.0   32.0   9.0

z = 4.0  0.0  0.0
     0.0  2.0  0.0
     0.0  0.0  0.5
```

b

band

Purpose

Extracts bands from a symmetric banded matrix.

Format

```
a = band(y,  n);
```

Input

y	$K \times K$ symmetric banded matrix.
n	scalar, number of subdiagonals.

Output

a	$K \times (N+1)$ matrix, 1 subdiagonal per column.
-----	--

Remarks

y can actually be a rectangular $P \times Q$ matrix. K is then defined as $\min(P, Q)$. It will be assumed that a is symmetric about the principal diagonal for $y[1:K, 1:K]$.

The subdiagonals of y are stored right to left in a , with the principal diagonal in the rightmost or $(N+1)$ th column of a . The upper left corner of a is unused; it is set to 0.

This compact form of a banded matrix is what **bandchol** expects.

Example

```
x = { 1 2 0 0,
      2 8 1 0,
      0 1 5 2,
      0 0 2 3 };

//Extract only the principal diagonal
b0 = band(x, 0);

//Extract the principal diagonal and the first subdiagonal
b1 = band(x, 1);

//Extract the principal diagonal and the first two
```

bandchol

```
subdiagonals
b2 = band(x,2);
```

After the code above:

	1		0	1		0	0	1
b0 =	8	b1 =	2	8	b2 =	0	2	8
	5		1	5		0	1	5
	3		2	3		0	2	3

See Also

[bandchol](#), [bandcholsol](#), [bandltsol](#), [bandrv](#), [bandsolpd](#)

bandchol

Purpose

Computes the Cholesky decomposition of a positive definite banded matrix.

Format

```
l = bandchol(a);
```

Input

a	KxN compact form matrix.
-----	--------------------------

Output

l	KxN compact form matrix, lower triangle of the Cholesky decomposition of a .
-----	--

Remarks

Given a positive definite banded matrix A , there exists a matrix L , the lower triangle of the Cholesky decomposition of A , such that $A = LL'$. a is the compact form of A ; see **band** for a description of the format of a .

l is the compact form of L . This is the form of matrix that **bandcholsol** expects.

Example

```
x = { 1 2 0 0,
      2 8 1 0,
      0 1 5 2,
      0 0 2 3 };

bx = band(x, 1);
bl = bandchol(bx);

l = chol(x);
```

After the code above:

	0	1		0	1		1	2	0	0		
bx =	2	8		bl =	2	2		l =	0	2	1	0
	1	5			1	2			0	0	2	1
	2	3			1	1			0	0	0	1

See Also

[band](#), [bandcholsol](#), [bandltsol](#), [bandrv](#), [bandsolpd](#)

bandcholsol

bandcholsol

Purpose

Solves the system of equations $Ax = b$ for x , given the lower triangle of the Cholesky decomposition of a positive definite banded matrix A .

Format

```
 $x = \text{bandcholsol}(b, \textit{l});$ 
```

Input

b	$K \times M$ matrix.
\textit{l}	$K \times N$ compact form matrix.

Output

x	$K \times M$ matrix.
-----	----------------------

Remarks

Given a positive definite banded matrix A , there exists a matrix L , the lower triangle of the Cholesky decomposition of A , such that $A = LL'$. \textit{l} is the compact form of L ; see **band** for a description of the format of \textit{l} .

b can have more than one column. If so, $Ax = b$ is solved for each column. That is,

$$A * x[:, i] = b[:, i]$$

Example

```
//Create matrix 'A' and right-hand side 'b'
```

b

```

A = { 1 2 0 0,
      2 8 1 0,
      0 1 5 2,
      0 0 2 3 };
b = { 1.3, 2.1, 0.7, 1.8 };

//Create banded matrix form of 'A'
Aband = band(A,1);

//Cholesky factorization of the banded 'A'
Lband = bandchol(Aband);

//Solve the system of equations
x = bandcholsol(b, Lband);

```

After the code above is run:

	0.000	1.000		1.495		1.300		1.300
Lband =	2.000	2.000	x =	-0.098	b =	2.100	A*x =	2.100
	0.500	2.179		-0.110		0.700		0.700
	0.918	1.469		0.673		1.800		1.800

See Also

[band](#), [bandchol](#), [bandltsol](#), [bandrv](#), [bandsolpd](#)

bandltsol

Purpose

Solves the system of equations $Ax = b$ for x , where A is a lower triangular banded matrix.

Format

```
x = bandltsol(b, A);
```

banditsol

Input

b	$K \times M$ matrix.
A	$K \times N$ compact form matrix.

Output

x	$K \times M$ matrix.
-----	----------------------

Remarks

A is a lower triangular banded matrix in compact form. See **band** for a description of the format of A .

b can have more than one column. If so, $Ax = b$ is solved for each column. That is,

$$A * x[:, i] = b[:, i];$$

Example

```
//Create matrix 'A' and right-hand side 'b'
A = { 1 2 0 0,
      2 8 1 0,
      0 1 5 2,
      0 0 2 3 };
b = { 1.3, 2.1, 0.7, 1.8 };

//Create a matrix containing the lower triangular part
//of 'A'
Alower = lowmat(A);

//Create banded matrix from of 'Alower'
Abandlow = band(Alower, 1);

//Solve the system of equations
```



```
x = bandltsol(b, Abandlow);
```

After the code above:

```

      1  0  0  0      0  1      1.300      1.3
1.3
Alower = 2  8  0  0  Aband = 2  8  x = -0.063  b = 2.1
Alower*x = 2.1
      0  1  5  0      1  5      0.153      0.7
0.7
      0  0  2  3      2  3      0.498      1.8
1.8
```

See Also

[band](#), [bandchol](#), [bandcholsol](#), [bandrv](#), [bandsolpd](#)

bandrv

Purpose

Creates a symmetric banded matrix, given its compact form.

Format

```
y = bandrv(a);
```

Input

a KxN compact form matrix.

Output

y KxK symmetrix banded matrix.

bandrv

Remarks

a is the compact form of a symmetric banded matrix, as generated by **band**. *a* stores subdiagonals right to left, with the principal diagonal in the rightmost (Nth) column. The upper left corner of *a* is unused. **bandchol** expects a matrix of this form.

y is the fully expanded form of *a*, a KxK matrix with N-1 subdiagonals.

Example

b

```
x = { 1 2 0 0,
      2 8 1 0,
      0 1 5 2,
      0 0 2 3 };

//Create a version of 'x' in band format
xBand = band(x,1);

//Expand the banded version of 'x' back to a full matrix
xNew = bandrv(xBand);
```

After the code above:

	0	1		1	2	0	0		1	2	0	0	
xBand =	2	8		x =	2	8	1	0	xNew =	2	8	1	0
	1	5			0	1	5	2		0	1	5	2
	2	3			0	0	2	3		0	0	2	3

See Also

[band](#), [bandchol](#), [bandcholsol](#), [bandltsol](#), [bandsolpd](#)

bandsolpd

Purpose

Solves the system of equations $Ax = b$ for x , where A is a positive definite banded matrix.

Format

$x = \text{bandsolpd}(b, A);$

Input

b	KxM matrix.
A	KxN compact form matrix.

Output

x	KxM matrix.
-----	-------------

Remarks

A is a positive definite banded matrix in compact form. See **band** for a description of the format of A .

b can have more than one column. If so, $Ax = b$ is solved for each column. That is,

$$A * x[:, i] = b[:, i]$$

See Also

[band](#), [bandchol](#), [bandcholsol](#), [bandltsol](#), [bandrv](#)

bar

bar

Purpose

Generates a bar graph. NOTE: This function is for the deprecated PQG graphics, use **plotBar** instead.

Library

pgraph

Format

bar(*val*, *ht*);

Input

<i>val</i>	Nx1 numeric vector, bar labels. If scalar 0, a sequence from 1 to rows (<i>ht</i>) will be created.
<i>ht</i>	NxK numeric vector, bar heights.

Global Input

<i>_pbarwid</i>	scalar, width and type of bars in bar graphs and histograms. The valid range is 0-1. If this is 0, the bars will be a single pixel wide. If this is 1, the bars will touch each other.
<i>_pbartyp</i>	<p>If this value is positive, the bars will overlap. If negative, the bars will be plotted side-by-side. The default is 0.5.</p> <p>Kx2 matrix.</p> <p>The first column controls the bar shading:</p> <p>0 no shading.</p>

- | | |
|---|-------------------------------------|
| 1 | dots. |
| 2 | vertical cross-hatch. |
| 3 | diagonal lines with positive slope. |
| 4 | diagonal lines with negative slope. |
| 5 | diagonal cross-hatch. |
| 6 | solid. |

The second column controls the bar color.

Remarks

Use **scale** or **ytics** to fix the scaling for the bar heights.

Example

In this example, three overlapping sets of bars will be created. The three heights for the i th bar are stored in $x[i,:]$.

```
library pgraph;
graphset;

t = seqa(0,1,10);
x = (t^2/2) .* (1~0.7~0.3);

_plegctl = { 1 4 };
_plegstr = "Accnt #1\000Accnt #2\000Accnt #3";
title("Theoretical Savings Balance");
xlabel("Years");
ylabel("Dollars x 1000");
_pbartyp = { 1 10 }; /* Set color of the bars */
_pnum = 2;

bar(t,x); /* Use t vector to label X axis. */
```

base10

Source

pbar.src

See Also

[asclabel](#), [xy](#), [logx](#), [logy](#), [loglog](#), [scale](#), [hist](#)

base10

Purpose

Breaks number into a number of the form `#.####...` and a power of 10.

Format

$\{ M, P \} = \text{base10}(x);$

Input

x scalar, number to break down.

Output

M scalar, in the range $-10 < M < 10$.

P scalar, integer power such that:

$$M \cdot 10^P = x$$

Example

`{ b, e } = base10(4500);`

After the code above:

```
b = 4.5  e = 3
```

and

```
b*10^e = 4.5*10^3 = 4500
```

Source

base10.src

begwind

Purpose

Initializes global graphic panel variables. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
begwind;
```

Remarks

This procedure must be called before any other graphic panel functions are called.

Source

pwindow.src

See Also

[endwind](#), [window](#), [makewind](#), [nextwind](#), [getwind](#)

besselj

besselj

Purpose

Computes a Bessel function of the first kind, $J_n(x)$.

Format

$y = \text{besselj}(n, x);$

Input

n	NxK matrix or P-dimensional array where the last two dimensions are NxK, the order of the Bessel function. Nonintegers will be truncated to an integer.
x	LxM matrix or P-dimensional array where the last two dimensions are LxM, ExE conformable with n .

Output

y	max(N,L) by max(K,M) matrix or P-dimensional array where the last two dimensions are max(N,L) by max(K,M).
-----	--

Example

```
//Create the sequence 0.1, 0.2, 0.3,...,19.9
x = seqa(0, 0.1, 200);

//Calculate a first order Bessel function
ord = 1;
y0 = besselj(ord, x);
```



```
//Calculate the first and second order Bessel function
ord = { 1 2 };
y = besselj(ord, x);

//Plot the output of the first and third order Bessel
//functions
plotXY(x, y);
```

In the code above, the calculation of both the first and second order Bessel functions assigns the return from the first order calculation to be the first column of y and the return from the calculation of the second order function to be the second column of y .

The **plotXY** function treats each incoming column as a separate line.

See Also

[bessely](#), [mbesseli](#)

bessely

Purpose

Computes a Bessel function of the second kind (Weber's function), $Y_n(x)$.

Format

$y = \text{bessely}(n, x);$

Input

n	NxK matrix or P-dimensional array where the last two dimensions are NxK, the order of the Bessel function. Nonintegers will be truncated to an integer.
x	LxM matrix or P-dimensional array where the last two dimensions are LxM, ExE conformable with n .

bessely

Output

y	max(N,L) by max(K,M) matrix or P-dimensional array where the last two dimensions are max(N,L) by max(K,M).
-----	--

Example

```
//Create the sequence 0.1, 0.2, 0.3, 0.4, 0.5
x = seqa(0.1, 0.1, 5);

//Create the sequence 1, 1.1, 1.2, 1.3, 1.4
x2 = seqa(1, 0.1, 5);

//Calculate a first order bessel function against 'x' and
//calculate a third order bessel function against 'x2'
//NOTE: The '~' provides horizontal concatenation
ord = { 1 3 };
y = bessely(ord, x~x2);
```

After the code above:

	-6.459	-5.822		0.100	1.000
	-3.324	-4.507		0.200	1.100
y =	-2.293	-3.590	x~x2 =	0.300	1.200
	-1.781	-2.930		0.400	1.300
	-1.471	-2.442		0.500	1.400

See Also

[besseli](#), [mbesseli](#)

beta

Purpose

Computes the standard Beta function, also called the Euler integral. The beta function is defined as:

$$B(x, y) = \int_0^1 t^{x-1} (1-t)^{y-1} dt$$

Format

```
f = beta(x, y);
```

Input

<i>x</i>	scalar or NxK matrix; <i>x</i> may be real or complex.
<i>y</i>	LxM matrix, ExE conformable with <i>x</i> .

Output

<i>f</i>	NxK matrix.
----------	-------------

Technical Notes

The Beta function's relationship with the Gamma function is:

$$\frac{\text{gamma}(\mathbf{x}) \times \text{gamma}(\mathbf{y})}{\text{gamma}(\mathbf{x} + \mathbf{y})}$$

See Also

[cdfBeta](#), [gamma](#), [gammacplx](#), [zeta](#)

box

box

Purpose

Graphs data using the box graph percentile method. NOTE: This function uses the deprecated PQG graphics. Use **plotBox** instead.

Library

pgraph

Format

box(*grp*, *y*);

Input

<i>grp</i>	1xM vector. This contains the group numbers corresponding to each column of <i>y</i> data. If scalar 0, a sequence from 1 to cols (<i>y</i>) will be generated automatically for the X axis.
<i>y</i>	NxM matrix. Each column represents the set of <i>y</i> values for an individual percentiles box symbol.

Global Input

<i>_pboxctl</i>	5x1 vector, controls box style, width, and color.
[1]	box width between 0 and 1. If zero, the box plot is drawn as two vertical lines representing the quartile ranges with a filled circle representing the 50th percentile.
[2]	box color. If this is set to 0, the colors may be individually controlled using the global variable

_pcolor.

[3] Min/max style for the box symbol. One of the following:

- 1 Minimum and maximum taken from the actual limits of the data. Elements 4 and 5 are ignored.
- 2 Statistical standard with the minimum and maximum calculated according to interquartile range as follows:

$$intqrang = 75th - 25th$$

$$min = 25th - 1.5 \text{ } intqrang$$

$$max = 75th + 1.5 \text{ } intqrang$$
 Elements 4 and 5 are ignored.
- 3 Minimum and maximum percentiles taken from elements 4 and 5.

[4] Minimum percentile value (0-100) if *_pboxctl*[3] = 3.

[5] Maximum percentile value (0-100) if *_pboxctl*[3] = 3.

_plctrl 1xM vector or scalar as follows:

- 0 Plot boxes only, no symbols.
- 1 Plot boxes and plot symbols which lie outside the *min* and *max* box values.
- 2 Plot boxes and all symbols.
- 1 Plot symbols only, no boxes.

These capabilities are in addition to the usual line control capabilities of *_plctrl*.

boxcox

<code>_pcolor</code>	1xM vector or scalar for symbol colors. If scalar, all symbols will be one color.
----------------------	---

Remarks

If missing values are encountered in the <i>y</i> data, they will be ignored during calculations and will not be plotted.

Source

`pbox.src`

boxcox

Purpose

Computes the Box-Cox function.

Format

$y = \text{boxcox}(x, \lambda);$

Input

<i>x</i>	MxN matrix or P-dimensional array where the last two dimensions are MxN.
<i>lambda</i>	KxL matrix or P-dimensional array where the last two dimensions are KxL, ExE conformable to <i>x</i> .

Output

<i>y</i>	max(M,L)xmax(N,K) or P-dimensional array where the last two dimensions are max(M,L)xmax(N,K).
----------	---

Remarks

Allowable range for x is: $x > 0$

The **boxcox** function computes:

$$\text{boxcox}(x) = (x^\lambda - 1) / \lambda$$

Example

```
x = { .2, .4, .8, 1, 1.2, 1.4 };  
lambda = .4;  
y = boxcox(x, lambda);
```

After the code above:

```
-1.187  
-0.767  
y = -0.213  
0.000  
0.189  
0.360
```

break

Purpose

Breaks out of a **do** or **for** loop.

Format

```
break;
```

break

Example

```
x = randn(4,4);

//Loop through each row of 'x' using 'r' as the loop
//counter
for r(1, rows(x), 1);
    //For each row, loop through its elements
    for c(1, cols(x), 1);
        if c == r;      /* Set the diagonal to 1 */
            x[r,c] = 1;
        elseif c > r; /* leave upper triangle as it is */
            break;      /* terminate inner loop */
        else;
            x[r,c] = 0; /* set lower triangle elements to 0 */
        endif;
    endfor;             /* break jumps to the statement after
                        this endfor */
endfor;
```

After running the code above, `x` should be a lower triangular matrix similar to below. Due to the use of random data, your matrix will have different non-zero elements above the diagonal.

1.000	1.288	-0.060	1.801
0.000	1.000	1.609	1.474
0.000	0.000	1.000	-0.768
0.000	0.000	0.000	1.000

Remarks

This command works just like in C.

See Also

[continue](#), [do](#), [for](#)

c

call

Purpose

Calls a function or procedure when the returned value is not needed and can be ignored, or when the procedure is defined to return nothing.

Format

```
call function_name(argument_list);  
call function_name;
```

Remarks

This is useful when you need to execute a function or procedure and do not need the value that it returns. It can also be used for calling procedures that have been defined to return nothing.

function_name can be any intrinsic **GAUSS** function, a procedure (**proc**), or any valid expression.

Example

```
// Create a positive definite matrix  
x = moment(rndn(100,4),0);  
// Call chol function  
call chol(x);  
// y is the determinant
```

cdfBeta

```
y = det1;
```

The above example is the fastest way to compute the determinant of a positive definite matrix. The result of **chol** is discarded and **det1** is used to retrieve the determinant that was computed during the call to **chol**.

See Also

[proc](#)

cdfBeta

Purpose

Computes the incomplete Beta function (i.e., the cumulative distribution function of the Beta distribution).

Format

```
y = cdfBeta(x, a, b);
```

Input

<i>x</i>	NxK matrix.
<i>a</i>	LxM matrix, ExE conformable with <i>x</i> .
<i>b</i>	PxQ matrix, ExE conformable with <i>x</i> and <i>a</i> .

Output

<i>y</i>	max(N,L,P) by max(K,M,Q) matrix.
----------	----------------------------------

Remarks

y is the integral from 0 to *x* of the beta distribution with parameters *a* and *b*.

Allowable ranges for the arguments are:

$$\begin{aligned} 0 &\leq x \leq 1 \\ a &> 0 \\ b &> 0 \end{aligned}$$

A -1 is returned for those elements with invalid inputs.

Example

```
x = { .1, .2, .3, .4 };
a = 0.5;
b = 0.3;
y = cdfBeta(x,a,b);
print "y = "      y;
```

After running above code,

```
y =
    0.14228251
    0.20662575
    0.26057158
    0.31087052
```

See Also

[cdfChic](#), [cdfFc](#), [cdfN](#), [cdfNc](#), [cdfTc](#), [gamma](#)

Technical Notes

cdfBeta has the following approximate accuracy:

		$\max(a, b)$	\leq	500	absolute error is approx. $\pm 5e-13$
500	$<$	$\max(a, b)$	\leq	10,000	absolute error is

cdfBetaInv

10,000

<

b)

max(a,

b)

<=

200,000

approx. ±5e-11

absolute error is

approx. ±1e-9

References

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men. *Theory of Probability and its Applications*. Vol. 8, No. 2, 1963, 129-55.

2. Boston N.E. and E.L. Battiste. "Remark on Algorithm 179 Incomplete Beta
Ratio." *Comm. ACM*. Vol. 17, No. 3, March 1974, 156-57.

3. Ludwig, O.G. "Algorithm 179 Incomplete Beta Ratio." *Comm. ACM*. Vol. 6, No.
6, June 1963, 314.

4. Mardia, K.V. and P.J. Zemroch. *Tables of the F- and related distributions with
algorithms*. Academic Press, New York, 1978. ISBN 0-12-471140-5.

5. Peizer, D.B. and J.W. Pratt. "A Normal Approximation for Binomial, F, Beta,
and Other Common, Related Tail Probabilities, I." *Journal of the American Stat-
istical Association*. Vol. 63, Dec. 1968, 1416-56.

6. Pike, M.C. and J.W. Pratt. "Remark on Algorithm 179 Incomplete Beta Ratio."
Comm. ACM. Vol. 10, No. 6, June 1967, 375-76.

cdfBetaInv

Purpose

Computes the quantile or inverse of the Beta cumulative distribution function.

Format

$x = \text{cdfBetaInv}(p, a, b);$

Input

p

a

NxK matrix, Nx1 vector or scalar. $0 < p < 1$.

ExE conformable with p . $0 < a$.

b ExE conformable with p . $0 < b$.

Output

x NxK matrix, Nx1 vector or scalar.

Remarks

For invalid inputs, **cdfBetaInv** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

Example

```
y = { 0.14228251, 0.20662575, 0.26057158, 0.31087052 };
a = 0.5;
b = 0.3;
p = cdfBeta(y,a,b);
print "p = "    p;
```

After running above code,

```
p =
    0.1
    0.2
    0.3
    0.4
```

See Also

[cdfBeta](#), [cdfBinomial](#), [cdfNegBinomial](#)

cdfBinomial

cdfBinomial

Purpose

Computes the binomial cumulative distribution function.

Format

```
p = cdfBinomial(successes, trials, prob);
```

Input

<i>successes</i>	NxK matrix, Nx1 vector or scalar. <i>successes</i> must be a positive number and < trials
<i>trials</i>	ExE conformable with <i>successes</i> . <i>trials</i> must be > <i>successes</i> .
<i>prob</i>	The probability of success on any given trial. ExE conformable with <i>successes</i> . $0 < \text{prob} < 1$.

Output

<i>p</i>	NxK matrix, Nx1 vector or scalar.
----------	-----------------------------------

Example

What are the chances that a baseball player with a long-term batting average of .317 could break Ichiro Suzuki's record of 270 hits in a season if he had as many at bats as Ichiro had that year, 704?

```
// The cumulative probability of our player  
//getting 270 or fewer hits in the season
```

```
p = cdfBinomial(270,704,.317);
p = 0.9999199430052614
```

Therefore the odds of this player breaking Ichiro's record:

```
1-p = 0.0000000000037863 or 0.0000000003786305%
```

Remarks

$$P(\mathbf{x} \leq \mathbf{k}) = \sum_{i=0}^k \binom{n}{i} p^i (1-p)^{n-i}$$

For invalid inputs, **cdfBinomial** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[cdfBinomialInv](#), [cdfNegBinomial](#), [pdfBinomial](#)

cdfBinomialInv

Purpose

Computes the binomial quantile or inverse cumulative distribution function.

Format

```
s = cdfBinomialInv(p, trials, prob);
```

cdfBinomialInv

Input

<i>p</i>	NxK matrix, Nx1 vector or scalar. $0 < p < 1$.
<i>trials</i>	ExE conformable with <i>p</i> . $trials > 0$.
<i>prob</i>	The probability of success on any given trial. ExE conformable with <i>p</i> . $0 < prob < 1$.

Output

<i>s</i>	The number of successes. NxK matrix, Nx1 vector or scalar.
----------	--

Example

What is a reasonable range of wins for a basketball team playing 82 games in a season, with a 60% chance of winning any game?

For our example we will define a reasonable range as falling between the top and bottom deciles.

```
range = { .10, .9 };  
s = cdfBinomialInv(range, 82, .6);  
print "s = " s;
```

After above code,

```
s =  
    43  
    55
```

This means that a team with a 60% chance of winning any one game would win between 43 and 55 games in 80% of seasons.

Remarks

For invalid inputs, `cdfBinomialInv` will return a scalar error code which, when its value is assessed by function `scalerr`, corresponds to the invalid input. If the first input is out of range, `scalerr` will return a 1; if the second is out of range, `scalerr` will return a 2; etc.

See Also

[cdfBinomial](#), [pdfBinomial](#), [cdfNegBinomial](#), [cdfNegBinomialInv](#)

cdfBvn

Purpose

Computes the cumulative distribution function of the standardized bivariate Normal density (lower tail).

Format

$c = \text{cdfBvn}(h, k, r);$

Input

h	NxK matrix, the upper limits of integration for variable 1.
k	LxM matrix, ExE conformable with h , the upper limits of integration for variable 2.
r	PxQ matrix, ExE conformable with h and k , the correlation coefficients between the two variables.

cdfBvn

Output

C

max(N,L,P) by max(K,M,Q) matrix, the result of the double integral from $-\infty$ to h and $-\infty$ to k of the standardized bivariate Normal density $f(x, y, r)$.

Remarks

The function integrated is:

$$f(x, y, r) = \frac{e^{-0.5w}}{2\pi\sqrt{1-r^2}}$$

with

$$w = \frac{x^2 - 2rxy + y^2}{1-r^2}$$

Thus, x and y have 0 means, unit variances, and correlation = r .

Allowable ranges for the arguments are:

$$-\infty \leq h \leq +\infty$$

$$-\infty \leq k \leq +\infty$$

$$-1 < r < 1$$

A -1 is returned for those elements with invalid inputs.

To find the integral under a general bivariate density, with x and y having nonzero means and any positive standard deviations, use the transformation equations:

$$h = (ht - ux) ./ sx;$$

$$k = (kt - uy)$$

where ux and uy are the (vectors of) means of x and y , sx and sy are the (vectors of) standard deviations of x and y , and ht and kt are the (vectors of) upper integration limits for the untransformed variables, respectively.

See Also

[cdfN](#), [cdfTvn](#)

Technical Notes

The absolute error for **cdfBvn** is approximately $\pm 5.0e-9$ for the entire range of arguments.

References

1. Daley, D.J. "Computation of Bi- and Tri-variate Normal Integral." *Appl. Statist.* Vol. 23, No. 3, 1974, 435-38.

2. Owen, D.B. "A Table of Normal Integrals." *Commun. Statist.-Simula. Computa.*, B9(4). 1980, 389-419.

cdfBvn2

Purpose

Returns the bivariate Normal cumulative distribution function of a bounded rectangle.

Format

$y = \text{cdfBvn2}(h, \quad dh, \quad k, \quad dk, r);$

Input

<i>h</i>	Nx1 vector, starting points of integration for variable 1.
<i>dh</i>	Nx1 vector, increments for variable 1.
<i>k</i>	Nx1 vector, starting points of integration for variable 2.
<i>dk</i>	Nx1 vector, increments for variable 2.

cdfBvn2

r

Nx1 vector, correlation coefficients between the two variables.

Output

y

Nx1 vector, the integral over the rectangle bounded by h , $h + dh$, k , and $k + dk$ of the standardized bivariate Normal distribution.

Remarks

Scalar input arguments are okay; they will be expanded to Nx1 vectors.

cdfBvn2 computes:

$$\text{cdfBvn}(h + dh, k + dk, r) + \text{cdfBvn}(h, k, r) - \text{cdfBvn}(h, k + dk, r) - \text{cdfBvn}(h + dh, k, r)$$

cdfBvn2 computes an error estimate for each set of inputs. The size of the error depends on the input arguments. If **trap 2** is set, a warning message is displayed when the error reaches $0.01 * \text{abs}(y)$. For an estimate of the actual error, see **cdfBvn2e**.

Example

Example 1

```
print cdfBvn2(1, -1, 1, -1, 0.5);  
1.4105101488974692e-001
```

Example 2

```
print cdfBvn2(1, -1e-15, 1, -1e-15, 0.5);  
4.9303806576313238e-32
```

Example 3

```
print cdfBvn2(1,-1e-45,1,-1e-45,0.5);
0.0000000000000000e+000
```

Example 4

```
trap 2,2;
print cdfBvn2(1,-1e-45,1,1e-45,0.5);

WARNING: Dubious accuracy from cdfBvn2:
0.000e+000 +/- 2.8e-060
0.0000000000000000e+000
```

Source

lncdfn.src

See Also

[cdfBvn2e](#), [lncdfbvn2](#)

cdfBvn2e

Purpose

Returns the bivariate Normal cumulative distribution function of a bounded rectangle.

Format

```
{ y, e } = cdfBvn2e(h, dh, k, dk, r);
```

Input

h Nx1 vector, starting points of integration for variable 1.

cdfBvn2e

dh	Nx1 vector, increments for variable 1.
k	Nx1 vector, starting points of integration for variable 2.
dk	Nx1 vector, increments for variable 2.
r	Nx1 vector, correlation coefficients between the two variables.

Output

y	Nx1 vector, the integral over the rectangle bounded by h , $h + dh$, k , and $k + dk$ of the standardized bivariate Normal distribution.
e	Nx1 vector, an error estimate.

Remarks

Scalar input arguments are okay; they will be expanded to Nx1 vectors. **cdfBvn2e** computes:

$$\mathbf{cdfBvn}(h + dh, k + dk, r) + \mathbf{cdfBvn}(h, k, r) - \mathbf{cdfBvn}(h, k + dk, r) - \mathbf{cdfBvn}(h + dh, k, r)$$

The real answer is $y \pm e$. The size of the error depends on the input arguments.

Example

Example 1

```
print
cdfBvn2e(1,-1,1,-1,0.5);

1.4105101488974692e-001
1.9927918166193113e-014
```

Example 2

```
print
cdfBvn2e(1,-1e-15,1,-1e-15,0.5);

7.3955709864469857e-032
2.8306169312687801e-030
```

Example 3

```
print
cdfBvn2e(1,-1e-45,1,-1e-45,0.5);

0.0000000000000000e+000
2.8306169312687770e-060
```

See Also

[cdfBvn2](#), [lncdfbvn2](#)

cdfCauchy

Purpose

Computes the cumulative distribution function for the Cauchy distribution.

Format

$y = \text{cdfCauchy}(x, a, b);$

Input

x	NxK matrix, an Nx1 vector or scalar.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x .

cdfCauchyInv

b	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . b must be greater than 0.
-----	---

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

The cumulative distribution function for the Cauchy distribution is defined as:

$$\frac{1}{2} + \frac{1}{\pi} \arctan\left(\frac{x-a}{b}\right)$$

See Also

[pdfCauchy](#)

cdfCauchyInv

Purpose

Computes the Cauchy inverse cumulative distribution function.

Format

$y = \text{cdfCauchyInv}(p, a, b);$

Input

p	NxK matrix, Nx1 vector or scalar. p must be greater than zero and less than 1.
a	Location parameter; NxK matrix, Nx1 vector or scalar,

b	ExE conformable with p . Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with p . b must be greater than 0.
-----	--

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

See Also

[pdfCauchy](#), [cdfCauchy](#)

cdfChic

Purpose

Computes the complement of the cdf of the chi-square distribution.

Format

```
 $y = \text{cdfChic}(x, n);$ 
```

Input

x	NxK matrix.
n	LxM matrix, ExE conformable with x .

Output

y	$\max(N,L)$ by $\max(K,M)$ matrix.
-----	------------------------------------

cdfChic

Remarks

y is the integral from x to ∞ of the chi-square distribution with n degrees of freedom.

The elements of n must all be positive integers. The allowable ranges for the arguments are:

```
x > 0
n > 0
```

A -1 is returned for those elements with invalid inputs.

This equals $1 - X_n^2(x)$. Thus, to get the chi-squared cdf, subtract **cdfChic**(x , n) from 1. The complement of the cdf is computed because this is what is most commonly needed in statistical applications, and because it can be computed with fewer problems of roundoff error.

Example

```
x = { .1, .2, .3, .4 };
n = 3;
y = cdfChic(x,n);
print "y = " y;
```

After running above code

```
0.991837
y = 0.977589
0.960028
0.940242
```

See Also

[cdfBeta](#), [cdfFc](#), [cdfNc](#), [cdfTc](#), [gamma](#)

Technical Notes

For $n \leq 1000$, the incomplete gamma function is used and the absolute error is approx. $\pm 6e-13$.

For $n > 1000$, a Normal approximation is used and the absolute error is $\pm 2e-8$.

For higher accuracy when $n > 1000$, use:

```
1 - cdfGam(0.5*x, 0.5*n);
```

References

1. Bhattacharjee, G.P. "Algorithm AS 32, the Incomplete Gamma Integral." *Applied Statistics*. Vol. 19, 1970, 285-87.
2. Mardia K.V. and P.J. Zemroch. *Tables of the F- and related distributions with algorithms*. Academic Press, New York, 1978. ISBN 0-12-471140-5.
3. Peizer, D.B. and J.W. Pratt. "A Normal Approximation for Binomial, F, Beta, and other Common, Related Tail Probabilities, I." *Journal of the American Statistical Association*. Vol. 63, Dec. 1968, 1416-56.

cdfChii

Purpose

Compute chi-square abscissae values given probability and degrees of freedom.

Format

```
c = cdfChii(p, n);
```

Input

p	MxN matrix, probabilities.
n	LxK matrix, ExE conformable with p , degrees of

cdfChii

freedom.

Output

C

max(M,L) by max(N,K) matrix, abscissae values for chi-squared distribution.

Example

The following generates a 3x3 matrix of pseudo-random numbers with a chi-squared distribution with expected value of 4:

```
//Set the rng seed for repeatable random numbers
rndseed 464578;

//Set the 'probabilities' input equal to a 3x3 matrix of
//uniform random numbers and the degrees of freedom' input
//to be a 3x3 matrix with each element equal to '4'
x = cdfChii(rndu(3,3),4+zeros(3,3));
```

After the code above:

```
0.934227 6.231914 4.227479
x = 2.647158 1.203957 10.559593
5.868060 1.368600 1.963283
```

Source

cdfchii.src

See Also

[gammaii](#)

C

cdfChinc

Purpose

Computes the cumulative distribution function for the noncentral chi-square distribution.

Format

```
 $y = \text{cdfChinc}(x, \nu, d);$ 
```

Input

x	Nx1 vector, values of upper limits of integrals, must be greater than 0.
ν	scalar, degrees of freedom, $\nu > 0$.
d	scalar, noncentrality parameter, $d > 0$.

This is the square root of the noncentrality parameter that sometimes goes under the symbol lambda. (See Scheffe, *The Analysis of Variance*, App. IV, 1959.)

Output

y	Nx1 vector.
-----	-------------

Remarks

y is the integral from 0 to x of the noncentral chi-square distribution with ν degrees of freedom and noncentrality d .

cdfChinc can return a vector of values, but the degrees of freedom and noncentrality parameter must be the same for all values of x .

cdfChincInv

For invalid inputs, **cdfChinc** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

Relation to **cdfChic**:

```
cdfChic(x, v) = 1 - cdfChinc(x, v, 0);
```

Example

```
x = { .5, 1, 5, 25 };  
print cdfChinc(x, 4, 2);
```

The code above returns:

```
0.0042086234  
0.016608592  
0.30954232  
0.99441140
```

See Also

[cdfFnc](#), [cdfTnc](#)

cdfChincInv

Purpose

Computes the quantile or inverse of noncentral chi-square cumulative distribution function.

Format

```
 $x = \text{cdfChincInv}(y, df, \text{nonc});$ 
```

Input

y	NxK matrix, Nx1 vector or scalar. The integral from 0 to x .
df	ExE conformable with y . The degrees of freedom. $df > 0$.
nonc	ExE conformable with y . The noncentrality parameter. Note: This is the <u>square root of the noncentrality parameter</u> that sometimes goes under the symbol λ . $\text{nonc} > 0$.

Output

x	NxK matrix, Nx1 vector or scalar. The upper limit of the integrals of the noncentral chi-square distribution with df degrees of freedom and noncentrality nonc .
-----	---

Remarks

Note: Input nonc is the square root of the noncentrality parameter that sometimes goes under the symbol λ .

For invalid inputs, **cdfChincInv** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[cdfChinc](#), [cdfChic](#), [cdfFnc](#), [cdfTnc](#)

cdfEmpirical

cdfEmpirical

Purpose

Computes the cumulative distribution function (cdf) of an empirical distribution.

Format

```
 $f$  = cdfEmpirical( $x$ );
```

Input

x	N x 1 vector.
-----	---------------

Output

f	(N+1) x 1 vector.
-----	-------------------

Remarks

f is the integral from $-\infty$ to x of the empirical distribution function.

Example

```
new ;  
cls ;  
rndseed 2223;  
  
// Create a random vector  
x = rndn(30,1) ;  
  
// Sort x for the first column  
x = sortc(x,1) ;  
  
// Get empirical cdf of x
```



```

f = cdfEmpirical(x);

// Add negative infinity (___INFN) for probability equal to
0.
print (___INFN|x)~f;

// Plot empirical distribution
plotCDFEmpirical(x);

// Get normal cdf of x
fnormal = cdfN(x);

// Plot theoretical distribution
plotAddXY(x, fnormal);

```

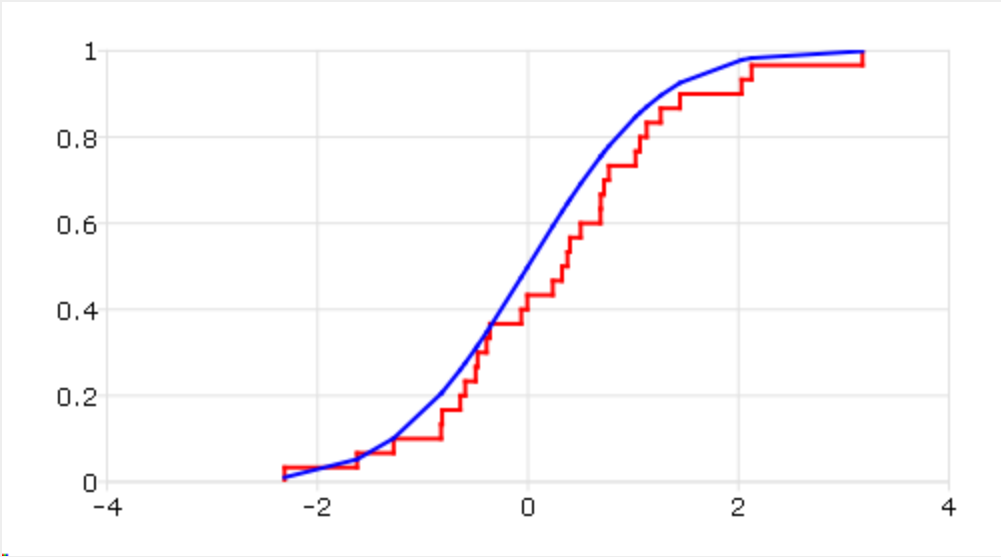
After above code,

-INF	0.00000000
-2.3124206	0.03333333
-1.6240227	0.06666667
-1.2763153	0.10000000
-0.82532512	0.13333333
-0.81574278	0.16666667
-0.64338729	0.20000000
-0.59625173	0.23333333
-0.49725006	0.26666667
-0.47855430	0.30000000
-0.39340284	0.33333333
-0.36201638	0.36666667
-0.063830011	0.40000000
-0.0064523646	0.43333333
0.23570074	0.46666667
0.32355136	0.50000000
0.37501508	0.53333333

cdfEmpirical

0.39847826	0.56666667
0.50039685	0.60000000
0.68900341	0.63333333
0.69132515	0.66666667
0.72246796	0.70000000
0.76893134	0.73333333
1.0221019	0.76666667
1.0638924	0.80000000
1.1274880	0.83333333
1.2610791	0.86666667
1.4445086	0.90000000
2.0295113	0.93333333
2.1240430	0.96666667
3.1784008	1.00000000

The plot is



See Also

[plotCDFEmpirical](#)

cdfExp

Purpose

Computes the cumulative distribution function for the exponential distribution.

Format

$$y = \text{cdfExp}(x, a, m);$$

Input

x	NxK matrix, an Nx1 vector or scalar.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . a must be less than x .
m	Mean parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . m must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

The cumulative distribution function for the exponential distribution is defined as

$$1 - \exp\left(-\frac{x-a}{b}\right)$$

See Also

[pdfExp](#)

cdfExpInv

cdfExpInv

Purpose

Computes the exponential inverse cumulative distribution function.

Format

$y = \text{cdfExpInv}(p, a, b);$

Input

p	NxK matrix, Nx1 vector or scalar. p must be greater than zero and less than 1.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with p .
b	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with p . b must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

See Also

[pdfExp](#), [cdfExp](#)

cdfFc

Purpose

Computes the complement of the cumulative distribution function of the F distribution.

Format

```
y = cdfFc(x, n1, n2);
```

Input

<i>x</i>	NxK matrix.
<i>n1</i>	LxM matrix, ExE conformable with <i>x</i> .
<i>n2</i>	PxQ matrix, ExE conformable with <i>x</i> and <i>n1</i> .

Output

<i>y</i>	max(N,L,P) by max(K,M,Q) matrix
----------	---------------------------------

Example

cdfFc can be used to calculate a p-value from an F-statistic.

```
n_obs = 100;  
n_vars = 5;  
f_stat = 2.4;  
p_value = cdfFc(f_stat, n_vars, n_obs - n_vars - 1);  
print p_value;
```

will return:

```
0.042803132
```

Remarks

y is the integral from *x* to ∞ of the *F* distribution with *n1* and *n2* degrees of freedom.

cdfFc

This equals

$$1 - G(x, n1, n2)$$

where G is the F cdf with $n1$ and $n2$ degrees of freedom. Thus, to get the F cdf, use:

$$1 - \text{cdfFc}(x, n1, n2);$$

The complement of the cdf is computed because this is what is most commonly needed in statistical applications, and because it can be computed with fewer problems of round-off error.

Allowable ranges for the arguments are:

$$\begin{aligned} x &> 0 \\ n1 &> 0 \\ n2 &> 0 \end{aligned}$$

A -1 is returned for those elements with invalid inputs.

For $\max(n1, n2) \leq 1000$, the absolute error is approx. $\pm 5e-13$. For $\max(n1, n2) > 1000$, Normal approximations are used and the absolute error is approx. $\pm 2e-6$.

For higher accuracy when $\max(n1, n2) > 1000$, use

$$\text{cdfBeta}(n2 / (n2 + n1 * x), n2/2, n1/2);$$

See Also

[cdfBeta](#), [cdfChic](#), [cdfN](#), [cdfNc](#), [cdfTc](#), [gamma](#)

References

1. Bol'shev, L.N. "Asymptotically Perason's Transformations." Teor. Veroyat. Pri-men. *Theory of Probability and its Applications*. Vol. 8, No. 2, 1963, 129-55.
2. Bosten, N.E. and E.L. Battiste. "Remark on Algorithm 179 Incomplete Beta

Ratio." *Comm. ACM*. Vol. 17, No. 3, March 1974, 156-57.

3. Kennedy, W.J., Jr. and J.E. Gentle. *Statistical Computing*. Marcel Dekker, Inc., New York, 1980.
4. Ludwig, O.G. "Algorithm 179 Incomplete Beta Ratio." *Comm. ACM*. Vol. 6, No. 6, June 1963, 314.
5. Mardia, K.V. and P.J. Zemroch. *Tables of the F- and related distributions with algorithms*. Academic Press, New York, 1978. ISBN 0-12-471140-5.
6. Peizer, D.B. and J.W. Pratt. "A Normal Approximation for Binomial, F, Beta, and other Common, Related Tail Probabilities, I." *Journal of the American Statistical Association*. Vol. 63, Dec. 1968, 1416-56.
7. Pike, M.C. and I.D. Hill, "Remark on Algorithm 179 Incomplete Beta Ratio." *Comm. ACM*. Vol. 10, No. 6, June 1967, 375-76.

cdfFnc

Purpose

Computes the cumulative distribution function of the noncentral F distribution.

Format

```
y = cdfFnc(x, n1, n2, d);
```

Input

x	$N \times 1$ vector, values of upper limits of integrals, $x > 0$.
$v1$	scalar, degrees of freedom of numerator, $n1 > 0$.
$v2$	scalar, degrees of freedom of denominator, $n2 > 0$.
d	scalar, noncentrality parameter, $d > 0$.

This is the square root of the noncentrality parameter that

cdfFncInv

sometimes goes under the symbol λ . (See Scheffe, *The Analysis of Variance*, App. IV, 1959.)

Output

y Nx1 vector.

Remarks

For invalid inputs, **cdfFnc** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

Technical Notes

Relation to cdfFc:

$$\text{cdfFc}(x, n1, n2) = 1 - \text{cdfFnc}(x, n1, n2, 0);$$

See Also

[cdfTnc](#), [cdfChinc](#)

cdfFncInv

Purpose

Computes the quantile or inverse of noncentral F cumulative distribution function.

Format

$x = \text{cdfFncInv}(y, \text{dfn}, \text{dfd}, \text{nonc});$

Input

y	NxK matrix, Nx1 vector or scalar.
dfn	ExE conformable with y . The degrees of freedom numerator. $dfn > 0$.
dfd	ExE conformable with y . The degrees of freedom denominator. $dfd > 0$.
$nonc$	ExE conformable with y . The noncentrality parameter. Note: This is the <u>square root of the noncentrality parameter</u> that sometimes goes under the symbol lambda. $nonc > 0$.

Output

x	NxK matrix, Nx1 vector or scalar. The upper limit of the integrals of the noncentral F distribution.
-----	--

Remarks

Note: Input $nonc$ is the square root of the noncentrality parameter that sometimes goes under the symbol lambda.

For invalid inputs, **cdfFncInv** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[cdfFnc](#), [cdfChinc](#), [cdfChic](#), [cdfTnc](#)

cdfGam

cdfGam

Purpose

Computes the incomplete gamma function.

Format

```
g = cdfGam(x, intlim);
```

Input

<i>x</i>	NxK matrix of data.
<i>intlim</i>	LxM matrix, ExE compatible with <i>x</i> , containing the integration limit.

Output

<i>g</i>	max(N,L) by max(K,M) matrix.
----------	------------------------------

Remarks

The incomplete gamma function returns the integral

$$\int_0^{int\ lim} \frac{e^{-t} t^{(x-1)}}{\gamma(x)} dt$$

The allowable ranges for the arguments are:

```
x > 0  
intlim > 0
```

A -1 is returned for those elements with invalid inputs.

Example

```
x = { 0.5 1 3 10 };
intlrim = sega(0, .2, 6);
g = cdfGam(x, intlrim);
print "intlrim = " intlrim;
print "g = " g;
```

After the code above:

```
intlrim =
0.00000000
0.20000000
0.40000000
0.60000000
0.80000000
1.00000000

g =
0.00000000      0.00000000      0.00000000
0.00000000
0.47291074      0.18126925      0.0011484812      2.3530688e-
014
0.62890663      0.32967995      0.0079263319      2.0098099e-
011
0.72667832      0.45118836      0.023115288      9.6697183e-
010
0.79409679      0.55067104      0.047422596      1.4331002e-
008
0.84270079      0.63212056      0.080301397      1.1142548e-
007
```

cdfGenPareto

This computes the integrals over the range from 0 to 1, in increments of 0.2, at the parameter values 0.5, 1, 3, 10.

Technical Notes

cdfGam has the following approximate accuracy:

```

          x < 500      : the absolute error is approx. ±6e-
13
    500 <= x <= 10,000 : the absolute error is approx. ±3e-
11
10,000 < x           : a Normal approximation is used and
                        the absolute error is approx. ±3e-
10
```

References

1. Bhattacharjee, G.P. "Algorithm AS 32, the Incomplete Gamma Integral." *Applied Statistics*. Vol. 19, 1970, 285-87.
2. Mardia, K.V. and P.J. Zemroch. *Tables of the F- and Related Distributions with Algorithms*. Academic Press, New York, 1978. ISBN 0-12-471140-5.
3. Peizer, D.B. and J.W. Pratt. "A Normal Approximation for Binomial, F, Beta, and other Common, Related Tail Probabilities, I." *Journal of the American Statistical Association*. Vol. 63, Dec. 1968, 1416-56.

cdfGenPareto

Purpose

Computes the cumulative distribution function for the Generalized Pareto distribution.

Format

```
y = cdfGenPareto(x,a,o,k);
```

Input

x	NxK matrix, an Nx1 vector or scalar.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x.
o	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x. o must be greater than 0.
k	Shape parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x.

Output

y	NxK matrix, Nx1 vector or scalar.
---	-----------------------------------

Remarks

The cumulative distribution function for the Generalized Pareto distribution is defined as:

$$f(x) = \begin{cases} \frac{1 - \left(1 + k \frac{(x-\mu)}{\sigma}\right)^{-1/k}}{1 - \exp\left(-\frac{(x-\mu)}{\sigma}\right)} & k \neq 0 \\ \frac{(x-\mu)}{\sigma} & k = 0 \end{cases}$$

See Also

[pdfGenPareto](#)

cdfHyperGeo

cdfHyperGeo

Purpose

Computes the cumulative distribution function for the hypergeometric distribution.

Format

$p = \text{cdfHyperGeo}(x, m, k, n);$

Input

x	NxK matrix, Nx1 vector or scalar. x must be a positive number and $< m$
m	The size of the population from which draws will be made. ExE conformable with x . m must be $> x$, k and n .
k	The number of marked items. ExE conformable with x . $0 < \text{prob} < 1$.
n	The number of items drawn from the population. ExE conformable with x . $0 < k < m$.

Output

p	The probability of drawing x or fewer marked items. NxK matrix, Nx1 vector or scalar.
-----	---

Example

You are given 120 hard drives, 14 of which are known to be bad. What is the probability of drawing 2 or fewer bad hard drive if you randomly select 12 drives?

```
p = cdfHyperGeo(2, 120, 14, 12);
```

After running the code above, p is equal to:

```
0.85284036
```

Continuing with the example above, what are the probabilities of drawing 4 or fewer bad hard drives if you draw 20 or 40 hard drives?

```
x = 4;  
n_total = 120;  
n_bad = 14;  
n_draw = { 20, 40 };  
p = cdfHyperGeo(x, n_total, n_bad, n_draw);  
print p;
```

After running the code above, p is equal to:

```
0.94307042  
0.47070798
```

Remarks

For invalid inputs, **cdfHyperGeo** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[pdfHyperGeo](#), [rndHyperGeo](#), [cdfBinomial](#)

cdfLaplace

cdfLaplace

Purpose

Computes the cumulative distribution function for the Laplace distribution.

Format

$y = \text{cdfLaplace}(x, a, b);$

Input

x	NxK matrix, an Nx1 vector or scalar.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x .
b	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . b must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

The cumulative distribution function for the Laplace distribution is defined as

$$F(x) = \begin{cases} \frac{1}{2} \exp(-\lambda(\mu - x)) & X \leq \mu \\ 1 - \frac{1}{2} \exp(-\lambda(\mu - x)) & X > \mu \end{cases}$$

See Also

[cdfLaplaceInv](#)

cdfLaplaceInv

Purpose

Computes the Laplace inverse cumulative distribution function.

Format

$y = \text{cdfLaplaceInv}(p, a, b);$

Input

p	NxK matrix, Nx1 vector or scalar. p must be greater than 0 and less than 1.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with p .
b	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with p . b must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

See Also

[cdfLaplace](#)

cdfLogistic

Purpose

Computes the cumulative distribution function for the logistic distribution.

cdfLogistic

Format

```
y = cdfLogistic(x, a, b);
```

Input

x	NxK matrix, an Nx1 vector or scalar.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x.
b	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x. b must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
---	-----------------------------------

Remarks

The cumulative distribution function for the logistic distribution is defined as:

$$F(x) = \frac{1}{1 + \exp(-z)}$$

where

$$z \equiv \frac{x - \mu}{\sigma}$$

See Also

[pdfLogistic](#)

cdfLogisticInv

Purpose

Computes the logistic inverse cumulative distribution function.

Format

```
 $y = \text{cdfLogisticInv}(p, a, b);$ 
```

Input

p	NxK matrix, Nx1 vector or scalar. p must be greater than 0 and less than 1.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with p .
b	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with p . b must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

See Also

[pdfLogistic](#), [cdfLogistic](#)

cdfMvn

Purpose

Computes multivariate Normal cumulative distribution function.

cdfMvn

Format

```
 $p = \text{cdfMvn}(\text{ulim}, \text{corr});$ 
```

Input

<i>ulim</i>	K x 1 vector or K x N matrix, abscissae. If <i>ulim</i> has more than one column, each column will be treated as a separate set of upper limits.
<i>corr</i>	K x K matrix, correlation matrix.

Output

<i>p</i>	N x 1 vector, $P(\mathbf{X} \leq \text{ulim} \text{corr})$. <i>p</i> will have as many elements as the input, <i>ulim</i> , has columns.
----------	---

Examples

Example 1: Uncorrelated variables

```
//Upper limits of integration
ulim = { 0, 0 };

//Identity matrix, indicates
//zero correlation between variables
corr = { 1 0,
        0 1 };

//Calculate cumulative probability of
//both variables being ≤ 0
p = cdfmvn(ulim, corr);

//Calculate joint probability of two
//variables with zero correlation,
```

```
//both, being  $\leq 0$   
p2 = cdfn(0) .* cdfn(0);
```

After the above code, both p and $p2$ should be equal to 0.25.

Example 2

```
//Upper limits of integration  
ulim = { -0.5, 1 };  
  
//Correlation matrix  
corr = { 1 0.26,  
         0.26 1 };  
  
//Calculate cumulative probability of  
//the first variable being  $\leq -0.5$   
//and the second variable being  $\leq 1$   
p = cdfmvn(ulim, corr);
```

After the above code, p should equal: 0.28025. It means :

$\Phi = P(-\infty < X_1 \leq -0.5, -\infty < X_2 \leq 1) \approx 0.28025$ when the correlation between two variables is 0.26.

Example 3: Compute the cdf at 3 separate pairs of points

```
//Upper limits of integration  
// $x_1 \leq -1$  and  $x_2 \leq -1.1$   
// $x_1 \leq 0$  and  $x_2 \leq 0.1$   
// $x_1 \leq 1$  and  $x_2 \leq 1.1$   
ulim = { -1 0 1,  
         -1.1 0.1 1.1 };  
  
//Correlation matrix  
corr = { 1 0.31,  
         0.31 1 };
```

cdfMvn

```
//Calculate cumulative probability of
//each pair of upper limits
p = cdfmvn(ulim, corr);
```

After the above code, p should equal:

```
0.040741382    0.31981965    0.74642007
```

which means that:

$$P(x_1 \leq -1 \text{ and } x_2 \leq -1.1) = 0.0407$$

$$P(x_1 \leq +0 \text{ and } x_2 \leq +0.1) = 0.3198$$

$$P(x_1 \leq 1 \text{ and } x_2 \leq 1.1) = 0.7464$$

Remarks

- **cdfMvn** evaluates the *MVN* integral with i -th row of \mathbf{x} (upper limits), where $1 \leq i \leq N$

$$\Phi(\mathbf{x}_i, \Sigma) = \frac{1}{\sqrt{|\Sigma|(2\pi)^K}} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \dots \int_{-\infty}^{x_{iK}} e^{-\frac{1}{2}\mathbf{z}'\Sigma^{-1}\mathbf{z}} d\mathbf{z}$$

where \mathbf{z} denotes K -dimensional multivariate normal distribution.

- The correlation matrix \mathbf{R} is defined by $\Sigma = \mathbf{D}\mathbf{R}\mathbf{D}$, where \mathbf{D} denotes the diagonal matrix which has the square roots of the diagonal entries for covariance matrix Σ on its diagonal.

Source

lncdfn.src

See Also

[cdfBvn](#), [cdfN](#), [lncdfmvn](#)

cdfMvnce

Purpose

Computes the upper tail of the multivariate Normal cumulative distribution function with error management.

Format

```
{y, err, retcode} = cdfMvnce(ctl, x, R, m);
```

Input

<i>ctl</i>	instance of a cdfmControl structure with members. <i>ctl.maxEvaluations</i> scalar, maximum number of evaluations. <i>ctl.absErrorTolerance</i> scalar absolute error tolerance. <i>ctl.relative</i> error tolerance.
<i>x</i>	NxK matrix, abscissae.
<i>R</i>	KxK matrix, correlation matrix.
<i>m</i>	Kx1 vector, non-centrality vector.

Output

<i>y</i>	Nx1 vector, $Pr(X \geq x R, m)$.
<i>err</i>	Nx1 vector, estimates of absolute error.
<i>retcode</i>	Nx1 vector, return codes, 0 normal completion with $err < ctl.absErrorTolerance$. 1 $err > ctl.absErrorTolerance$

	and <i>ctl.maxEvaluations</i> exceeded; increase <i>ctl.maxEvaluations</i> to decrease error.
2	$K > 100$ or $K < 1$.
3	R not positive semi-definite.
<i>missing</i>	R not properly defined.

Examples

Example 1: Uncorrelated variables

```
//Lower limits of integration for K dimensional multivariate
distribution
x = { 0  0 };

//Identity matrix, indicates
//zero correlation between variables
R = { 1 0,
      0 1 };

//Define non-centrality vector
m = { 0, 0 };

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//both variables being  $\geq 0$ 
{p, err, retcode} = cdfMvnce(ctl, x, R, m );

//Calculate joint probablity of two
//variables with zero correlation,
//both, being  $\geq 0$ 
p2 = cdfnc(0) .* cdfnc(0);
```


After the above code, both p and $p2$ should be equal to 0.25.

$$\Phi = P(0 \leq X_1 < \infty \text{ and } 0 \leq X_2 < \infty) \approx 0.25.$$

Example 2: Compute the upper tail of multivariate normal cdf at 3 separate pairs of lower limits

```
//Lower limits of integration
// x1 ≥ -1 and x2 ≥ -1.1
// x1 ≥ 0 and x2 ≥ 0.1
// x1 ≥ 1 and x2 ≥ 1.1
x = {  -1   -1.1,
       0    0.1,
       1    1.1 };

//Correlation matrix
R = {   1   0.31,
       0.31  1 };

//Define non-centrality vector
m = { 0, 0 };

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//each pair of lower limits
{p, err, retcode} = cdfMvnce(ctl, x, R, m);
```

After the above code, p should equal:

```
0.74642007
0.27999181
0.04074138
```

which means that:

$$P(x_1 \geq -1 \text{ and } x_2 \geq -1.1) = 0.7464$$

$$P(x_1 \geq +0 \text{ and } x_2 \geq +0.1) = 0.2800$$

$$P(x_1 \geq 1 \text{ and } x_2 \geq 1.1) = 0.0407$$

Example 3: Compute the upper tail of non central multivariate normal cdf

```
//Lower limits of integration
// x1 ≥ -1 and x2 ≥ -1.1
// x1 ≥ 0 and x2 ≥ 0.1
// x1 ≥ 1 and x2 ≥ 1.1
x = { -1    -1.1,
      0     0.1,
      1     1.1 };

//Correlation matrix
R = {    1  0.31,
      0.31  1 };

//Define non-centrality vector , Kx1
m = {    1,
      -2.5 };

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//each pair of lower limits
{p, err, retcode} = cdfMvnce(ctl, x, R, m);
```

After the above code, *p* should equal:

```
0.08046686
0.00455354
0.00014231
```

which means with non-central vector, the multivariate normal cdf are:

$$P(x_1 \geq -1 \text{ and } x_2 \geq -1.1) = 0.0805$$

$$P(x_1 \geq +0 \text{ and } x_2 \geq +0.1) = 0.0046$$

$$P(x_1 \geq 1 \text{ and } x_2 \geq 1.1) = 0.0001$$

Remarks

- The **cdfMvnce** evaluates the upper tail of *MVN* integral, where $1 \leq i \leq N$

$$\Phi(x_i; \Sigma) = \frac{1}{\sqrt{|\Sigma|}(2\pi)^K} \int_{x_{i1}}^{\infty} \int_{x_{i2}}^{\infty} \dots \int_{x_{iK}}^{\infty} e^{-\frac{1}{2}z' \Sigma^{-1}z} dz$$

For the non-central *MVN*, we have

$$\Phi(x_i; \Sigma, \delta) = \frac{1}{\sqrt{|\Sigma|}(2\pi)^K} \int_{x_{i1}}^{\infty} \int_{x_{i2}}^{\infty} \dots \int_{x_{iK}}^{\infty} e^{-\frac{1}{2}(z-\delta)' \Sigma^{-1}(z-\delta)} dz$$

where z denotes K -dimensional multivariate normal distribution, δ denotes the $K \times 1$ non-centrality vector with $-\infty < \delta_k < \infty$.

- The correlation matrix R is defined by $\Sigma = DRD$, where D denotes the diagonal matrix which has the square roots of the diagonal entries for covariance matrix Σ on its diagonal.

Source

`cdfm.src`

See Also

[cdfMvn2e](#), [cdfMvnce](#), [cdfMvte](#)

References

1. Genz, A. and F. Bretz, "Numerical computation of multivariate t-probabilities with application to power calculation of multiple contrasts", *Journal of Statistical Computation and Simulation*, 63:361-378, 1999.

2. Genz, A., "Numerical computation of multivariate normal probabilities", *Journal of Computational and Graphical Statistics*, 1:141-149, 1992.

cdfMvne

Purpose

Computes multivariate Normal cumulative distribution function with error management.

Format

```
{y, err, retcode} = cdfMvne(ctl, x, R, m);
```

Input

c	ctl	instance of a cdmControl structure with members.	
		<i>ctl.maxEvaluations</i>	scalar, maximum number of evaluations.
		<i>ctl.absErrorTolerance</i>	scalar absolute error tolerance.
		<i>ctl.relative</i>	error tolerance.
	x	NxK matrix, abscissae.	
	R	KxK matrix, correlation matrix.	
	m	Kx1 vector, non-centrality vector.	

Output

<i>y</i>	Nx1 vector, $Pr(X \leq x R, m)$.
<i>err</i>	Nx1 vector, estimates of absolute error.
<i>retcode</i>	Nx1 vector, return codes.
0	normal completion with <i>err</i> < <i>ctl.absErrorTolerance</i> .
1	<i>err</i> > <i>ctl.absErrorTolerance</i> and <i>ctl.maxEvaluations</i> exceeded; increase <i>ctl.maxEvaluations</i> to decrease error
2	$K > 100$ or $K < 1$
3	<i>R</i> not positive semi-definite
<i>missing</i>	<i>R</i> not properly defined

Examples

Example 1: Uncorrelated variables

```
//Upper limits of integration for K dimensional multivariate
distribution
x = { 0  0 };

//Identity matrix, indicates
//zero correlation between variables
R = { 1 0,
      0 1 };

//Define non-centrality vector
m = { 0, 0 };

//Define control structure
struct cdfmControl ctl;
```

cdfMvne

```
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//both variables being  $\leq 0$ 
{p, err, retcode} = cdfMvne(ctl, x, R, m);

//Calculate joint probability of two
//variables with zero correlation,
//both, being  $\leq 0$ 
p2 = cdfn(0) .* cdfn(0);
```

After the above code, both p and $p2$ should be equal to 0.25.

$$\Phi = P(-\infty < X_1 \leq 0 \text{ and } -\infty < X_2 \leq 0) \approx 0.25.$$

Example 2: Compute the multivariate normal cdf at 3 separate pairs of upper limits

```
//Upper limits of integration
//x1  $\leq$  -1 and x2  $\leq$  -1.1
//x1  $\leq$  0 and x2  $\leq$  0.1
//x1  $\leq$  1 and x2  $\leq$  1.1
x = { -1   -1.1,
      0    0.1,
      1    1.1 };

//Correlation matrix
R = { 1  0.31,
      0.31 1 };

//Define non-centrality vector
m = { 0, 0 };

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();
```

```
//Calculate cumulative probability of
//each pair of upper limits
{ p, err, retcode } = cdfMvne(ctl, x, R, m);
```

After the above code, p should equal:

```
0.040741382
0.31981965
0.74642007
```

which means that:

$$P(x_1 \leq -1 \text{ and } x_2 \leq -1.1) = 0.0407$$

$$P(x_1 \leq +0 \text{ and } x_2 \leq +0.1) = 0.3198$$

$$P(x_1 \leq 1 \text{ and } x_2 \leq 1.1) = 0.7464$$

Example 3: Compute the non central multivariate normal cdf

```
//Upper limits of integration
//x1 ≤ -1 and x2 ≤ -1.1
//x1 ≤ 0 and x2 ≤ 0.1
//x1 ≤ 1 and x2 ≤ 1.1
x = {  -1   -1.1,
        0    0.1,
        1    1.1 };

//Correlation matrix
R = {  1  0.31,
      0.31  1 };

//Define non-centrality vector , Kx1
m = {  1,
      -2.5 };

//Define control structure
```

cdfMvne

```
struct cdfmControl ctl;  
ctl = cdfmControlCreate();  
  
//Calculate cumulative probability of  
//each pair of upper limits  
{p, err, retcode} = cdfMvne(ctl, x, R, m);
```

After the above code, p should equal:

```
0.02246034  
0.15854761  
0.49998320
```

which means with non-central vector, the multivariate normal cdf are:

$$P(x_1 \leq -1 \text{ and } x_2 \leq -1.1) = 0.0225$$

$$P(x_1 \leq +0 \text{ and } x_2 \leq +0.1) = 0.1585$$

$$P(x_1 \leq 1 \text{ and } x_2 \leq 1.1) = 0.5000$$

Remarks

- **cdfMvne** evaluates the *MVN* integral, where $1 \leq i \leq N$

$$\Phi(x_i, \Sigma) = \frac{1}{\sqrt{|\Sigma|(2\pi)^K}} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \dots \int_{-\infty}^{x_{iK}} e^{-\frac{1}{2}z' \Sigma^{-1}z} dz$$

For the non-central *MVN* we have

$$\Phi(x_i, \Sigma, \delta) = \frac{1}{\sqrt{|\Sigma|(2\pi)^K}} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \dots \int_{-\infty}^{x_{iK}} e^{-\frac{1}{2}(z-\delta)' \Sigma^{-1}(z-\delta)} dz$$

where \mathbf{z} denotes K -dimensional multivariate normal distribution, $\boldsymbol{\delta}$ denotes the $K \times 1$ non-centrality vector with $-\infty < \delta_k < \infty$.

- The correlation matrix \mathbf{R} is defined by $\boldsymbol{\Sigma} = \mathbf{D}\mathbf{R}\mathbf{D}$, where \mathbf{D} denotes the diagonal matrix which has the square roots of the diagonal entries for covariance matrix $\boldsymbol{\Sigma}$ on its diagonal.

Source

`cdm.src`

See Also

[cdfMvne](#), [cdfMvn2e](#), [cdfMvte](#)

References

1. Genz, A. and F. Bretz, "Numerical computation of multivariate t-probabilities with application to power calculation of multiple contrasts," *Journal of Statistical Computation and Simulation*, 63:361-378, 1999.
2. Genz, A., "Numerical computation of multivariate normal probabilities," *Journal of Computational and Graphical Statistics*, 1:141-149, 1992.

cdfMvn2e

Purpose

Computes the multivariate Normal cumulative distribution function with error management over the range $[a, b]$.

Format

```
{y, err, retcode} = cdfMvn2e(ctl, a, b, R, m);
```

Input

<i>ctl</i>	instance of a cdfmControl structure with members.
<i>ctl.maxEvaluations</i>	scalar, maximum number of evaluations.
<i>ctl.absErrorTolerance</i>	scalar absolute error tolerance.
<i>ctl.relative</i>	error tolerance.
<i>a</i>	NxK matrix, lower limits.
<i>b</i>	NxK matrix, upper limits.
<i>R</i>	KxK matrix, correlation matrix.
<i>m</i>	Kx1 vector, non-centrality vector.

Output

<i>y</i>	Nx1 vector, $Pr(X \geq a \text{ and } X \leq b R, m)$.
<i>err</i>	Nx1 vector, estimates of absolute error.
<i>retcode</i>	Nx1 vector, return codes.
0	normal completion with <i>err</i> < <i>ctl.absErrorTolerance</i> .
1	<i>err</i> > <i>ctl.absErrorTolerance</i> and <i>ctl.maxEvaluations</i> exceeded; increase <i>ctl.maxEvaluations</i> to decrease error.
2	K > 100 or K < 1.
3	<i>R</i> not positive semi-definite.
<i>missing</i>	<i>R</i> not properly defined.

Examples

Example 1: Uncorrelated variables

```

//Lower limits of integration for K dimensional multivariate
distribution
a = {-1e4 -1e4};

//Upper limits of integration for K dimensional multivariate
distribution
b = {0 0};

//Identity matrix, indicates
//zero correlation between variables
R = { 1 0,
      0 1 };

//Define non-centrality vector
m = {0, 0};

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//both variables being from -1e4 to 0
{p, err, retcode} = cdfMvn2e(ctl, a, b, R, m );

```

After the above code, both p equal to 0.25.

$$\Phi = P(-10000 \leq X_1 \leq 0 \text{ and } -10000 \leq X_2 \leq 0) \approx 0.25.$$

Example 2: Compute the multivariate normal cdf at 3 separate pairs of upper limits

```

//Limits of integration
//-5 ≤ x1 ≤ -1 and -8 ≤ x2 ≤ -1.1
//-10 ≤ x1 ≤ 0 and -10 ≤ x2 ≤ 0.1
//0 ≤ x1 ≤ 1 and 0 ≤ x2 ≤ 1.1
a = { -5 -8,

```

cdfMvn2e

```
        -20 -10,  
        0    0 };  
b = {  -1 -1.1,  
      0  0.1,  
      1  1.1 };  
  
//Correlation matrix  
R = { 1 0.31,  
      0.31 1};  
  
//Define non-centrality vector  
m = {0, 0};  
  
//Define control structure  
struct cdfmControl ctl;  
ctl = cdfmControlCreate();  
  
//Calculate cumulative probability of  
//each pair of limits  
{p, err, retcode} = cdfMvn2e(ctl, a, b, R, m);
```

After the above code, p should equal:

```
0.04074118  
0.31981965  
0.13700266
```

which means that:

$$P(-5 \leq x_1 \leq -1 \text{ and } -8 \leq x_2 \leq -1.1) = 0.0407$$

$$P(-20 \leq x_1 \leq 0 \text{ and } -10 \leq x_2 \leq 0.1) = 0.3198$$

$$P(0 \leq x_1 \leq 1 \text{ and } 0 \leq x_2 \leq 1.1) = 0.1370$$

Example 3: Compute the non central multivariate normal cdf

```

//Limits of integration
// $-5 \leq x_1 \leq -1$  and  $-8 \leq x_2 \leq -1.1$ 
// $-10 \leq x_1 \leq 0$  and  $-10 \leq x_2 \leq 0.1$ 
// $0 \leq x_1 \leq 1$  and  $0 \leq x_2 \leq 1.1$ 
a = { -5  -8,
      -20 -10,
        0   0 };
b = {  -1 -1.1,
        0  0.1,
        1  1.1 };

//Correlation matrix
R = {   1  0.31,
      0.31  1 };

//Define non-centrality vector , Kx1
m  = {   1,
      -2.5 };

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//each pair of upper limits
{p, err, retcode} = cdfMvn2e(ctl, a, b, R, m);

```

After the above code, p should equal:

```

0.02246034
0.15854761
0.00094761

```

which means with non-central vector, the multivariate normal cdf are:

cdfMvn2e

$$P(-5 \leq x_1 \leq -1 \text{ and } -8 \leq x_2 \leq -1.1) = 0.0225$$

$$P(-20 \leq x_1 \leq 0 \text{ and } -10 \leq x_2 \leq 0.1) = 0.1585$$

$$P(0 \leq x_1 \leq 1 \text{ and } 0 \leq x_2 \leq 1.1) = 0.0009$$

Remarks

- **cdfMvn2e** evaluates the following non-central *MVN* integral, where $1 \leq i \leq N$

$$\Phi(a_i, b_i, \Sigma, \delta) = \frac{1}{\sqrt{|\Sigma|}(2\pi)^K} \int_{a_{i1}}^{b_{i1}} \int_{a_{i2}}^{b_{i2}} \dots \int_{a_{iK}}^{b_{iK}} e^{-\frac{1}{2}(z-\delta)' \Sigma^{-1}(z-\delta)} dz$$

where z denotes K -dimensional multivariate normal distribution, δ denotes the $K \times 1$ non-centrality vector with $-\infty < \delta_k < \infty$.

- The correlation matrix R is defined by $\Sigma = DRD$, where D denotes the diagonal matrix which has the square roots of the diagonal entries for covariance matrix Σ on its diagonal.

Source

`cdfm.src`

See Also

[cdfMvne](#), [cdfMvnce](#), [cdfMvt2e](#)

References

1. Genz, A. and F. Bretz, "Numerical computation of multivariate t-probabilities with application to power calculation of multiple contrasts," *Journal of Statistical Computation and Simulation*, 63:361-378, 1999.
2. Genz, A., "Numerical computation of multivariate normal probabilities," *Journal of*

Computational and Graphical Statistics, 1:141-149, 1992.

cdfMvtce

Purpose

Computes complement (upper tail) of multivariate Student's t cumulative distribution function with error management.

Format

```
{y, err, retcode} = cdfMvtce(ctl, x, R, m, v);
```

Input

<i>ctl</i>	instance of a cdfmControl structure with members. <i>ctl.maxEvaluations</i> scalar, maximum number of evaluations. <i>ctl.absErrorTolerance</i> scalar absolute error tolerance. <i>ctl.relErrorTolerance</i> tolerance.
<i>x</i>	NxK matrix, abscissae. K is the dimension of multivariate Student's t distribution. N is the number of <i>MVT</i> cdf integrals
<i>R</i>	KxK matrix, correlation matrix.
<i>m</i>	Kx1 vector, noncentralities.
<i>v</i>	scalar, degrees of freedom.

Output

<i>y</i>	Nx1 vector, $Pr(X \geq x R, m)$.
<i>err</i>	Nx1 vector, estimates of absolute error.

<i>retcode</i>	Nx1 vector, return codes.
0	normal completion with <i>err</i> < <i>ctl.absErrorTolerance</i> .
1	<i>err</i> > <i>ctl.absErrorTolerance</i> and <i>ctl.maxEvaluations</i> exceeded; increase <i>ctl.maxEvaluations</i> to decrease error.
2	$K > 100$ or $K < 1$.
3	<i>R</i> not positive semi-definite.
<i>missing</i>	<i>R</i> not properly defined.

Examples

Example 1: Uncorrelated variables

```
//Lower limits of integration for K dimensional multivariate
Student's t distribution
x = { 0  0 };

//Identity matrix, indicates
//zero correlation between variables
R = { 1 0,
      0 1 };

//Define non-centrality vector
m = {0, 0};

//Define degree of freedom
v = 3;

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();
```



```
//Calculate cumulative probability of
//both variables being ≥ 0
{ p, err, retcode } = cdfMvtce(ctl, x, R, m, v );

//Calculate joint probability of two
//variables with zero correlation,
//both, being ≥ 0
p2 = cdftc(0, v) .* cdftc(0, v);
```

After the above code, both p and $p2$ should be equal to 0.25.

$$T = P(0 \leq X_1 < \infty \text{ and } 0 \leq X_2 < \infty) \approx 0.25.$$

Example 2: Compute the upper tail of multivariate student's t cdf at 3 separate pairs of lower limits

```
//Lower limits of integration
//x1 ≥ -1 and x2 ≥ -1.1
//x1 ≥ 0 and x2 ≥ 0.1
//x1 ≥ 1 and x2 ≥ 1.1
x = {  -1   -1.1,
       0    0.1,
       1    1.1 };

//Correlation matrix
R = { 1  0.31,
      0.31 1};

//Define non-centrality vector
m = { 0, 0 };

//Define degree of freedom
v = 3;

//Define control structure
```

cdfMvtce

```
struct cdfmControl ctl;  
ctl = cdfmControlCreate();  
  
//Calculate cumulative probability of  
//each pair of lower limits  
{ p, err, retcode } = cdfMvtce(ctl, x, R, m, v);
```

After the above code, p should equal:

```
0.69617932  
0.28156926  
0.06752203
```

which means that:

$$P(x_1 \geq -1 \text{ and } x_2 \geq -1.1) = 0.6962$$

$$P(x_1 \geq +0 \text{ and } x_2 \geq +0.1) = 0.2816$$

$$P(x_1 \geq 1 \text{ and } x_2 \geq 1.1) = 0.0675$$

Example 3: Compute the upper tail of non central multivariate student's t cdf

```
//Lower limits of integration  
//x1 ≥ -1 and x2 ≥ -1.1  
//x1 ≥ 0 and x2 ≥ 0.1  
//x1 ≥ 1 and x2 ≥ 1.1  
x = { -1    -1.1,  
       0     0.1,  
       1     1.1 };  
  
//Correlation matrix  
R = {    1  0.31,  
      0.31    1 };  
  
//Define non-centrality vector, Kx1  
m = { 1,
```

```
        -2.5 };  
  
//Define degree of freedom  
v = 3;  
  
//Define control structure  
struct cdfmControl ctl;  
ctl = cdfmControlCreate();  
  
//Calculate cumulative probability of  
//each pair of lower limits  
{ p, err, retcode } = cdfMvtce(ctl, x, R, m, v);
```

After the above code, p should equal:

```
0.08623943  
0.00468427  
0.00049538
```

which means with non-central vector, the multivariate student's t cdf are:

$$P(x_1 \geq -1 \text{ and } x_2 \geq -1.1) = 0.0862$$

$$P(x_1 \geq +0 \text{ and } x_2 \geq +0.1) = 0.0047$$

$$P(x_1 \geq 1 \text{ and } x_2 \geq 1.1) = 0.0005$$

Remarks

- The central multivariate Student's t upper tail cdf for the i -th row of x is defined by

$$\begin{aligned}
 T(x_i; \Sigma, \nu) &= \frac{\Gamma\left(\frac{\nu+K}{2}\right)}{\Gamma\left(\frac{\nu}{2}\right)\sqrt{|\Sigma|}(\nu\pi)^K} \int_{x_{i1}}^{\infty} \int_{x_{i2}}^{\infty} \cdots \int_{x_{iK}}^{\infty} \left(1 + \frac{z' \Sigma^{-1} z}{\nu}\right)^{-\frac{\nu+K}{2}} dz \\
 &\equiv \frac{2^{1-\frac{\nu}{2}}}{\Gamma\left(\frac{\nu}{2}\right)} \int_0^{\infty} s^{\nu-1} e^{-\frac{s^2}{2}} \Phi\left(\frac{s x_i}{\sqrt{\nu}}, \infty; \Sigma\right) ds
 \end{aligned}$$

where $\nu \in \mathbb{R}^+$ is a scale (or degree of freedom) parameter, \mathbf{z} is a K -dimensional Student's t multivariate distribution, and

$$\Phi(x_i; \Sigma) = \frac{1}{\sqrt{|\Sigma|}(2\pi)^K} \int_{x_{i1}}^{\infty} \int_{x_{i2}}^{\infty} \cdots \int_{x_{iK}}^{\infty} e^{-\frac{1}{2} z' \Sigma^{-1} z} dz$$

For the non-central multivariate Student's t distribution cdf, we have

$$T(x_i; \Sigma, \nu, \delta) = \frac{\Gamma\left(\frac{\nu+K}{2}\right)}{\Gamma\left(\frac{\nu}{2}\right)\sqrt{|\Sigma|}(\nu\pi)^K} \int_{x_{i1}}^{\infty} \int_{x_{i2}}^{\infty} \cdots \int_{x_{iK}}^{\infty} \left(1 + \frac{(z-\delta)' \Sigma^{-1} (z-\delta)}{\nu}\right)^{-\frac{\nu+K}{2}} dz$$

where δ denotes the $K \times 1$ non-centrality vector with $-\infty < \delta_k < \infty$.

Another form of non-central multivariate Student's t distribution cdf is

$$T(x_i; \Sigma, \nu, \delta) = \frac{2^{1-\frac{\nu}{2}}}{\Gamma\left(\frac{\nu}{2}\right)} \int_0^{\infty} s^{\nu-1} e^{-\frac{s^2}{2}} \Phi\left(\frac{s x_i}{\sqrt{\nu}} - \delta; \Sigma\right) ds$$

- The correlation matrix \mathbf{R} is defined by covariance matrix Σ , $\Sigma = \mathbf{D} \mathbf{R} \mathbf{D}$, where \mathbf{D} denotes the diagonal matrix which has the square roots of the diagonal entries for Σ on its diagonal.

Source

cdfm.src

See Also

[cdfMvt2e](#), [cdfMvte](#), [cdfMvne](#)

1. Genz, A. and F. Bretz, "Numerical computation of multivariate t-probabilities with application to power calculation of multiple contrasts," *Journal of Statistical Computation and Simulation*, 63:361-378, 1999.
2. Genz, A., "Numerical computation of multivariate normal probabilities," *Journal of Computational and Graphical Statistics*, 1:141-149, 1992.

cdfMvte

Purpose

Computes multivariate Student's t cumulative distribution function with error management.

Format

```
{y, err, retcode} = cdfMvte(ctl, x, R, m, v);
```

Input

<i>ctl</i>	instance of a cdmControl structure with members.	
	<i>ctl.maxEvaluations</i>	scalar, maximum number of evaluations.
	<i>ctl.absErrorTolerance</i>	scalar absolute error tolerance.
	<i>ctl.relErrorTolerance</i>	tolerance.
<i>x</i>	NxK matrix, abscissae. K is the dimension of multivariate Student's t distribution. N is the number of <i>MVT</i> cdf integrals	
<i>R</i>	KxK matrix, correlation matrix.	
<i>m</i>	Kx1 vector, noncentralities.	

cdfMvte

<code>v</code>	scalar, degrees of freedom.
----------------	-----------------------------

Output

<code>y</code>	Nx1 vector, $Pr(X \leq x R,m)$.
<code>err</code>	Nx1 vector, estimates of absolute error.
<code>retcode</code>	Nx1 vector, return codes.
<code>0</code>	normal completion with <code>err < ctl.absErrorTolerance</code> .
<code>1</code>	<code>err > ctl.absErrorTolerance</code> and <code>ctl.maxEvaluations</code> exceeded; increase <code>ctl.maxEvaluations</code> to decrease error.
<code>2</code>	<code>K > 100</code> or <code>K < 1</code> .
<code>3</code>	<code>R</code> not positive semi-definite.
<code>missing</code>	<code>R</code> not properly defined.

Examples

Example 1: Uncorrelated variables

```
//Upper limits of integration for K dimensional multivariate
Student's t distribution
x = { 0  0 };

//Identity matrix, indicates
//zero correlation between variables
R = { 1 0,
      0 1 };

//Define non-centrality vector
m = { 0, 0 };
```

```
//Define degree of freedom
v = 3;

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//both variables being ≤ 0
{ p, err, retcode } = cdfMvte(ctl, x, R, m, v );

//Calculate joint probability of two
//variables with zero correlation,
//both, being ≤ 0
p2 = (1 - cdfctc(0, v)) .* (1 - cdfctc(0, v));
```

After the above code, both p and $p2$ should be equal to 0.25.

$$T = P(-\infty < X_1 \leq 0 \text{ and } -\infty < X_2 \leq 0) \approx 0.25.$$

Example 2: Compute the multivariate student's t cdf at 3 separate pairs of upper limits

```
//Upper limits of integration
//x1 ≤ -1 and x2 ≤ -1.1
//x1 ≤ 0 and x2 ≤ 0.1
//x1 ≤ 1 and x2 ≤ 1.1
x = { -1    -1.1,
      0     0.1,
      1     1.1 };

//Correlation matrix
R = { 1    0.31,
      0.31 1};
```

cdfMvte

```
//Define non-centrality vector
m = {0, 0};

//Define degree of freedom
v = 3;

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//each pair of upper limits
{p, err, retcode} = cdfMvte(ctl, x, R, m, v);
```

After the above code, p should equal:

```
0.06752203
0.31824308
0.69617932
```

which means that:

$$P(x_1 \leq -1 \text{ and } x_2 \leq -1.1) = 0.0675$$

$$P(x_1 \leq +0 \text{ and } x_2 \leq +0.1) = 0.3182$$

$$P(x_1 \leq 1 \text{ and } x_2 \leq 1.1) = 0.6962$$

Example 3: Compute the non central multivariate student's t cdf

```
//Upper limits of integration
//x1 ≤ -1 and x2 ≤ -1.1
//x1 ≤ 0 and x2 ≤ 0.1
//x1 ≤ 1 and x2 ≤ 1.1
x = { -1  -1.1,
      0   0.1,
      1   1.1 };
```



```
//Correlation matrix
R = { 1  0.31,
      0.31 1};

//Define non-centrality vector , Kx1
m = { 1,
      -2.5 };

//Define degree of freedom
v = 3;

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//each pair of upper limits
{ p, err, retcode } = cdfMvte(ctl, x, R, m, v);
```

After the above code, p should equal:

```
0.03571301
0.15854358
0.46919524
```

which means with non-central vector, the multivariate student's t cdf are:

$$P(x_1 \leq -1 \text{ and } x_2 \leq -1.1) = 0.0357$$

$$P(x_1 \leq +0 \text{ and } x_2 \leq +0.1) = 0.1585$$

$$P(x_1 \leq 1 \text{ and } x_2 \leq 1.1) = 0.4692$$

Remarks

- The central multivariate Student's t cdf for the i -th row of x is defined by

$$\begin{aligned}
 T(x_i; \Sigma, \nu) &= \frac{\Gamma\left(\frac{\nu+K}{2}\right)}{\Gamma\left(\frac{\nu}{2}\right)\sqrt{|\Sigma|}(\nu\pi)^K} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \cdots \int_{-\infty}^{x_{iK}} \left(1 + \frac{z'\Sigma^{-1}z}{\nu}\right)^{-\frac{\nu+K}{2}} dz \\
 &\equiv \frac{2^{1-\frac{\nu}{2}}}{\Gamma\left(\frac{\nu}{2}\right)} \int_0^\infty s^{\nu-1} e^{-\frac{s^2}{2}} \Phi\left(\frac{sx_i}{\sqrt{\nu}}; \Sigma\right) ds
 \end{aligned}$$

where $\nu \in \mathbb{R}^+$ is a scale (or degree of freedom) parameter, \mathbf{z} is a K -dimensional Student's t multivariate distribution, and

$$\Phi(x_i; \Sigma) = \frac{1}{\sqrt{|\Sigma|}(2\pi)^K} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \cdots \int_{-\infty}^{x_{iK}} e^{-\frac{1}{2}z'\Sigma^{-1}z} dz$$

For the non-central multivariate Student's t distribution cdf, we have

$$T(x_i; \Sigma, \nu, \delta) = \frac{\Gamma\left(\frac{\nu+K}{2}\right)}{\Gamma\left(\frac{\nu}{2}\right)\sqrt{|\Sigma|}(\nu\pi)^K} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \cdots \int_{-\infty}^{x_{iK}} \left(1 + \frac{(z-\delta)'\Sigma^{-1}(z-\delta)}{\nu}\right)^{-\frac{\nu+K}{2}} dz$$

where δ denotes the $K \times 1$ non-centrality vector with $-\infty < \delta_k < \infty$.

Another form of non-central multivariate Student's t distribution cdf is

$$T(x_i; \Sigma, \nu, \delta) = \frac{2^{1-\frac{\nu}{2}}}{\Gamma\left(\frac{\nu}{2}\right)} \int_0^\infty s^{\nu-1} e^{-\frac{s^2}{2}} \Phi\left(\frac{sx_i}{\sqrt{\nu}} - \delta; \Sigma\right) ds$$

- The correlation matrix \mathbf{R} is defined by covariance matrix Σ , $\Sigma = \mathbf{D}\mathbf{R}\mathbf{D}$, where \mathbf{D} denotes the diagonal matrix which has the square roots of the diagonal entries for Σ on its diagonal.

Source

cdfm.src

See Also

[cdfMvte](#), [cdfMvt2e](#), [cdfMvnce](#)

1. Genz, A. and F. Bretz, "Numerical computation of multivariate t-probabilities with application to power calculation of multiple contrasts," *Journal of Statistical Computation and Simulation*, 63:361-378, 1999.
2. Genz, A., "Numerical computation of multivariate normal probabilities," *Journal of Computational and Graphical Statistics*, 1:141-149, 1992.

cdfMvt2e

Purpose

Computes multivariate Student's t cumulative distribution function with error management over [a,b].

Format

```
{y, err, retcode} = cdfMvt2e(ctl, a, b, R, m, v);
```

Input

<i>ctl</i>	instance of a cdfmControl structure with members.
<i>ctl.maxEvaluations</i>	scalar, maximum number of evaluations.
<i>ctl.absErrorTolerance</i>	scalar absolute error tolerance.
<i>ctl.relErrorTolerance</i>	tolerance.
<i>a</i>	NxK matrix, lower limits. K is the dimension of multivariate Student's t distribution. N is the number of <i>MVT</i> cdf integrals.
<i>b</i>	NxK matrix, upper limits.
<i>R</i>	KxK matrix, correlation matrix.

<i>m</i>	Kx1 vector, noncentralities.
<i>v</i>	scalar, degrees of freedom.

Output

<i>y</i>	Nx1 vector, a $Pr(X \geq a \text{ and } X \leq b R, m)$.
<i>err</i>	Nx1 vector, estimates of absolute error.
<i>retcode</i>	Nx1 vector, return codes.
0	normal completion with <i>err</i> < <i>ctl.absErrorTolerance</i> .
1	<i>err</i> > <i>ctl.absErrorTolerance</i> and <i>ctl.maxEvaluations</i> exceeded; increase <i>ctl.maxEvaluations</i> to decrease error.
2	K > 100 or K < 1.
3	R not positive semi-definite.
<i>missing</i>	R not properly defined.

Examples

Example 1: Uncorrelated variables

```
//Lower limits of integration for K dimensional multivariate
distribution
a = { -1e4 -1e4 };

//Upper limits of integration for K dimensional multivariate
distribution
b = { 0 0 };

//Identity matrix, indicates
//zero correlation between variables
```

```

R = { 1 0,
      0 1 };

//Define non-centrality vector
m = { 0, 0 };

//Define degree of freedom
v = 3;

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//both variables being from -1e4 to 0
{ p, err, retcode } = cdfMvt2e(ctl, a, b, R, m, v );

```

After the above code, both p equal to 0.25.

$$T = P(-\infty < X_1 \leq 0 \text{ and } -\infty < X_2 \leq 0) \approx 0.25.$$

Example 2: Compute the multivariate student's t cdf at 3 separate pairs of upper limits

```

//Limits of integration
//-5 ≤ x1 ≤ -1 and -8 ≤ x2 ≤ -1.1
//-20 ≤ x1 ≤ 0 and -10 ≤ x2 ≤ 0.1
//0 ≤ x1 ≤ 1 and 0 ≤ x2 ≤ 1.1
a = { -5 -8,
      -20 -10,
        0 0 };
b = { -1 -1.1,
        0 0.1,
        1 1.1 };

```

cdfMvt2e

```
//Correlation matrix
R = {      1 0.31,
      0.31      1};

//Define non-centrality vector
m = { 0, 0 };

//Define degree of freedom
v = 3;

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//each pair of limits
{ p, err, retcode } = cdfMvt2e(ctl, a, b, R, m, v);
```

After the above code, p should equal:

```
0.06226091
0.31743546
0.12010880
```

which means that:

$$\begin{aligned}P(-5 \leq x_1 \leq -1 \text{ and } -8 \leq x_2 \leq -1.1) &= 0.0623 \\P(-20 \leq x_1 \leq +0 \text{ and } -10 \leq x_2 \leq +0.1) &= 0.3174 \\P(0 \leq x_1 \leq 1 \text{ and } 0 \leq x_2 \leq 1.1) &= 0.1201\end{aligned}$$

Example 3: Compute the non central multivariate student's t cdf

```
//Limits of integration
//-5 ≤ x1 ≤ -1 and -8 ≤ x2 ≤ -1.1
//-20 ≤ x1 ≤ 0 and -10 ≤ x2 ≤ 0.1
//0 ≤ x1 ≤ 1 and 0 ≤ x2 ≤ 1.1
```

```

a = {   -5  -8,
      -20 -10,
         0   0 };
b = {   -1 -1.1,
         0  0.1,
         1  1.1 };

//Correlation matrix
R = { 1      0.31,
      0.31    1 };

//Define non-centrality vector , Kx1
m = { 1,
      -2.5 };

//Define degree of freedom
v = 3;

//Define control structure
struct cdfmControl ctl;
ctl = cdfmControlCreate();

//Calculate cumulative probability of
//each pair of limits
{ p, err, retcode } = cdfMvt2e(ctl, a, b, R, m, v);

```

After the above code, p should equal:

```

0.02810292
0.15190018
0.00092484

```

which means with non-central vector, the multivariate student's t cdf are:

$$P(-5 \leq x_1 \leq -1 \text{ and } -8 \leq x_2 \leq -1.1) = 0.0281$$

$$P(-20 \leq x_1 \leq +0 \text{ and } -10 \leq x_2 \leq +0.1) = 0.1519$$

$$P(0 \leq x_1 \leq 1 \text{ and } 0 \leq x_2 \leq 1.1) = 0.0009$$

Remarks

- The central multivariate Student's t cdf for the i-th row of a and b is defined by

$$\begin{aligned} T(a_i, b_i, \Sigma, \nu) &= \frac{\Gamma\left(\frac{\nu+K}{2}\right)}{\Gamma\left(\frac{\nu}{2}\right)\sqrt{|R|}(\nu\pi)^K} \int_{a_{i1}}^{b_{i1}} \int_{a_{i2}}^{b_{i2}} \cdots \int_{a_{iK}}^{b_{iK}} \left(1 + \frac{z'\Sigma^{-1}z}{\nu}\right)^{-\frac{\nu+K}{2}} dz \\ &\equiv \frac{2^{1-\frac{\nu}{2}}}{\Gamma\left(\frac{\nu}{2}\right)} \int_0^\infty s^{\nu-1} e^{-\frac{s^2}{2}} \Phi\left(\frac{sa_i}{\sqrt{\nu}}, \frac{sb_i}{\sqrt{\nu}}; \Sigma\right) ds \end{aligned}$$

where $\nu \in \mathbb{R}^+$ is a scale (or degree of freedom) parameter, z is a K -dimensional Student's t multivariate distribution, and

$$\Phi(a_i, b_i; \Sigma) = \frac{1}{\sqrt{|\Sigma|}(2\pi)^K} \int_{a_{i1}}^{b_{i1}} \int_{a_{i2}}^{b_{i2}} \cdots \int_{a_{iK}}^{b_{iK}} e^{-\frac{1}{2}z'\Sigma^{-1}z} dz$$

For the non-central Student's t multivariate distribution cdf, we have

$$T(a_i, b_i; \Sigma, \nu, \delta) = \frac{\Gamma\left(\frac{\nu+K}{2}\right)}{\Gamma\left(\frac{\nu}{2}\right)\sqrt{|\Sigma|}(\nu\pi)^K} \int_{a_{i1}}^{b_{i1}} \int_{a_{i2}}^{b_{i2}} \cdots \int_{a_{iK}}^{b_{iK}} \left(1 + \frac{(z-\delta)'\Sigma^{-1}(z-\delta)}{\nu}\right)^{-\frac{\nu+K}{2}}$$

where δ denotes the $K \times 1$ non-centrality vector with $-\infty < \delta_k < \infty$.

Another form of non-central multivariate Student's t distribution cdf is

$$T(a_i, b_i; \Sigma, \nu, \delta) = \frac{2^{1-\frac{\nu}{2}}}{\Gamma(\frac{\nu}{2})} \int_0^\infty s^{\nu-1} e^{-\frac{s^2}{2}} \Phi\left(\frac{sa_i}{\sqrt{\nu}} - \delta, \frac{sb_i}{\sqrt{\nu}} - \delta; \Sigma\right) ds$$

- The correlation matrix R is defined by covariance matrix Σ , $\Sigma = DRD$, where D denotes the diagonal matrix which has the square roots of the diagonal entries for Σ on its diagonal.

See Also

[cdfMvte](#), [cdfMvtce](#), [cdfMvn2e](#)

Source

`cdfm.src`

1. Genz, A. and F. Bretz, "Numerical computation of multivariate t-probabilities with application to power calculation of multiple contrasts," *Journal of Statistical Computation and Simulation*, 63:361-378, 1999.
2. Genz, A., "Numerical computation of multivariate normal probabilities," *Journal of Computational and Graphical Statistics*, 1:141-149, 1992.

cdfN, cdfNc

Purpose

cdfN computes the cumulative distribution function (cdf) of the Normal distribution. **cdfNc** computes 1 minus the cdf of the Normal distribution.

Format

```
n = cdfN(x);
nc = cdfNc(x);
```

cdfN, cdfNc

Input

x	$N \times K$ matrix.
-----	----------------------

Output

n	$N \times K$ matrix.
-----	----------------------

nc	$N \times K$ matrix.
------	----------------------

Remarks

n is the integral from $-\infty$ to x of the Normal density function, and nc is the integral from x to $+\infty$.

Note that:

$$\mathbf{cdfN}(x) + \mathbf{cdfNc}(x) = 1$$

However, many applications expect $\mathbf{cdfN}(x)$ to approach 1, but never actually reach it. Because of this, we have capped the return value of \mathbf{cdfN} at $1 - \text{machine epsilon}$, or approximately $1 - 1.11\text{e-}16$. As the relative error of \mathbf{cdfN} is about $\pm 5\text{e-}15$ for $\mathbf{cdfN}(x)$ around 1, this does not invalidate the result. What it does mean is that for $\mathbf{abs}(x) > (\text{approx.}) 8.2924$, the identity does not hold true. If you have a need for the uncapped value of \mathbf{cdfN} , the following code will return it:

```
n = cdfN(x);  
if n >= 1-eps;  
    n = 1;  
endif;
```

where the value of machine epsilon is obtained as follows:

```
x = 1;  
do while 1-x /= 1;
```

```

    eps = x;
    x = x/2;
endo;

```

Note that this is an alternate definition of machine epsilon. Machine epsilon is usually defined as the smallest number such that $1 + \text{machine epsilon} > 1$, which is about $2.23\text{e-}16$. This defines machine epsilon as the smallest number such that $1 - \text{machine epsilon} < 1$, or about $1.11\text{e-}16$.

The **erf** and **erfc** functions are also provided, and may sometimes be more useful than **cdfN** and **cdfNc**.

Example

```

x = { -2 -1 0 1 2 };
n = cdfN(x);
nc = cdfNc(x);

```

After above code,

```

x  = -2.0000000  -1.0000000  0.0000000  1.0000000  2.0000000
n  =  0.0227501  0.15865525  0.5000000  0.8413447  0.9772498
nc =  0.9772498  0.84134475  0.5000000  0.1586552  0.0227501

```

See Also

[erf](#), [erfc](#), [cdfBeta](#), [cdfChic](#), [cdfFc](#), [cdfFc](#), [gamma](#)

Technical Notes

For the integral from ∞ to x :

$x \leq -37$

cdfN
underflows
and 0.0 is
returned

cdfN, cdfNc

-36	<	x	<	-10	cdfN has a relative error of approx. $\pm 5e-12$
-10	<	x	<	0	cdfN has a relative error of approx. $\pm 1e-13$
0	<	x			cdfN has a relative error of approx. $\pm 5e-15$

For **cdfNc**, i.e., the integral from x to $+\infty$, use the above accuracies but change x to $-x$.

References

1. Adams, A.G. "Remark on Algorithm 304 Normal Curve Integral." *Comm. ACM*. Vol. 12, No. 10, Oct. 1969, 565-66.
2. Hill, I.D. and S.A. Joyce. "Algorithm 304 Normal Curve Integral." *Comm. ACM*. Vol. 10, No. 6, June 1967, 374-75.
3. Holmgren, B. "Remark on Algorithm 304 Normal Curve Integral." *Comm. ACM*. Vol. 13, No. 10, Oct. 1970.
4. Mardia, K.V. and P.J. Zemroch. *Tables of the F- and Related Distributions with Algorithms*. Academic Press, New York, 1978, ISBN 0-12-471140-5.

cdfNegBinomial

Purpose

Computes the cumulative distribution function for the negative binomial distribution.

Format

```
 $p = \text{cdfNegBinomial}(f, s, prob);$ 
```

Input

f	$N \times K$ matrix, $N \times 1$ vector or scalar. $0 < f$.
s	$E \times E$ conformable with f . $0 < s$.
$prob$	The probability of success on any given trial. $E \times E$ conformable with f . $0 < prob < 1$.

Output

p	$N \times K$ matrix, $N \times 1$ vector or scalar. The probability of observing f failures before observing s .
-----	--

Example

Pat is required to sell candy bars to raise money for the 6th grade field trip. There are thirty houses in the neighborhood, and Pat is not supposed to return home until five candy bars have been sold. So the child goes door to door, selling candy bars. At each house, there is a 0.4 probability of selling one candy bar and a 0.6 probability of selling nothing.

What's the probability that Pat finishes on or before reaching the eighth house?

cdfNegBinomial

```
// f is number of failure times, f = 0, 1, 2, 3
f = seqa(0,1,4);

// p is the probability of selling the last candy bar

// the probability of selling each candy bar is 0.4, suc-
cess number = 5
p = cdfNegBinomial(f, 5, 0.4);

// since the success number is 5, so the total number is f
+ 5
f = f + 5;

print "After nth try, the probability =";
print f~p;
```

After running above code, the probability that Pat finishes on or before reaching the eighth house is 0.1736704 or 17.36704%.

After nth try, the probability =

5.0000000	0.010240000
6.0000000	0.040960000
7.0000000	0.096256000
8.0000000	0.17367040

Remarks

For invalid inputs, **cdfNegBinomial** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[cdfBinomial](#), [cdfBinomialInv](#), [cdfNegBinomialInv](#)

cdfNegBinomialInv

Purpose

Computes the quantile or inverse negative binomial cumulative distribution function.

Format

```
 $f = \text{cdfNegBinomialInv}(p, s, prob);$ 
```

Input

p	NxK matrix, Nx1 vector or scalar. $0 < f < 1$.
s	ExE conformable with p . $0 < s$.
$prob$	The probability of success on any given trial. ExE conformable with p . $0 < prob < 1$.

Output

f	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Example

Pat is required to sell candy bars to raise money for the 6th grade field trip. There are thirty houses in the neighborhood, and Pat is not supposed to return home until five candy bars have been sold. So the child goes door to door, selling candy bars. At each house, there is a 0.4 probability of selling one candy bar and a 0.6 probability of selling nothing.

If we know the probability that Pat finishes selling the last candy bar is 17.36704%, then how many times of selling nothing?

cdfN2

```
// p is the probability of selling the last candy bar
p = 0.1736704;

// f is number of failure times
f = cdfNegBinomialInv(p, 5, 0.4);

print "selling nothing times =";
print f;
```

After running above code, the number of failure times is

```
selling nothing times =
3.0000000
```

Remarks

For invalid inputs, **cdfNegBinomialInv** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[cdfBinomial](#), [cdfBinomialInv](#), [cdfNegBinomial](#)

cdfN2

Purpose

Computes interval of Normal cumulative distribution function.

Format

```
y = cdfN2(x, dx);
```

Input

x	MxN matrix, abscissae.
dx	KxL matrix, ExE conformable to x , intervals.

Output

y	max(M,K) by max(N,L) matrix, the integral from x to $x + dx$ of the Normal distribution, i.e., Pr ($x < X < x + dx$)
-----	---

Remarks

The relative error is:

$ x \leq 1$	and	$dx \leq 1$	$\pm 1e - 14$
$1 < x < 37$	and	$ dx < 1/ x $	$\pm 1e - 13$
$\min(x, x + dx) > -37$	and	$y > 1e - 300$	$\pm 1e - 11$ or better

A relative error of $\pm 1e-14$ implies that the answer is accurate to better than ± 1 in the 14th digit.

Example

```
print cdfN2(0, 1.96);
```

cdfNi

```
0.4750021048517795
```

```
print cdfN2(1,0.5);
```

```
9.1848052662599017e-02
```

```
print cdfN2(20,1e-2);
```

```
5.0038115018684521e-90
```

```
print cdfN2(-5,2);
```

```
1.3496113800582164e-03
```

```
print cdfN2(-5,0.15);
```

```
3.3065580013000255e-07
```

Source

lncdfn.src

See Also

[lncdfn2](#)

cdfNi

Purpose

Computes the inverse of the cdf of the Normal distribution.

Format

```
x = cdfNi(p);
```

Input

p	NxK real matrix, Normal probability levels, $0 \leq p \leq 1$.
-----	---

Output

x	NxK real matrix, Normal deviates, such that: $\text{cdfN}(x) = p$.
-----	--

Remarks

$\text{cdfN}(\text{cdfNi}(p)) = p$ to within the errors given below:				
		\leq	$4.6\text{e-}308$	-37.5 is returned
	p			
$4.6\text{e-}308$	$<$	$<$	$5\text{e-}24$	accurate to ± 5 in 12th digit
	p			
$5\text{e-}24$	$<$	$<$	0.5	accurate to ± 1 in 13th digit
	p			
0.5	$<$	$<$	$1 - 2.22045\text{e-}16$	accurate to ± 5 in 15th digit
	p			
		\geq	$1 - 2.22045\text{e-}16$	8.12589 is returned
	p			

See Also

[cdfN](#)

cdfPoisson

cdfPoisson

Purpose

Computes the Poisson cumulative distribution function.

Format

```
p = cdfPoisson(x, lambda);
```

Input

<i>x</i>	NxK matrix, Nx1 vector or scalar. <i>x</i> must be a positive whole number.
<i>lambda</i>	ExE conformable with <i>x</i> . The mean parameter.

Output

<i>p</i>	NxK matrix, Nx1 vector or scalar.
----------	-----------------------------------

Remarks

For invalid inputs, **cdfPoisson** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

Example

Suppose that a hospital emergency department sees an average of 200 patients during the Friday evening shift. What is the probability that they will see fewer than 250 patients during any one Friday evening shift.

```
p = cdfPoisson(250,200);
```

After running above code,

```
p = 0.99971538 or 99.715%
```

See Also

[cdfPoissonInv](#), [pdfPoisson](#), [cdfBinomial](#), [cdfNegBinomial](#)

cdfPoissonInv

Purpose

Computes the quantile or inverse Poisson cumulative distribution function.

Format

```
x = cdfPoissonInv(p, lambda);
```

Input

<i>p</i>	NxK matrix, Nx1 vector or scalar. $0 < p < 1$.
<i>lambda</i>	ExE conformable with <i>p</i> . The mean parameter.

Output

<i>x</i>	NxK matrix, Nx1 vector or scalar.
----------	-----------------------------------

Example

Suppose that a hospital emergency department sees an average of 200 patients during the Friday evening shift. If the hospital wants to have enough staff on hand to handle the patient load on 95% of Friday evenings, how many patients do they need staff on hand for?

cdfRayleigh

```
x = cdfPoissonInv(.95,200);
```

After running above code, the hospital should expect to see 224 or few patients on 95% of Friday evenings.

```
x = 224
```

Remarks

For invalid inputs, **cdfPoissoninv** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[cdfPoisson](#), [pdfPoisson](#), [cdfBinomial](#), [cdfNegBinomial](#),

cdfRayleigh

Purpose

Computes the Rayleigh cumulative distribution function.

Format

```
y = cdfRayleigh(x,b);
```

Input

x	NxK matrix, an Nx1 vector or scalar. x must be greater than or equal to 0.
b	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . b must be greater than 0.

Output

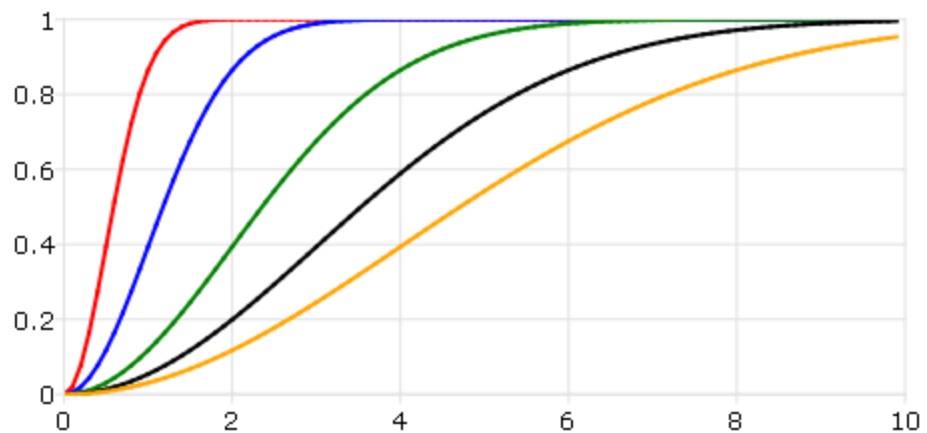
y $N \times K$ matrix, $N \times 1$ vector or scalar.

Example

Here is an example show the Rayleigh cumulative distribution plot with different scale parameters.

```
x = seqa(0,0.1,100);  
b = 0.5~1~2~3~4;  
y = cdfRayleigh(x,b);  
plotxy(x,y);
```

After running above code,



Remarks

The Rayleigh cumulative distribution function is defined as

cdfRayleighInv

$$1 - \exp\left(\frac{-x^2}{2\sigma^2}\right)$$

See Also

[cdfRayleighInv](#), [pdfRayleigh](#)

cdfRayleighInv

Purpose

Computes the Rayleigh inverse cumulative distribution function.

Format

$x = \text{cdfRayleighInv}(p, b);$

Input

p	NxK matrix, Nx1 vector or scalar. p must be greater than 0 and less than 1.
b	Shape parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with p . b must be greater than 0.

Output

x	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

$\text{cdfRayleighInv}(\text{cdfRayleigh}(x, b), b) = x$

See Also

[pdfRayleigh](#), [cdfRayleigh](#)

cdfTc

Purpose

Computes the complement of the cdf of the Student's t distribution.

Format

```
 $y = \text{cdfTc}(x, \ n);$ 
```

Input

x	$N \times K$ matrix.
n	$L \times M$ matrix, $E \times E$ conformable with x .

Output

y	$\max(N,L)$ by $\max(K,M)$ matrix.
-----	------------------------------------

Remarks

y is the integral from x to ∞ of the t distribution with n degrees of freedom.

Allowable ranges for the arguments are:

$$\begin{aligned} -\infty &\leq x \leq +\infty \\ n &> 0 \end{aligned}$$

A -1 is returned for those elements with invalid inputs.

cdfTc

This equals:

$$1 - F(x, n)$$

where F is the t cdf with n degrees of freedom. Thus, to get the t cdf, subtract **cdfTc**(x , n) from 1. The complement of the cdf is computed because this is what is most commonly needed in statistical applications, and because it can be computed with fewer problems of roundoff error.

Example

```
x = { .1, .2, .3, .4 };  
n = 3;  
y = cdfTc(x, n);
```

After running above code,

```
y =  
0.46332617  
0.42713516  
0.39188165  
0.35796758
```

See Also

[cdfTci](#)

Technical Notes

For results greater than 0.5e-30, the absolute error is approx. $\pm 1\text{e-}14$ and the relative error is approx. $\pm 1\text{e-}12$. If you multiply the relative error by the result, then take the minimum of that and the absolute error, you have the maximum actual error for any result. Thus, the actual error is approx. $\pm 1\text{e-}14$ for results greater than 0.01. For results less than 0.01, the actual error will be less. For example, for a result of 0.5e-30, the actual error is only $\pm 0.5\text{e-}42$.

References

1. Abramowitz, M. and I.A. Stegun, eds. *Handbook of Mathematical Functions*. 7th ed. Dover, New York, 1970. ISBN 0-486-61272-4.

2. Hill, G.W. "Algorithm 395 Student's t-Distribution." **Comm. ACM**. Vol. 13, No. 10, Oct. 1970.

3. Hill, G.W. "Reference Table: Student's t-Distribution Quantiles to 20D." *Division of Mathematical Statistics Technical Paper No. 35*. Commonwealth Scientific and Industrial Research Organization, Australia, 1972.

cdfTci

Purpose

Computes the inverse of the complement of the Student's *t* cdf.

Format

$x = \text{cdfTci}(p, \ n);$

Input

p	$N \times K$ real matrix, complementary Student's <i>t</i> probability levels, $0 \leq p \leq 1$.
n	$L \times M$ real matrix, degrees of freedom, $n > 1$, n need not be integral. $E \times E$ conformable with p .

Output

x	$\max(N,L)$ by $\max(K,M)$ real matrix, Student's <i>t</i> deviates, such that $\text{cdfTc}(x, \ n) = p$.
-----	---

cdfTnc

Remarks

$\text{cdfTnc}(\text{cdfTci}(p, \ n)) = p$ to within the errors given below:

$0.5\text{e-}30$	$<$	p	$<$	0.01	accurate to ± 1 in 12th digit
0.01	$<$	p			accurate to $\pm 1\text{e-}14$

Extreme values of arguments can give rise to underflows, but no overflows are generated.

See Also

[cdfTc](#)

cdfTnc

Purpose

The integral under noncentral Student's t distribution, from $-\infty$ to x . It can return a vector of values, but the degrees of freedom and noncentrality parameter must be the same for all values of x .

Format

$$y = \text{cdfTnc}(x, \ v, \ d);$$

Input

x	Nx1 vector, values of upper limits of integrals.
v	scalar, degrees of freedom, $v > 0$.
d	scalar, noncentrality parameter.
This is the <u>square root of the noncentrality parameter</u> that	

sometimes goes under the symbol λ . (See Scheffe, *The Analysis of Variance*, App. IV, 1959.)

Output

 y

$N \times 1$ vector, integrals from $-\infty$ to x of noncentral t .

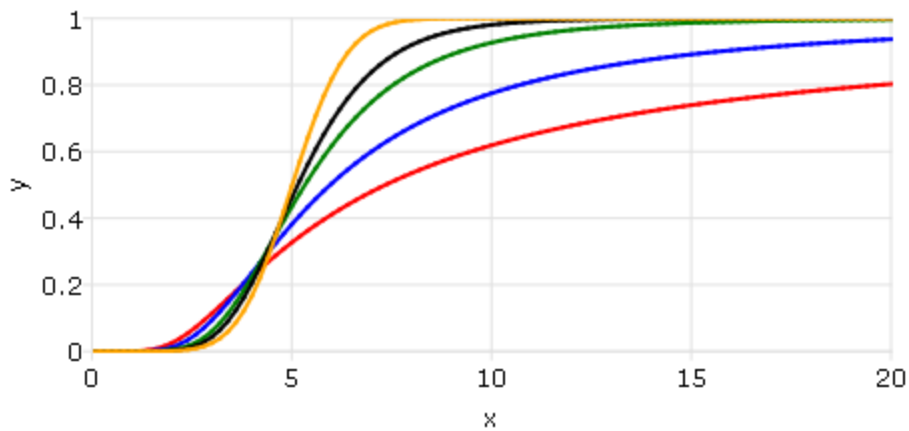
Examples

Example 1 noncentral t distributions with different parameters.

```
// sigma = 5
x = seqa(0,0.2,101);
sigma = 5;
df = 1~2~5~10~100;
y = cdfTnc(x, df, sigma');
plotxy(x, y);
```

After running above code,

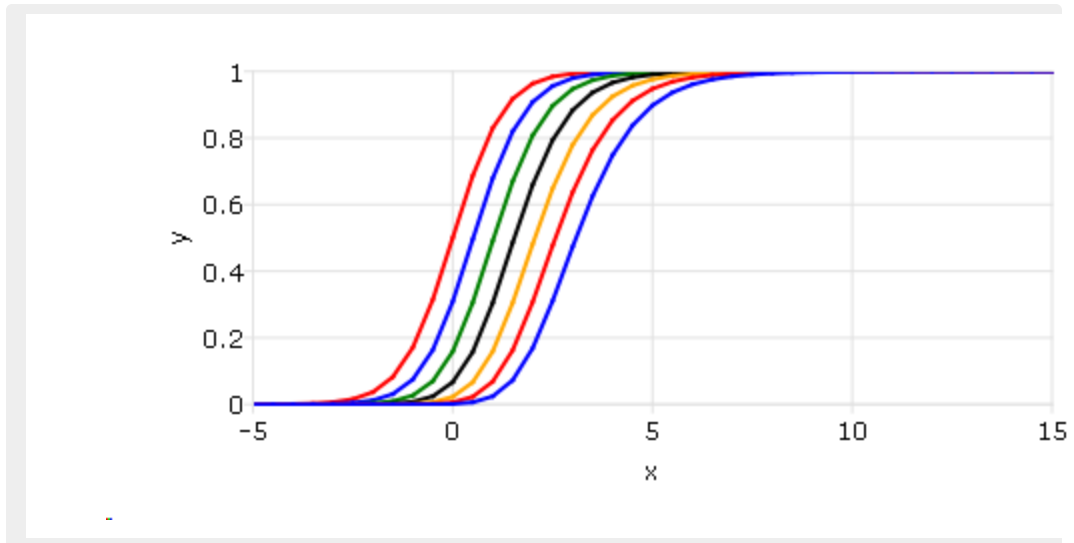
cdfTnc



Example 2 noncentral t distributions with different degree of freedoms.

```
// df = 10  
x = seqa(-5,0.5,41);  
sigma = seqa(0, 0.5, 7);  
df = 10;  
y = cdfTnc(x, df, sigma');  
plotxy(x, y);
```

After running above code,



Remarks

$$\text{cdfTc}(x, v) = 1 - \text{cdfTnc}(x, v, 0).$$

See Also

[cdfFnc](#), [cdfChinc](#)

cdfTvn

Purpose

Computes the cumulative distribution function of the standardized trivariate Normal density (lower tail).

Format

```
c = cdfTvn(x1, x2, x3, rho12, rho23, rho13);
```

Input

<i>x1</i>	Nx1 vector of upper limits of integration for variable 1.
<i>x2</i>	Nx1 vector of upper limits of integration for variable 2.
<i>x3</i>	Nx1 vector of upper limits of integration for variable 3.
<i>rho12</i>	scalar or Nx1 vector of correlation coefficients between the two variables <i>x1</i> and <i>x2</i> .
<i>rho23</i>	scalar or Nx1 vector of correlation coefficients between the two variables <i>x2</i> and <i>x3</i> .
<i>rho13</i>	scalar or Nx1 vector of correlation coefficients between the two variables <i>x1</i> and <i>x3</i> .

Output

<i>c</i>	Nx1 vector containing the result of the triple integral from $-\infty$ to <i>x1</i> , $-\infty$ to <i>x2</i> , and $-\infty$ to <i>x3</i> of the standardized trivariate Normal density.
----------	--

Remarks

Allowable ranges for the arguments are:	
	$-\infty < \mathbf{x1} < \infty$
	$-\infty < \mathbf{x2} < \infty$
	$-\infty < \mathbf{x3} < \infty$
	$-1 < \mathbf{rho12} < 1$
	$-1 < \mathbf{rho23} < 1$
	$-1 < \mathbf{rho13} < 1$

c

In addition, ρ_{12} , ρ_{23} and ρ_{13} must come from a legitimate positive definite matrix. A -1 is returned for those rows with invalid inputs.

A separate integral is computed for each row of the inputs.

The first 3 arguments (x_1 , x_2 , x_3) must be the same length, N. The second 3 arguments (ρ_{12} , ρ_{23} , ρ_{13}) must also be the same length, and this length must be N or 1. If it is 1, then these values will be expanded to apply to all values of x_1 , x_2 , x_3 . All inputs must be column vectors.

To find the integral under a general trivariate density, with x_1 , x_2 , and x_3 having nonzero means and any positive standard deviations, transform by subtracting the mean and dividing by the standard deviation. For example:

$$x_1 = (x_1 - \text{mean}(x_1)) / \text{std}(x_1)$$

The absolute error for **cdfTvn** is approximately $\pm 2.5\text{e-}8$ for the entire range of arguments.

See Also

[cdfN](#), [cdfBvn](#)

References

1. Daley, D.J. "Computation of Bi- and Tri-variate Normal Integral." *Appl. Statist.* Vol. 23, No. 3, 1974, 435-38.
2. Steck, G.P. "A Table for Computing Trivariate Normal Probabilities." *Ann. Math. Statist.* Vol. 29, 780-800.

cdfWeibull

Purpose

Computes the cumulative distribution function for the Weibull distribution.

cdfWeibull

Format

```
y = cdfWeibull(x,k,lambda);
```

Input

<i>x</i>	NxK matrix, Nx1 vector or scalar. <i>x</i> must be greater than 0.
<i>k</i>	Shape parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with <i>x</i> . <i>k</i> must be greater than 0.
<i>lambda</i>	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with <i>x</i> . <i>lambda</i> must be greater than 0.

Output

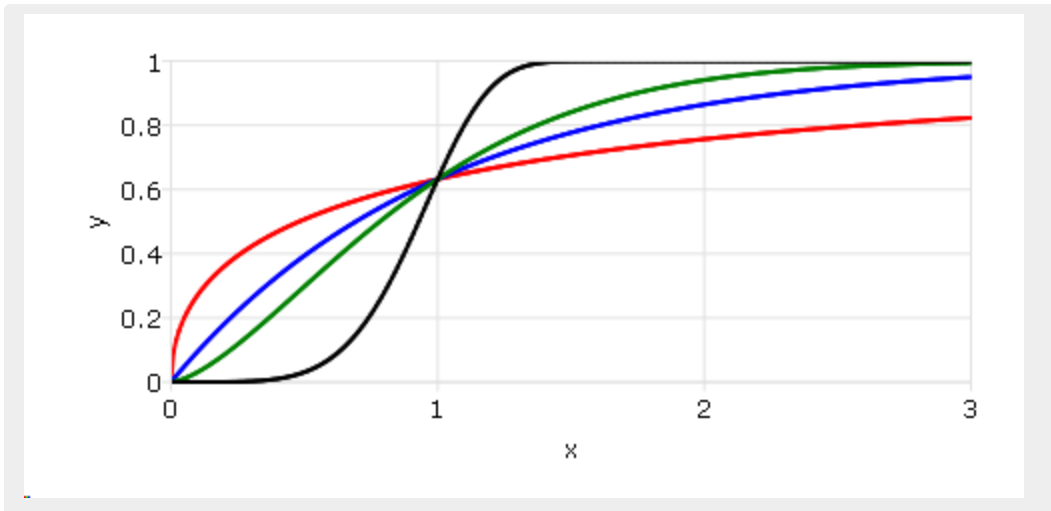
<i>y</i>	NxK matrix, Nx1 vector or scalar.
----------	-----------------------------------

Example

Calculate the cdf for the Weibull distribution with different shape parameters.

```
// lambda = 1
x = sega(0,0.01,301);
k = 0.5~1~1.5~5;
lambda = 1;
y = cdfWeibull(x, k, lambda);
plotxy(x, y);
```

After running above code,



Remarks

The Weibull cumulative distribution function is defined as:

$$f(x; k, \lambda) = 1 - e^{-(x/\lambda)^k}$$

See Also

[pdfWeibull](#), [cdfWeibullInv](#)

cdfWeibullInv

Purpose

Computes the Weibull inverse cumulative distribution function.

Format

```
y = cdfWeibullInv(p,k,lambda);
```

cdir

Input

p	NxK matrix, Nx1 vector or scalar. p must be greater than 0 and less than 1.
k	Shape parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . k must be greater than 0.
$lambda$	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . $lambda$ must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

See Also

[pdfWeibull](#), [cdfWeibull](#)

cdir

Purpose

Returns the current directory.

Format

$y = \text{cdir}(s);$

Input

s	string, if the first character is 'A'-'Z' and the second character is a colon ':' then that drive will be used. If not, the current default drive will be used.
-----	---

Output

y	string containing the drive and full path name of the current directory on the specified drive.
-----	---

Remarks

If the current directory is the root directory, the returned string will end with a backslash, otherwise it will not.

A null string or scalar zero can be passed in as an argument to obtain the current drive and path name.

Example

If the current working directory is C:\gauss:

```
x = cdir(0);  
y = cdir("d:");  
print x;  
print y;
```

The code above will return:

```
C:\gauss  
d:
```

ceil

Purpose

Round up toward $+\infty$.

ceil

Format

```
y = ceil(x);
```

Input

x NxK matrix.

Output

y NxK matrix.

Remarks

This rounds every element in the matrix *x* to an integer. The elements are rounded up toward $+\infty$.

Example

```
x = 10*rndn(2,2);  
y = ceil(x);
```

After the code above, the matrices *x* and *y* should hold values similar to below. Answers will vary due to the use of random numbers as the input to the **ceil** function.

```
    x =  8.73383   -0.783488    y =  9.0000000   0.0000000  
        13.1106    7.155113       14.000000   8.0000000
```

See Also

[floor](#), [trunc](#)

changeDir

Purpose

Changes the working directory within a program.

Format

```
d = changeDir(s);
```

Input

<i>s</i>	string, directory to change to.
----------	---------------------------------

Output

<i>d</i>	string, new working directory, or null string if change failed.
----------	---

See Also

[chdir](#), [cdir](#)

chdir

Purpose

Changes working directory in interactive mode.

Format

```
chdir dirst;
```

chiBarSquare

Input

<i>dirstr</i>	literal or ^string, directory to change to.
---------------	---

Remarks

This is for interactive use. Use **ChangeDir** in a program.

If the directory change fails, **chdir** prints an error message.

See Also

[changedir](#), [cdir](#)

chiBarSquare

Purpose

Compute compute the probability for a chi-bar square statistic from an hypothesis involving parameters under constraints.

Format

```
SLprob = chiBarSquare(SL, H, a, b, c, d, bounds);
```

Input

<i>SL</i>	scalar, chi-bar square statistic
<i>H</i>	KxK matrix, positive covariance matrix
<i>a</i>	MxK matrix, linear equality constraint coefficients
<i>b</i>	Mx1 vector, linear equality constraint constants
These arguments specify the linear equality constraints of the following type:	

	$a * X = b$ <p>where x is the $K \times 1$ parameter vector.</p>
c	$M \times K$ matrix, linear inequality constraint coefficients.
d	$M \times 1$ vector, linear inequality constraint constants.
	These arguments specify the linear inequality constraints of the following type:
	$c * X \geq d$ <p>where x is the $K \times 1$ parameter vector.</p>
$bounds$	$K \times 2$ matrix, bounds on parameters. The first column contains the lower bounds, and the second column the upper bounds.

Output

$SLprob$	scalar, probability of SL .
----------	-------------------------------

Remarks

See Silvapulle and Sen, <i>Constrained Statistical Inference</i> , page 75 for further details about this function. Let
$Z_{p \times 1} \sim N(0, V)$
where V is a positive definite covariance matrix. Define
$x^{-2}(V, C) = Z'V^{-1}Z - \min_{\theta \in C} (Z - \theta)'V^{-1}(Z - \theta)$

C

chol

C is a closed convex cone describing a set of constraints. **ChiBarSquare** computes the probability of this statistic given V and C .

Example

```
V = { 0.0005255598 -0.0006871606 -0.0003191342,  
      -0.0006871606 0.0037466205 0.0012285813,  
      -0.0003191342 0.0012285813 0.0009081412 };  
  
SL = 3.860509;  
  
Bounds = { 0 200, 0 200, 0 200 };  
  
vi = invpd(v);  
  
SLprob = chiBarSquare(SL,Vi,0,0,0,0,bounds);
```

After running above code,

```
SLprob = 0.10885000
```

Source

hypotest.src

chol

Purpose

Computes the Cholesky decomposition of a symmetric, positive definite square matrix.

Format

```
y = chol(x);
```

Input

x	$N \times N$ matrix.
-----	----------------------

Output

y	$N \times N$ matrix containing the Cholesky decomposition of x .
-----	--

Remarks

y is the "square root" matrix of x . That is, it is an upper triangular matrix such that $x = y'y$.

chol does not check to see that the matrix is symmetric. **chol** will look only at the upper half of the matrix including the principal diagonal.

If the matrix x is symmetric but not positive definite, either an error message or an error code will be generated, depending on the lowest order bit of the trap flag:

trap 0	Print error message and terminate program.
---------------	--

trap 1	Return scalar error code 10.
---------------	------------------------------

See **scalerr** and [trap](#) for more details about error codes.

Example

```
//'moment' calculates x'*x with options for handling missing data
x = moment (rndn(100,4),0);
y = chol(x);
```

choldn

```
//y'y is equivalent to y'*y
ypy = y'y;
```

	95.2801	8.6983	3.7248	1.5449		9.7612	0.8911	0.3816	0.1583
x =	8.6983	83.4547	-6.1455	-12.5551	y =	0.0000	9.0918	-0.7133	-1.3964
	3.7248	-6.1455	87.6666	-3.0284		0.0000	0.0000	9.3280	-0.4379
	1.5449	-12.5551	-3.0284	90.8311		0.0000	0.0000	0.0000	9.4162

	95.2801	8.6983	3.7248	1.5449
ypy =	8.6983	83.4547	-6.1455	-12.5551
	3.7248	-6.1455	87.6666	-3.0284
	1.5449	-12.5551	-3.0284	90.8311

See Also

[crout](#), [solpd](#)

choldn

Purpose

Performs a Cholesky downdate of one or more rows on an upper triangular matrix.

Format

`r = choldn(C, x);`

Input

<i>C</i>	KxK upper triangular matrix.
<i>x</i>	NxK matrix, the rows to downdate <i>C</i> with.

Output

r KxK upper triangular matrix, the downdated matrix.

Remarks

If **trap 1** is set, **choldn** returns scalar error code 60, otherwise it terminates the program with an error message.

C should be a Cholesky factorization.

```
choldn(C, x);
```

is equivalent to

```
chol(C'C - x'x);
```

but **choldn** is numerically much more stable.

WARNING: it is possible to render a Cholesky factorization non-positive definite with **choldn**. You should keep an eye on the ratio of the largest diagonal element of r to the smallest--if it gets very large, r may no longer be positive definite. This ratio is a rough estimate of the condition number of the matrix.

Example

```
let C[3,3] = 20.16210005 16.50544413 9.86676135
              0 11.16601462 2.97761666
              0 0 11.65496052;
let x[2,3] = 1.76644971 7.49445820 9.79114666
              6.87691156 4.41961438 4.32476921;
r = choldn(C,x);

      18.8706 15.3229 8.0495
r = 0.0000 9.3068 -2.1201
      0.0000 0.0000 7.6288
```

cholsol

See Also

[cholup](#), [chol](#)

cholsol

Purpose

Solves a system of linear equations given the Cholesky factorization of the system.

Format

$x = \text{cholsol}(b, C);$

Input

b	NxK matrix.
C	NxN matrix.

Output

x	NxK matrix.
-----	-------------

Remarks

C is the Cholesky factorization of a linear system of equations A . x is the solution for $Ax = b$. b can have more than one column. If so, the system is solved for each column, i.e., $A * x[:, i] = b[:, i]$.

Since $A^{-1} = I/A$ and **eye**(N) creates an identity matrix of size N :

`cholsol (eye (N) , C) ;`

is equivalent to:

```
invpd(A);
```

Thus, if you have the Cholesky factorization of A , **cholsol** is the most efficient way to obtain the inverse of A .

Example

```
//Assign the right-hand side 'b' and the Cholesky
//factorization 'C'
b = { 0.03177513, 0.41823100, 1.70129375 };
C = { 1.73351215 1.53201723 1.78102499,
      0 1.09926365 0.63230050,
      0 0 0.67015361 };

//Solve the system of equations
x = cholsol(b,C);

//Note: C'C is equivalent to C'*C
A = C'C;

//Solve the system of equations
x2 = b/A;

      -1.9440      -1.9440
x = -1.5269  x2 = -1.5269
      3.2158      3.2158
```

See Also

[chol](#)

cholup

cholup

Purpose

Performs a Cholesky update of one or more rows on an upper triangular matrix.

Format

```
r = cholup(C, x);
```

Input

<i>C</i>	KxK upper triangular matrix.
<i>x</i>	NxK matrix, the rows to update <i>C</i> with.

Output

<i>r</i>	KxK upper triangular matrix, the updated matrix.
----------	--

Remarks

C should be a Cholesky factorization.

cholup(*C*, *x*) is equivalent to **chol**(*C'**C* + *x'**x*), but **cholup** is numerically much more stable.

Example

```
let C[3,3] = 18.87055964 15.3229443  8.04947012
              0  9.30682813 -2.12009339
              0  0              7.62878355;
let x[2,3] = 1.76644971 7.49445820 9.79114666
              6.87691156 4.41961438 4.32476921;
r = cholup(C,x);
```

C


```
      20.162100      16.505444      9.8667614
r = 0.0000000      11.166015      2.9776167
      0.0000000      0.0000000      11.654961
```

See Also

[choldn](#)

chrs

Purpose

Converts a matrix of ASCII values into a string containing the appropriate characters.

Format

```
y = chrs(x);
```

Input

x	NxK matrix.
---	-------------

Output

y	string of length N*K containing the characters whose ASCII values are equal to the values in the elements of x.
---	---

Remarks

This function is useful for embedding control codes in strings and for creating variable

chr

length strings when formatting printouts, reports, etc.

Example

```
//42 is the ascii value for an asterisk '*'
print chr(42);
```

The code above returns:

```
*
```

chr can be used to create an interactive program in which the user is prompted for keyboard input which the code uses to make decisions.

```
//Print a string to prompt the user for input
print "Choose a parameter: Enter [a,b,c]";

//Wait for the user to enter a keystroke and assign the
//ASCII value of that key to 'param'
param = keyw;

//Convert the ASCII value to a string
paramString = chr(param);

if paramString == "a";
    print "You have chosen:" "a";
    //execute code for this choice
elseif paramString == "b";
    print "You have chosen:" "b";
    //execute code for this choice
elseif paramString == "c";
    print "You have chosen:" "c";
    //execute code for this choice
endif;
```

See Also

[vals](#), [fios](#), [stof](#)

clear

Purpose

Clears space in memory by setting matrices equal to scalar zero.

Format

```
clear x, y;
```

Remarks

If your program is running out of memory, or uses considerable system resources, using `clear` to deallocate large matrices after they are no longer needed may allow it to run more efficiently.

```
clear x;
```

is equivalent to

```
x = 0;
```

Matrix names are retained in the symbol table after they are cleared.

Matrices can be `clear`'ed even though they have not previously been defined. `clear` can be used to initialize matrices to scalar 0.

Example

```
A = randn(1000, 1000);  
//Code that uses 'A' would be here  
//Free memory holding 'A'
```

clearg

```
clear A;
```

See Also

[clearg](#), [new](#), [show](#), [delete](#)

clearg

Purpose

Clears global symbols by setting them equal to scalar zero.

Format

```
clearg a, b, c;
```

Output

a, b, c scalar global matrices containing 0.

Remarks

It is considered a best practice to avoid using global variables inside of procedures when possible.

```
clearg x;
```

is equivalent to

```
x = 0;
```

where *x* is understood to be a global symbol. `clearg` can be used to initialize symbols not previously referenced. This command can be used inside of procedures to clear global matrices. It will ignore any locals by the same name.

See Also

[clear](#), [delete](#), [new](#), [show](#), [local](#)

close

Purpose

Closes a **GAUSS** file.

Format

```
y = close(handle);
```

Input

<i>handle</i>	scalar, the file handle given to the file when it was opened with the open , create , or fopen command.
---------------	--

Output

<i>y</i>	scalar, 0 if successful, -1 if unsuccessful.
----------	--

Remarks

handle is the scalar file handle created when the file was opened. It will contain an integer which can be used to refer to the file.

close will close the file specified by *handle*, and will return a 0 if successful and a -1 if not successful. The handle itself is not affected by **close** unless the return value of **close** is assigned to it.

If *f1* is a file handle and it contains the value 7, then after:

close

```
call close (f1) ;
```

the file will be closed but *f1* will still have the value 7. The best procedure is to do the following:

```
f1 = close (f1) ;
```

This will set *f1* to 0 upon a successful close.

It is important to set unused file handles to zero because both `open` and `create` check the value that is in a file handle before they proceed with the process of opening a file. During `open` or `create`, if the value that is in the file handle matches that of an already open file, the process will be aborted and a File already open error message will be given. This gives you some protection against opening a second file with the same handle as a currently open file. If this happened, you would no longer be able to access the first file.

An advantage of the `close` function is that it returns a result which can be tested to see if there were problems in closing a file. The most common reason for having a problem in closing a file is that the disk on which the file is located is no longer in the disk drive--or the handle was invalid. In both of these cases, `close` will return a -1.

Files are not automatically closed when a program terminates. This allows users to run a program that opens files, and then access the files from interactive mode after the program has been run. Files are automatically closed when **GAUSS** exits to the operating system or when a program is terminated with the `end` statement. `stop` will terminate a program but not close files.

As a rule it is good practice to make `end` the last statement in a program, unless further access to the open files is desired from interactive mode. You should close files as soon as you are done writing to them to protect against data loss in the case of abnormal termination of the program due to a power or equipment failure.

The danger in not closing files is that anything written to the files may be lost. The disk directory will not reflect changes in the size of a file until the file is closed and system buffers may not be flushed.

Example

```
new;
cls;

// Create 'mydata' matrix
mydata = sega(1,1,3);

// Using saved function to save mydata matrix into
'mydata.dat' file
saved(mydata, "mydata.dat", "x");

// Set a random seed
rndseed 855;

// Open 'mydata.dat' file
open f1 = mydata for append;

// Create an appended data set 'x'
x = randu(3,1);

y = writer(f1,x);
f1 = close(f1);

data_new = loadadd("mydata.dat");

print "mydata = " mydata;
print "x = " x;
print "data_new = " data_new;
```

After running above code,

closeall

```
1.0000000
mydata =
1.0000000
2.0000000
3.0000000
x =
0.33589398
0.62804541
0.017829664
data_new =
1.0000000
2.0000000
3.0000000
0.33589398
0.62804541
0.017829664
```

The first 1 means the "mydata.dat" file is closed.

See Also

[closeall](#)

closeall

Purpose

Closes all currently open **GAUSS** files.

Format

```
closeall;
closeall list_of_handles;
```


Remarks

`list_of_handles` is a comma-delimited list of file handles.

`closeall` with no specified list of handles will close all files. The file handles will not be affected. The main advantage of using `closeall` is ease of use; the file handles do not have to be specified, and one statement will close all files.

When a list of handles follows `closeall`, all files are closed and the file handles listed are set to scalar 0. This is safer than `closeall` without a list of handles because the handles are cleared.

It is important to set unused file handles to zero because both `open` and `create` check the value that is in a file handle before they proceed with the process of opening a file. During `open` or `create`, if the value that is in the file handle matches that of an already open file, the process will be aborted and a File already open error message will be given. This gives you some protection against opening a second file with the same handle as a currently open file. If this happened, you would no longer be able to access the first file.

Files are not automatically closed when a program terminates. This allows users to run a program that opens files, and then access the files from interactive mode after the program has been run. Files are automatically closed when **GAUSS** exits to the operating system or when a program is terminated with the `end` statement. `stop` will terminate a program but not close files.

As a rule it is good practice to make `end` the last statement in a program, unless further access to the open files is desired from interactive mode. You should close files as soon as you are done writing to them to protect against data loss in the case of abnormal termination of the program due to a power or equipment failure.

The danger in not closing files is that anything written to the files may be lost. The disk directory will not reflect changes in the size of a file until the file is closed and system buffers may not be flushed.

closeall

Example

```
new;
cls;

// Create 'mydata' matrix
mydata = sega(1,1,3);

// Using saved function to save mydata matrix into
'mydata.dat' file
saved(mydata, "mydata.dat", "x");
open f1 = dat1 for read;
open f2 = dat1 for update;
x = readr(f1, rowsf(f1));
x = sqrt(x);
call writer(f2, x);
closeall f1, f2;

// Check the new data file
mydata_new = loadd("mydata.dat");
print "mydata = " mydata;
print "x = " x;
print "mydata_new = " mydata_new;
```

After running the above code,

```
      1.0000000
mydata =
      1.0000000
      2.0000000
      3.0000000
x =
      1.0000000
      1.4142136
```

```
1.7320508
mydata_new =
1.0000000
1.4142136
1.7320508
```

The first 1 means the "mydata.dat" file is closed.

See Also

[close](#), [open](#)

cls

Purpose

Clears the program input/output window.

Format

```
cls;
```

Remarks

This command clears the window and locates the cursor at the upper left hand corner of the window. It is sometimes useful to put a `cls` statement at the beginning of a program that prints a report to the screen so that you have fewer lines of data to look at.

See Also

[locate](#)

code

code

Purpose

Allows a new variable to be created (coded) with different values depending upon which one of a set of logical expressions is true.

Format

```
y = code(logical, new_vals);
```

Input

<i>logical</i>	NxK matrix of 1's and 0's. Each column of this matrix is created by a logical expression using "dot" conditional and boolean operators. Each of these expressions should return a column vector result. The columns are horizontally concatenated to produce <i>logical</i> . If more than one of these vectors contains a 1 in any given row, the code function will terminate with an error message.
<i>new_vals</i>	(K+1)x1 vector containing the values to be assigned to the new variable.

Output

<i>y</i>	Nx1 vector containing the new values.
----------	---------------------------------------



Example

Example 1

Suppose we have a vector of blood pressure data that we want to separate into two classes. Class 1 will contain the observations with a blood pressure value below 120. The others will belong to class 2.

```
//Blood pressure data
x = { 91,
      121,
      99,
      135,
      110,
      155 };

//Values for the classes
new_val = { 1,
            2 };

//Create a vector containing a 1 for every element
//which is less than 120, or a 0 otherwise
logical = x .< 120;

//Create a new vector which contains the class
//assignment for each element in 'x'
x_class = code(logical, new_val);
```

After the code above:

x = 91	logical = 1	x_class = 1
121	0	2
99	1	1
135	0	2
110	1	1
155	0	2

Example 2

Continuing with the blood pressure example from above, we will now create a new categorical variable with 3 levels. Level 1 will contain observations less than or equal to 100. Level 2 will contain observations greater than 100 and less than or equal to 120. Level 3 will contain observations greater than 120.

```
//Blood pressure data
x = { 91,
      121,
      99,
      135,
      110,
      155 };

//Values for the classes
new_val = { 1,
            2,
            3 };

//Create a vector containing a 1 for every element
//which is 100 or less, or a 0 otherwise
logical_1 = x .<= 100;

//Create a vector containing a 1 for every element
//which is between 100 and 120, or a 0 otherwise
logical_2 = x .> 100 .and x .<= 120;

//Form a 2 column logical vector using
//horizontal concatenation
logical = logical_1 ~ logical_2;

//Create a new vector which contains the class
```

```
//assignment for each element in 'x'
x_class = code(logical, new_val);
```

After the code above:

x = 91	logical = 1 0	x_class = 1
121	0 0	3
99	1 0	1
135	0 0	3
110	0 1	2
155	0 0	3

Remarks

For every row in *logical*, if a 1 is in the first column, the first element of *new_vals* is used. If a 1 is in the second column, the second element of *new_vals* is used, and so on. If there are only zeros in the row, the last element of *new_vals* is used. This is the default value.

If there is more than one 1 in any row of *logical*, the function will terminate with an error message.

Source

datatran.src

See Also

[recode](#), [reclassifyCuts](#), [reclassify](#), [substute](#), [rescale](#), [dummy](#)

code (dataloop)

Purpose

Creates new variables with different values based on a set of logical expressions.

code (dataloop)

Format

```
code [[#]] [[$]] var [[default defval]] with
  val_1 for expression_1,
  val_2 for expression_2,
  .
  .
  .
  val_n for expression_n;
```

Input

<i>var</i>	literal, the new variable name.
<i>defval</i>	scalar, the default value if none of the expressions are TRUE.
<i>val</i>	scalar, value to be used if corresponding expression is TRUE.
<i>expression</i>	logical scalar-returning expression that returns nonzero TRUE or zero FALSE.

Remarks

If '\$' is specified, the new variable will be considered a character variable. If '#' or nothing is specified, the new variable will be considered numeric.

The logical expressions must be mutually exclusive, i.e., only one may return TRUE for a given row (observation).

Any variables referenced must already exist, either as elements of the source data set, as externs, or as the result of a previous `make`, `vector`, or `code` statement.

If no default value is specified, 999 is used.

Example

```
code agecat default 5 with
1 for age < 21,
2 for age >= 21 and age < 35,
3 for age >= 35 and age < 50,
4 for age >= 50 and age < 65;
```

```
code $ sex with
"MALE" for gender == 1,
"FEMALE" for gender == 0;
```

See Also

[recode \(dataloop\)](#)

cols

Purpose

Returns the number of columns in a matrix.

Format

```
 $p$  = cols( $x$ );
```

Input

x	$N \times K$ matrix or sparse matrix.
-----	---------------------------------------

Output

p	number of columns in x .
-----	----------------------------

colsf

Remarks

Use **getorders** to return both the number of rows and columns in one call. If *x* is an empty matrix, **rows**(*x*) and **cols**(*x*) both return 0.

Example

```
//Create a 100x3 matrix of uniform random numbers
x = randu(100,3);

//Find out how many columns are in 'x'
p = cols(x);
```

After the code above:

```
p = 3
```

See Also

[rows](#), [colsf](#), [getorders](#), [show](#)

colsf

Purpose

Returns the number of columns in a **GAUSS** data (*.dat*) file or **GAUSS** matrix (*.fmt*) file.

Format

```
yf = colsf(fh);
```

Input

<i>fh</i>	file handle of an open file.
-----------	------------------------------

Output

yf number of columns in the file that has the handle fh .

Remarks

In order to call `colsf` on a file, the file must be open.

Example

```
//Create a file with 10 columns
create fp = myfile with x,10,4;

//Calculate the number of rows of the file created above
nCols = colsf(fp);
```

The result will be

```
nCols = 10
```

See Also

[rowsf](#), [cols](#), [show](#)

combine

Purpose

Computes combinations of N things taken K at a time.

Format

```
 $y$  = combine( $N$ ,  $K$ );
```

combinate

Input

N	scalar.
K	scalar.

Output

y	$M \times K$ matrix, where M is the number of combinations of N things taken K at a time.
-----	---

Remarks

"Things" are represented by a sequence of integers from 1 to N , and the integers in each row of y are the combinations of those integers taken K at a time.

Example

```
//Calculate all combinations of 4 items chosen 2 at a time
n = 4;
k = 2;
y = combinate (n,k) ;

print y;
```

The code above will create the following output:

1.0000	2.0000
1.0000	3.0000
1.0000	4.0000
2.0000	3.0000
2.0000	4.0000
3.0000	4.0000



See Also

[combined](#), [numCombinations](#)

combined

Purpose

Writes combinations of N things taken K at a time to a **GAUSS** data set.

Format

```
ret = combined(fname, vnames, N, K);
```

Input

<i>fname</i>	string, file name.
<i>vname</i>	1x1 or Kx1 string array, names of columns in data set. If 1x1 string, names will have column number appended. If null string, names will be X1, X2, ...
<i>N</i>	scalar.
<i>K</i>	scalar.

Output

<i>ret</i>	scalar, if data set was successfully written, <i>ret</i> = number of rows written to data set. Otherwise, one of the following: <div><div><i>0</i></div><div>file already exists.</div></div> <div><div><i>-1</i></div><div>data set couldn't be created.</div></div> <div><div><i>-n</i></div><div>the (<i>n</i>-1)th write to the data set failed.</div></div>
------------	---

combined

Remarks

The rows of the data set in `fname` contain sequences of the integers from 1 to `N` in combinations taken `K` at a time.

Example

```
//Note: The '$|' operator vertically concatenates strings
vnames = "Jim"$|"Harry"$|"Susan"$|"Wendy";

//Create a dataset file named 'couples', containing all
//combinations of the names in 'vnames' taken 2 at a time
k = 2;
m = combined("couples","Spouse 1"$|"Spouse 2", rows
(vnames),k);

print m "rows were written to the dataset";
```

After the above code,

```
6.0000 rows were written to the dataset
```

Continuing from the code above:

```
//Open the file written above
open f0 = "couples";

//Read in m=6 rows of the dataset into 'y'
y = readr(f0,m);
print "y = " y;

//Get the variable names from the dataset and assign them
//to 'names'
names = getnamef(f0);
f0=close(f0);
print $names ;
```

```

for i(1, rows(y),1);
    print vnames[y[i,.]]';
endfor;

```

will produce the following output:

```

y =
    1.0000000    2.0000000
    1.0000000    3.0000000
    1.0000000    4.0000000
    2.0000000    3.0000000
    2.0000000    4.0000000
    3.0000000    4.0000000

    Spouse 1    Spouse 2
        Jim      Harry
        Jim      Susan
        Jim      Wendy
    Harry      Susan
    Harry      Wendy
    Susan      Wendy

```

The first row of the print output 'Jim Harry' is the first and second element of *vnames*, because the first row of *y* is equal to '1 2'. The fourth row of the `print` output is 'Harry Susan', because the fourth row of *y* is '2 3' and 'Harry' is the second element of *vnames* while 'Susan' is the third element.

See Also

[combine](#), [numCombinations](#)

comlog

comlog

Purpose

Controls logging of interactive mode commands to a disk file.

Format

```
comlog [[file=filename]] [[on|off|reset]];
```

Input

filename literal or ^string.

Remarks

`comlog` on turns on command logging to the current file. If the file already exists, subsequent commands will be appended.

`comlog` off closes the log file and turns off command logging.

`comlog` reset turns on command logging to the current log file, resetting the log file by deleting any previous commands.

Interactive mode statements are always logged into the file specified in the `log_file` configuration variable, regardless of the state of `comlog`.

The command `comlog file= filename` selects the file but does not turn on logging.

The command `comlog off` will turn off logging. The filename will remain the same. A subsequent `comlog` on will cause logging to resume. A subsequent `comlog` reset will cause the existing contents of the log file to be destroyed and a new file created.

The command `comlog` by itself will cause the name and status of the current log file to be printed in the window.

compile

Purpose

Compiles a source file to a compiled code file. See also **COMPILER**, CHAPTER 1.

Format

```
compile source fname;
```

Input

<i>source</i>	literal or ^string, the name of the file to be compiled.
<i>fname</i>	literal or ^string, optional, the name of the file to be created. If not given, the file will have the same filename and path as <i>source</i> . It will have a <code>.gcg</code> extension.

Remarks

- The *source* file will be searched for in the *src_path* if the full path is not specified and it is not present in the current directory.
- The *source* file is a regular text file containing a **GAUSS** program. There can be references to global symbols, **Run-Time Library** references, etc.
- If there are `library` statements in *source*, they will be used during the compilation to locate various procedures and symbols used in the program. Since all of these library references are resolved at compile time, the `library` statements are not transferred to the compiled file. The compiled file can be run without activating any libraries.
- If you do not want extraneous stuff saved in the compiled image, put a `new` at

compile

the top of the *source* file or execute a [new](#) in interactive mode before compiling.

- The program saved in the compiled file can be run with the [run](#) command. If no extension is given, the [run](#) command will look for a file with the correct extension for the version of **GAUSS**. The *src_path* will be used to locate the file if the full path name is not given and it is not located on the current directory.
- When the compiled file is [run](#), all previous symbols and procedures are deleted before the program is loaded. It is therefore unnecessary to execute a [new](#) before [run](#)'ning a compiled file.
- If you want line number records in the compiled file you can put a [#lineson](#) statement in the *source* file or turn line tracking on from the main **GAUSS** menu, Tools->Preferences->Advanced.
- Don't try to include compiled files with [#include](#).
- **GAUSS** compiled files are platform and bit-size specific. For example, a file compiled with **GAUSS** for Windows 64-bit will not run under **GAUSS** for Windows 32-bit or on Linux 64-bit

Example

```
compile qxy.e;
```

In this example, the *src_path* would be searched for *qxy.e*, which would be compiled to a file called *qxy.gcg* on the same subdirectory *qxy.e* was found.

```
compile qxy.e xy;
```

In this example, the *src_path* would be searched for *qxy.e* which would be compiled to a file called *xy.gcg* on the current subdirectory.

See Also

[run](#), [use](#), [saveall](#)

complex

Purpose

Converts a pair of real matrices to a complex matrix.

Format

```
 $z = \text{complex}(xr, \ xi);$ 
```

Input

xr	$N \times K$ real matrix, the real elements of z .
xi	$N \times K$ real matrix or scalar, the imaginary elements of z .

Output

z	$N \times K$ complex matrix.
-----	------------------------------

Example

```
x = { 4 6,  
      9 8 };  
  
y = { 3 5,  
      1 7 };  
  
t = complex(x, y);
```

After the code above, t will be equal to:

con

```
4 + 3i 6 + 5i
9 + 1i 8 + 7i
```

See Also

[imag](#), [real](#)

con

Purpose

Requests input from the keyboard (console), and returns it in a matrix.

Format

```
x = con(r, c);
```

Input

<i>r</i>	scalar, row dimension of matrix.
<i>c</i>	scalar, column dimension of matrix.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix.
----------	-----------------------------

Remarks

con gets input from the active window. **GAUSS** will not "see" any input until you press ENTER, so follow each entry with an ENTER.

r and *c* may be any scalar-valued expressions. Nonintegers will be truncated to an integer.

c

If r and c are both set to 1, **con** will cause a question mark to appear in the window, indicating that it is waiting for a scalar input.

Otherwise, **con** will cause the following prompt to appear in the window:

```
- [1,1]
```

indicating that it is waiting for the [1,1] element of the matrix to be inputted. The - means that **con** will move horizontally through the matrix as you input the matrix elements. To change this or other options, or to move to another part of the matrix, use the following commands:

u	up one row	U	first row
d	down one row	D	last row
l	left one column	L	first column
r	right one column	R	last column
t	first element		
b	last element		
g #, #	goto element		
g #	goto element of vector		
h	move horizontally, default		
v	move vertically, default		
exttt\	move diagonally, default		
s	show size of matrix		

con

n	display element as numeric, default
c	display element as character
e	exp (1)
p	pi
.	missing value
?	show help screen
x	exit

If the desired matrix is 1xN or Nx1, then **con** will automatically exit after the last element has been entered, allowing you to input the vector quickly.

If the desired matrix is NxK, you will need to type 'x' to exit when you have finished entering the matrix data. If you exit before all elements have been entered, unspecified elements will be zeroed out.

Use a leading single quote for character input.

Example

```
n = con(1,1);  
print rndn(n,n);
```

If you enter 2 at the **con** generated prompt:

```
? 2
```

the code above will return a 2x2 random matrix, similar to:

```
-1.2505596      1.6322417  
-1.0894098      0.74763307
```

In this example, the **cond** function is used to obtain the size of a square matrix of Normal random variables which is to be printed out.

See Also

[cons](#), [let](#), [load](#)

cond

Purpose

Computes the condition number of a matrix using the singular value decomposition.

Format

```
c = cond(x);
```

Input

<i>x</i>	NxK matrix.
----------	-------------

Output

<i>c</i>	scalar, an estimate of the condition number of <i>x</i> . This equals the ratio of the largest singular value to the smallest. If the smallest singular value is zero or not all of the singular values can be computed, the return value is 10^{300} .
----------	---

conj

Example

```
x = { 4 2 6,  
      8 5 7,  
      3 8 9 };
```

```
y = conj(x);
```

will assign *y* to equal:

```
y = 9.8436943
```

Source

svd.src

conj

Purpose

Returns the complex conjugate of a matrix.

Format

```
y = conj(x);
```

Input

<i>x</i>	NxK matrix.
----------	-------------

Output

<i>y</i>	NxK matrix, the complex conjugate of <i>x</i> .
----------	---

Remarks

Compare **conj** with the transpose (') operator.

Example

```
x = { 1+9i    2,
      4+4i    5i,
      7i 8-2i };
y = conj(x);
```

$$x = \begin{bmatrix} 1 + 9i & 2 \\ 4 + 4i & 0 + 5i \\ 0 + 7i & 8 - 2i \end{bmatrix} \quad y = \begin{bmatrix} 1 - 9i & 2 \\ 4 - 4i & 0 - 5i \\ 0 - 7i & 8 + 2i \end{bmatrix}$$

cons

Purpose

Retrieves a character string from the keyboard.

Format

```
x = cons();
```

Output

x	string, the characters entered from the keyboard
-----	--

Remarks

x is assigned the value of a character string typed in at the keyboard. The program will pause to accept keyboard input. The maximum length of the string that can be entered

ConScore

is 254 characters. The program will resume execution when the ENTER key is pressed.

Example

```
x = cons ();
```

At the cursor enter:

```
probability
```

Now *x* will be equal to:

```
x = "probability";
```

See Also

[con](#)

ConScore

Purpose

Compute local score statistic and its probability for hypotheses involving parameters under constraints

Format

```
{ SL, SLprob } = ConScore(H, G, grad, a, b, c, d, bounds, psi);
```

Input

H

KxK matrix, Hessian of loglikelihood with respect to

c

	parameters.
G	$K \times K$ matrix, cross-product matrix of the first derivatives by observation. If not available set to H .
$grad$	$K \times 1$ vector, gradient of loglikelihood with respect to parameters.
a	$M \times K$ matrix, linear equality constraint coefficients.
b	$M \times 1$ vector, linear equality constraint constants.
	These arguments specify the linear equality constraints of the following type:
	$a * X = b$
	where X is the $K \times 1$ parameter vector.
c	$M \times K$ matrix, linear inequality constraint coefficients.
d	$M \times 1$ vector, linear inequality constraint constants.
	These arguments specify the linear inequality constraints of the following type:
	$c * X \geq d$
	where X is the $K \times 1$ parameter vector.
$bounds$	$K \times 2$ matrix, bounds on parameters. The first column contains the lower bounds, and the second column the upper bounds.
psi	indices of the set of parameters in the hypothesis.

Output

SL	scalar, local score statistic of hypothesis.
$SLprob$	scalar, probability of SL .

ConScore

Remarks

ConScore computes the local score statistic for the hypothesis $H(\theta) = 0$ vs. $H(\theta) \geq 0$, where θ is the vector of estimated parameters, and $H()$ is a constraint function of the parameters.

First, the model with $H(\theta) = 0$ is estimated, and the Hessian and optionally the cross-product of the derivatives is computed. Also, the gradient vector is computed.

Next, the constraint arguments are set to $H(\theta) \geq 0$.

Example

This example is from Silvapulle and Sen, *Constrained Statistical Inference*, page 181-3. It computes the local score statistic and probability for an ARCH model. It tests the null hypothesis of no arch effects against the alternative of arch effects subject to their being constrained to be positive.

The Hessian, H, cross-product matrix, G, and the gradient vector, grad, are generated by an estimation using **sqpSolveMT** where the model is an ARCH model with the arch parameters constrained to be zero.

```
#include sqpsolveMT.sdf
/* data */
struct DS d0;
d0 = reshape(dsCreate,2,1);

load z0[] = aoi.asc;
z = packr(lagn(251*ln(trimr(z0,1,0)./trimr
(z0,0,1)),0|1|2|3|4));
d0[1].dataMatrix = z[:,1];
d0[2].dataMatrix = z[:,2:5];

/* control structure */
struct sqpsolveMTControl c0;
c0 = sqpSolveMTControlCreate;
```

```

/* constraints setting arch parameter equal
to zero for H(theta) = 0 */

c0.A = zeros(3,6) ~ eye(3);
c0.B = zeros(3,1);

c0.covType = 2; /* causes cross-product of
                 Jacobian to be computed which
                 is needed for ConScore */

struct PV p0;
p0 = pvPack(pvCreate,.08999, "constant");
p0 = pvPack(p0,.25167|-0.12599|.09164|.07517,
            "phi");
p0 = pvPack(p0,3.22713, "omega");
p0 = pvPack(p0,0|0|0, "arch");

struct sqpsolvemtOut out0;
out0 = sqpsolvemt(&lpr,p0,d0,c0);

/* set up constraints for H(theta) >= 0 */
bounds = { -1e256 1e256,
            -1e256 1e256,
            -1e256 1e256,
            -1e256 1e256,
            -1e256 1e256,
            -1e256 1e256,
            0 1e256,
            0 1e256,
            0 1e256 };
H = out0.hessian;
G = out0.xproduct;
// minus because -logl in log-likelihood

```

continue

```
grad = -out0.gradient;

psi = { 7, 8, 9 };

{ SL, SLprob } = ConScore(H,G,grad,0,0,0,0,bounds,psi);
```

will assign the variables *SL* and *SLprob* as follows:

```
SL = 3.8605086

SLprob = 0.10410000
```

Source

hypotest.src

continue

Purpose

Jumps to the top of a **do** or **for** loop.

Format

```
continue;
```

Example

```
rndseed 8989;
x = rndn(4,4);
//Loop through each row of 'x' using 'r' as the loop
//counter
for r(1, rows(x), 1);
```

C

```

//Loop through each element in our current row
for c(1, cols(x), 1); /* continue jumps here */
    //If we are on the diagonal skip the rest of the
    //inner loop
    if c == r;
        continue;
    endif;
    //Set the non-diagonal elements to 0
    x[r,c] = 0;
endfor;
endfor;

```

Before the loops, x looks like:

0.01055555	-0.045969063	0.12701699	
1.6454828			
1.2380373	0.53988699	1.1556776	-
0.53575797			
0.14056238	0.11221419	0.91500922	-
2.2910169			
1.4278412	-0.96476892	0.22852569	-
1.6014053			

After the loops above, x looks like:

0.01055555	0.00000000	0.00000000	
0.00000000			
0.00000000	0.53988699	0.00000000	
0.00000000			
0.00000000	0.00000000	0.91500922	
0.00000000			
0.00000000	0.00000000	0.00000000	-
1.6014053			

contour

Remarks

This command works just as in C.

contour

Purpose

Graphs a matrix of contour data. Note: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
contour(x, y, z);
```

Input

<i>x</i>	1xK vector, the X axis data. K must be odd.
<i>y</i>	Nx1 vector, the Y axis data. N must be odd.
<i>z</i>	NxK matrix, the matrix of height data to be plotted.

Global Input

<i>_plev</i>	Kx1 vector, user-defined contour levels for contour . Default 0.
<i>_pzclr</i>	Nx1 or Nx2 vector. This controls the Z level colors. See surface for a complete description of how to set this global.

C

Remarks

A vector of evenly spaced contour levels will be generated automatically from the `z` matrix data. Each contour level will be labeled. For unlabeled contours, use **zticks**.

To specify a vector of your own unequal contour levels, set the vector `_plev` before calling **contour**.

To specify your own evenly spaced contour levels, see **zticks**.

Source

`pcontour.src`

See Also

[surface](#)

conv

Purpose

Computes the convolution of two vectors.

Format

```
c = conv(b, x, f, l);
```

Input

<code>b</code>	Nx1 vector.
<code>x</code>	Lx1 vector.
<code>f</code>	scalar, the first convolution to compute.
<code>l</code>	scalar, the last convolution to compute.

convtsatostr

Output

c	Qx1 result, where: $Q = (l - f + 1)$ If f is 0, the first to the l 'th convolutions are computed. If l is 0, the f 'th to the last convolutions are computed. If f and l are both zero, all the convolutions are computed.
-----	---

Remarks

If x and b are vectors of polynomial coefficients, this is the same as multiplying the two polynomials.

See Also

[polymult](#)

convtsatostr

Purpose

Converts a 1x1 string array to a string.
--

Format

$str = \text{convtsatostr}(sa);$

Input

sa	1x1 string array.
------	-------------------

Output

str	string, sa converted to a string.
-------	-------------------------------------

See Also

[convertstrtosa](#)

convertstrtosa

Purpose

Converts a string to a 1x1 string array.

Format

```
sa = convertstrtosa(str);
```

Input

str	string.
-----	---------

Output

sa	1x1 string array, str converted to a string array.
----	--

Example

```
str = "This is a string";
z = convertstrtosa(str);
```

You can check the types of your variables by viewing them on the **GAUSS** data page, or by using the [show](#) command. If the code above was executed at startup, running the [show](#) command would return:

24 bytes	str	STRING
16 char		

corrmm,corrvc,corrxx

40 bytes	z	STRING ARRAY
1,1		

See Also

[convertsatostr](#)

corrmm,corrvc,corrxx

Purpose

Computes an unbiased estimate of a correlation matrix.

Format

```
cx = corrmm(m);
cx = corrvc(vc);
cx = corrxx(x);
```

Input

m	KxK moment ($x'x$) matrix. A constant term MUST have been the first variable when the moment matrix was computed.
vc	KxK variance-covariance matrix (of data or parameters).
x	NxK matrix of data.

Output

cx	PxP correlation matrix. For corrmm , P = K-1. For corrvc and corrxx , P = K.
----	---

Example

```

rndseed    8989;
x1 = rndn(3,3);
x2 = ones(3,1)~x1;
print "x1 :" x1 ;
print "x2 :" x2;

```

After the above code, *x1* and *x2* look like:

```

x1 :
      0.010555555      -0.045969063      0.12701699
      1.6454828        1.2380373        0.53988699
      1.1556776       -0.53575797       0.14056238

x2 :
      1.0000000      0.010555555     -0.045969063
0.12701699
      1.0000000      1.6454828        1.2380373
0.53988699
      1.0000000      1.1556776       -0.53575797
0.14056238

```

Continuing from above code,

```

print "corrxx(x1) :" corrxx(x1);
print "corrmm(x2'x2) :" corrmm(x2'x2);
print "corrvc(varCovMS(x2'x2)):" corrvc(varCovMS(x2'x2));

```

After the above code,

```

corrxx(x1) :
      1.0000000      0.52196856      0.75039768

```

corrms,corrxs

	0.52196856	1.0000000	0.95548228
	0.75039768	0.95548228	1.0000000
corrmm(x2'x2) :			
	1.0000000	0.52196856	0.75039768
	0.52196856	1.0000000	0.95548228
	0.75039768	0.95548228	1.0000000
corrvc(varCovMS(x2'x2)) :			
	1.0000000	0.52196856	0.75039768
	0.52196856	1.0000000	0.95548228
	0.75039768	0.95548228	1.0000000

Remarks

The correlation matrix is the standardized version of the unbiased estimator of the population variance-covariance matrix. It is computed using the moment matrix of deviations about the mean divided by the number of observations minus one $N - 1$. For the observed correlation/covariance matrix which uses N rather than $N - 1$, see **corrms** and **corrxs**.

Source

corr.src

See Also

[momentd](#), [corrms](#), [corrxs](#), [varCovX](#), [varCovM](#)

corrms,corrxs

Purpose

Computes the observed correlation matrix.

Format

```
CX = corrms(m);  
CX = corrxs(x);
```

Input

<i>m</i>	KxK moment ($x'x$) matrix. A constant term MUST have been the first variable when the moment matrix was computed.
<i>x</i>	NxK matrix of data.

Output

<i>CX</i>	PxP correlation matrix. For corrms , $P = K-1$. For corrxs , $P = K$.
-----------	---

Example

```
randseed    8989;  
x1 = rndn(3,3);  
x2 = ones(3,1)~x1;  
print "x1 : " x1 ;  
print "x2 : " x2;
```

After the above code, *x1* and *x2* look like:

```
x1 :  
      0.01055555      -0.04596906      0.12701699  
      1.6454828       1.2380373      0.53988699  
      1.1556776      -0.53575797      0.14056238  
x2 :  
      0.01055555      -0.04596906      0.12701699  
      1.6454828       1.2380373      0.53988699  
      1.1556776      -0.53575797      0.14056238
```

corrms,corrxs

```
          1.0000000      0.01055555      -0.045969063
0.12701699
          1.0000000      1.6454828      1.2380373
0.53988699
          1.0000000      1.1556776      -0.53575797
0.14056238
```

Continuing from above code,

```
print "corrxs(x1)  :" corrxs(x1) ;
print "corrms(x2'x2)  :" corrms(x2'x2) ;
```

After the above code,

```
corrxs(x1)  :
          1.0000000      0.52196856      0.75039768
          0.52196856      1.0000000      0.95548228
          0.75039768      0.95548228      1.0000000
corrms(x2'x2)  :
          1.0000000      0.52196856      0.75039768
          0.52196856      1.0000000      0.95548228
          0.75039768      0.95548228      1.0000000
```

Remarks

The correlation matrix is the standardized version of the correlation/covariance matrix computed from the input data, that is, it divides the sample size, N , rather than $N - 1$. For an unbiased estimate correlation/covariance matrix which uses $N - 1$, use **corrmm** or **corrxx**.

Source

corrs.src

See Also

[momentd](#), [corrmm](#), [corrxx](#), [varCovX](#), [varCovM](#)

COS

Purpose

Returns the cosine of its argument.

Format

```
y = cos(x);
```

Input

x	NxK matrix.
---	-------------

Output

y	NxK matrix containing the cosines of the elements of x.
---	---

Remarks

For real matrices, x should contain angles measured in radians.

To convert degrees to radians, multiply the degrees by $\pi/180$.

Example

```
//Create a sequence starting at 0 and increasing by pi/4  
x = seqa(0, pi/4, 5);
```

cosh

```
y = cos(x);
```

	0.0000	1.0000
	0.7854	0.7071
x = 1.5708	y = 0.0000	
	2.3562	-0.7071
	3.1416	-1.0000

See Also

[atan](#), [atan2](#), [pi](#)

cosh

Purpose

Computes the hyperbolic cosine.

Format

```
y = cosh(x);
```

Input

x NxK matrix.

Output

y NxK matrix containing the hyperbolic cosines of the elements of x.

Example

```
x = { -0.5, -0.25, 0, 0.25, 0.5, 1 };
x = x * pi;
y = cosh(x);
```

```
      -1.5708      2.5092
      -0.7854      1.3246
x = 0.0000  y = 1.0000
      0.7854      1.3246
      1.5708      2.5092
      3.1416     11.5920
```

Source

trig.src

counts

Purpose

Counts the numbers of elements of a vector that fall into specified ranges.

Format

```
c = counts(x, v);
```

Input

x	$N \times 1$ vector containing the numbers to be counted.
v	$P \times 1$ vector containing breakpoints specifying the ranges within which counts are to be made. The vector v MUST be sorted in ascending order.

counts

Output

c

$P \times 1$ vector, the counts of the elements of x that fall into the regions:

$$\begin{aligned} & x \leq v[1], \\ & v[1] < x \leq v[2], \\ & \cdot \\ & \cdot \\ & \cdot \\ & v[p-1] < x \leq v[p] \end{aligned}$$

Remarks

If the maximum value of x is greater than the last element (the maximum value) of v , the sum of the elements of the result, c , will be less than N , the total number of elements in x .

If

$$\begin{array}{rcl} & 1 & \\ & 2 & \\ & 3 & \\ & 4 & 4 \\ x = 5 & v = 5 & \\ & 6 & 8 \\ & 7 & \\ & 8 & \\ & 9 & \end{array}$$

then

$$\begin{array}{rcl} & 4 & \\ c = 1 & & \\ & 3 & \end{array}$$

c

The first category can be a missing value if you need to count missings directly. Also $+\infty$ or $-\infty$ are allowed as breakpoints. The missing value must be the first breakpoint if it is included as a breakpoint and infinities must be in the proper location depending on their sign. $-\infty$ must be in the $[2,1]$ element of the breakpoint vector if there is a missing value as a category as well, otherwise it has to be in the $[1,1]$ element. If $+\infty$ is included, it must be the last element of the breakpoint vector.

Example

```
x = { 1.5, 3, 5, 4, 1, 3 };  
v = { 0, 2, 4 };  
c = countwts(x,v);
```

1.5		
3	0	0
x = 2	v = 2	c = 2
4	4	3
1		
3		

countwts

Purpose

Returns a weighted count of the numbers of elements of a vector that fall into specified ranges.

Format

```
c = countwts(x, v, w);
```

countwts

Input

x	$N \times 1$ vector, the numbers to be counted.
v	$P \times 1$ vector, the breakpoints specifying the ranges within which counts are to be made. This MUST be sorted in ascending order (lowest to highest).
w	$N \times 1$ vector, containing weights.

Output

c	<p>$P \times 1$ vector containing the weighted counts of the elements of x that fall into the regions:</p> <div>$\begin{aligned} & x < v[1], \\ v[1] \leq x < v[2], \\ & \cdot \\ & \cdot \\ & \cdot \\ v[p-1] \leq x < v[p] \end{aligned}$</div> <p>That is, when $x[i]$ falls into region j, the weight $w[i]$ is added to the jth counter.</p>
-----	---

Remarks

c	<p>If any elements of x are greater than the last element of v, they will not be counted.</p> <p>Missing values are not counted unless there is a missing in v. A missing value in v MUST be the first element in v.</p>
-----	--

Example

	$x = \{ 1, 3, 2, 4, 1, 3 \};$
--	-------------------------------

```
w = { .25, 1, .333, .1, .25, 1 };
v = { 0, 1, 2, 3, 4 };
c = countwts(x,v,w);
```

```
0.0000000
0.5000000
c = 0.3330000
2.0000000
0.1000000
```

create

Purpose

Creates and opens a **GAUSS** data set for subsequent writing.

Format

```
create [[vflag]] [[-w32]] [[complex]] fh = filename with
vnames, col, dtyp, vtyp;
create [[vflag]] [[-w32]] [[complex]] fh = filename using comfile;
```

Input

<i>vflag</i>	literal, version flag.
-v89	obsoleted, use -v96.
-v92	obsoleted, use -v96.
-v96	supported on all platforms.

For details on the various versions, see **FOREIGN LANGUAGE INTERFACE**, CHAPTER 1. The default format can be specified in `gauss.cfg` by setting the `dat_fmt_version` configuration variable. The

create

	default, <code>v96</code> , should be used.
<i>filename</i>	literal or <code>^string</code> <i>filename</i> is the name to be given to the file on the disk. The name can include a path if the directory to be used is not the current directory. This file will automatically be given the extension <code>.dat</code> . If an extension is specified, the <code>.dat</code> will be overridden. If the name of the file is to be taken from a string variable, the name of the string must be preceded by the <code>^</code> (caret) operator.
<code>create... with...</code>	
<i>vnames</i>	literal or <code>^string</code> or <code>^character matrix</code> . <i>vnames</i> controls the names to be given to the columns of the data file. If the names are to be taken from a string or character matrix, the <code>^</code> (caret) operator must be placed before the name of the string or character matrix. The number of columns parameter, <i>col</i> , also has an effect on the way the names will be created. See below and see the examples for details on the ways names are assigned to a data file.
<i>col</i>	scalar expression. <i>col</i> is a scalar expression containing the number of columns in the data file. If <i>col</i> is 0, the number of columns will be controlled by the contents of <i>vnames</i> . If <i>col</i> is positive, the file will contain <i>col</i> columns and the names to be given each column will be created as necessary depending on the <i>vnames</i> parameter. See the examples.
<i>dtyp</i>	scalar expression. <i>dtyp</i> is the precision used to store the data. This is a scalar expression containing 2, 4, or 8, which is the number of bytes per element.

	2	signed integer		
	4	single precision		
	8	double precision		

Data Type	Digits		Range	
integer	4	-32768	< X <	32768
single	6-7	8.43×10^{-37}	< X <	$3.37 \times 10^{+38}$
double	15-16	4.19×10^{-307}	< X <	1.67×10^{308}

If the integer type is specified, numbers will be rounded to the nearest integer as they are written to the data set. If the data to be written to the file contains character data, the precision must be 8 or the character information will be lost.

vtyp matrix, types of variables.

The types of the variables in the data set. If **rows** (*vtyp*)***cols**(*vtyp*) < *col*, only the first element is used. Otherwise nonzero elements indicate a numeric variable and zero elements indicate character variables.

create... using...

comfile literal or ^string.

comfile is the name of a command file that contains the information needed to create the file. The default extension for the command file is *.gcf*, which can be overridden.

There are three possible commands in this file:

```
numvar  n str;
```

create

```
outvar  varlist;  
outtyp  dtyp;
```

numvar and **outvar** are alternate ways of specifying the number and names of the variables in the data set to be created.

When **numvar** is used, *n* is a constant which specifies the number of variables (columns) in the data file and *str* is a string literal specifying the prefix to be given to all the variables. Thus:

```
numvar 10 xx;
```

says that there are 10 variables and that they are to be named *xx01* through *xx10*. The numeric part of the names will be padded on the left with zeros as necessary so the names will sort correctly:

xx1,	...	xx9	1-9 names
xx01,	...	xx10	10-99 names
xx001,	...	xx100	100-999 names
xx0001,	...	xx1000	1000-8100 names

If *str* is omitted, the variable prefix will be "X".

When **outvar** is used, *varlist* is a list of variable names, separated by spaces or commas. For instance:

```
outvar x1, x2, zed;
```

specifies that there are to be 3 variables per row of the data set, and that they are to be named *X1*, *X2*, *ZED*, in

that order.

outtyp specifies the precision. It can be a constant: 2, 4, or 8, or it can be a literal: I, F, or D. For an explanation of the available data types, see *dtyp* in [create...with...](#) previously.

The **outtyp** statement does not have to be included. If it is not, then all data will be stored in 4 bytes as single precision floating point numbers.

Output

fh

scalar.

fh is the file handle which will be used by most commands to refer to the file within **GAUSS**. This file handle is actually a scalar containing an integer value that uniquely identifies each file. This value is assigned by **GAUSS** when the [create](#) (or [open](#)) command is executed.

Remarks

If the *complex* flag is included, the new data set will be initialized to store complex number data. Complex data is stored a row at a time, with the real and imaginary halves interleaved, element by element.

The *-w32* flag is an optimization for Windows. It is ignored on all other platforms. **GAUSS** 7.0 and later use Windows system file write commands that support 64-bit file sizes. These commands are slower on Windows XP than the 32-bit file write commands that were used in **GAUSS** 6.0 and earlier. If you include the *-w32* flag, successive writes to the file indicated by *fh* will use 32-bit Windows write commands, which will be faster on Windows XP. Note, however, that the *-w32* flag does not support 64-bit file sizes.

create

Example

```
let vnames = age sex educat wage occ;
create fl = simdat with ^vnames,0,8;

obs = 0; nr = 1000;
do while obs < 10000;
    data = rndn(nr, colsf(fl));
    if writer(fl,data) /= nr;
        print "Disk Full";
    end;
endif;
obs = obs+nr;
endo;

closeall fl;
```

This example uses `create... with...` to create a double precision data file called `simdat.dat` on the default drive with 5 columns. The **writer** command is used to write 10000 rows of Normal random numbers into the file. The variables (columns) will be named: *AGE*, *SEX*, *EDUCAT*, *WAGE*, *OCC*.

Here are some examples of the variable names that will result when using a character vector of names in the argument to the `create` function.

```
vnames = { AGE PAY SEX JOB };
typ = { 1, 1, 0, 0 };
create fp = mydata with ^vnames,0,8,typ;
```

The names in the this example will be: *AGE*, *PAY*, *SEX*, *JOB*.

AGE and *PAY* are numeric variables, *SEX* and *JOB* are character variables.

```
create fp = mydata with ^vnames,3,2;
```

The names will be: *AGE*, *PAY*, *SEX*.

```
create fp = mydata with ^vnames,8,2;
```

The names will now be: *AGE*, *PAY*, *SEX*, *JOB1*, *JOB2*, *JOB3*, *JOB4*, *JOB5*.

If a literal is used for the *vnames* parameter, the number of columns should be explicitly given in the *col* parameter and the names will be created as follows:

```
create fp = mydata with var,4,2;
```

Giving the names: *VAR1*, *VAR2*, *VAR3*, *VAR4*.

The next example assumes a command file called *comd.gcf* containing the following lines, created using a text editor:

```
outvar age, pay, sex;
outtyp i;
```

Then the following program could be used to write 100 rows of random integers into a file called *smpl.dat* in the subdirectory called */gauss/data*:

```
filename = "/gauss/data/smpl";
create fh = ^filename using comd;
x = rndn(100,3)*10;
if writer(fh,x) /= rows(x);
    print "Disk Full";
end;
endif;
closeall fh;
```

For platforms using the backslash as a path separator, remember that two backslashes ("\\") are required to enter one backslash inside of double quotes. This is because a backslash is the escape character used to embed special characters in strings.

See Also

[datacreate](#), [datacreatecomplex](#), [open](#), [readr](#), [writer](#), [eof](#), [close](#), [output](#), [iscplxf](#)

crossprd

crossprd

Purpose

Computes the cross-products (vector products) of sets of 3x1 vectors.

Format

$z = \text{crossprd}(x, y);$

Input

x	3xK matrix, each column is treated as a 3x1 vector.
y	3xK matrix, each column is treated as a 3x1 vector.

Output

z	3xK matrix, each column is the cross-product (sometimes called vector product) of the corresponding columns of x and y .
-----	--

Remarks

The cross-product vector z is orthogonal to both x and y . **sumc**($x .* z$) and **sumc**($y .* z$) will be Kx1 vectors, all of whose elements are 0 (except for rounding error).

Example

```
x = { 10  4,
      11 13,
      14 13 };
y = { 3 11,
      5 12,
```

```
    7  9  };  
z = crossprd(x,y);
```

After the above code,

```
    7  -39  
z = -28  107  
    17  -95
```

Source

crossprd.src

crout

Purpose

Computes the Crout decomposition of a square matrix without row pivoting, such that: $X = LU$.

Format

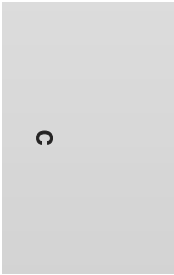
```
y = crout(x);
```

Input

x	NxN square nonsingular matrix.
---	--------------------------------

Output

y	NxN matrix containing the lower (<i>L</i>) and upper (<i>U</i>) matrices of the Crout decomposition of <i>x</i> . The main diagonal of <i>y</i> is the main diagonal of the lower matrix <i>L</i> . The upper matrix has an implicit main diagonal of
---	---



crout

ones. Use **lowmat** and **upmat1** to extract the L and U matrices from y .

Remarks

Since it does not do row pivoting, it is intended primarily for teaching purposes. See **croutp** for a decomposition with pivoting.

Example

```
X = { 1 2 -1,
      2 3 -2,
      1 -2 1 };

//Perform crout decomposition
y = crout(x);

//Extract lower triangle of 'y' and assign it to 'L'
L = lowmat(y);

//Extract upper triangle of 'y', fill the diagonal with
//ones and assign it to 'U'
U = upmat1(y);
```

After the code above:

1.0	2.0	-1.0	1.0	0.0	0.0	1.0	2.0	-1.0
2.0	-1.0	0.0	2.0	-1.0	0.0	0.0	1.0	0.0
1.0	-4.0	2.0	1.0	-4.0	2.0	0.0	0.0	1.0

See Also

[croutp](#), [chol](#), [lowmat](#), [lowmat1](#), [lu](#), [upmat](#), [upmat1](#)

croutp

Purpose

Computes the Crout decomposition of a square matrix with partial (row) pivoting.

Format

```
 $y = \text{croutp}(x);$ 
```

Input

x	$N \times N$ square nonsingular matrix.
-----	---

Output

y	$(N+1) \times N$ matrix containing the lower (L) and upper (U) matrices of the Crout decomposition of a permuted x . The $N+1$ row of the matrix y gives the row order of the y matrix. The matrix must be reordered prior to extracting the L and U matrices. Use lowmat and upmat1 to extract the L and U matrices from the reordered y matrix.
-----	---

Example

This example illustrates a procedure for extracting L and U of the permuted x matrix. It continues by sorting the result of LU to compare with the original matrix x .

```
x = { 1 2 -1,  
      2 3 -2,  
      1 -2 1 };
```

croutp

```
y = croutp(x);
```

If we view 'y', we will see:

```
      1.0000      0.5000      0.28571
y =  2.0000      1.5000     -1.0000
      1.0000     -3.5000     -0.57142
      2.0000      3.0000      1.0000
```

```
//This bottom row is the permutation index vector
//Calculate how many rows in 'y'
r = rows(y);

//Extract the index row and transpose it into a column
//vector
index = y[r,:].';
```

Viewing 'indx' will reveal:

```
      2
index = 3
      1
```

```
//Rearrange the rows of 'y' based upon the index vector
z = y[index,:];

// obtain L and U of permuted matrix X
L = lowmat(z);
U = upmat1(z);

//Horizontally concatenate the index vector and the product
//of L*U then pass that result into the 'sortc' function
//which will sort this result based upon the first column
//(which is the index vector)
```

```
q = sortc(index~(L*U),1);

//Remove the index vector, which we added by way of
//horizontal concatenation in the statement just above
x2 = q[:,2:cols(q)];
```

Now at the end of this example, `x2` is equal to `x`.

See Also

[crou](#), [chol](#), [lowmat](#), [lowmat1](#), [upmat](#), [upmat1](#)

csrcol,csrlin

Purpose

Returns the position of the cursor.

Format

```
y = csrcol;
y = csrlin;
```

Output

<code>y</code>	scalar, row or column value.
----------------	------------------------------

Remarks

`y` will contain the current column or row position of the cursor on the screen. The upper left corner is (1,1).

csrcol returns the column position of the cursor. **csrlin** returns the row position.

csvReadM

The `locate` command allows the cursor to be positioned at a specific row and column.

`csrcol` returns the cursor column with respect to the current output line, i.e., it will return the same value whether the text is wrapped or not. `csrln` returns the cursor line with respect to the top line in the window.

Example

```
r = csrln;  
c = csrcol;  
  
//Clear the program input/output window  
cls;  
  
//Re-position the cursor to its location before the program  
//input/output window was cleared  
locate r,c;
```

In this example the screen is cleared without affecting the cursor position.

See Also

[cls](#), [locate](#)

csvReadM

Purpose

Reads data from CSV file into a **GAUSS** matrix.

Format

```
mat = csvReadM(file);  
mat = csvReadM(file, row_range);  
mat = csvReadM(file, row_range, col_range);  
mat = csvReadM(file, row_range, col_range, delimiter);
```

Input

<i>file</i>	string, name of CSV file.
<i>row_range</i>	Optional input: scalar, or 2x1 matrix. The first element of <i>row_range</i> will specify the first row of the file to read. If there is a second element in <i>row_range</i> , it will specify the last row to read from the file. If there is no second element in <i>row_range</i> , GAUSS will read to the end of the file. If <i>row_range</i> is not passed in, all rows will be read from the file. Default = 1.
<i>col_range</i>	Optional input: scalar, or 2x1 matrix. The first element of <i>col_range</i> will specify the first column of the file to read. If there is a second element in <i>col_range</i> , it will specify the last column to read from the file. If there is no second element in <i>col_range</i> , GAUSS will read to the end of the file. If <i>col_range</i> is not passed in, all columns will be read from the file. Default = 1.
<i>delimiter</i>	Optional input: string. The character used to separate elements in the file. Examples include: <ul style="list-style-type: none">• space " "

csvReadM

- tab `"\t"`
- semi-colon `";"`
- comma `","` (Default)

Output

mat

matrix, data read from the CSV file.

Examples

Example 1: Basic Example

Read all contents from the file `housing.csv` located in your GAUSS examples directory.

```
//Get file name with full path
file = getGAUSSHome() $+ "examples/housing.csv";

//Read entire contents of 'housing.csv'
housing = csvReadM(file);

//Print the first 5 rows of all columns
print housing[1:5, .];
```

The code above will produce the following output. Notice that the first row contains all missing values. This is because the first row of the file `housing.csv` contains a header. **csvReadM** reads in textual data as missing values by default. You can easily remove any rows that contain all missing values with the function *packr* (which stands for "pack rows").

.
3104.00	4.00	2.00	0.00	279.90	
2048.00					

1173.00	2.00	1.00	0.00	146.50
912.00				
3076.00	4.00	2.00	0.00	237.70
1654.00				
1608.00	3.00	2.00	0.00	200.00
2068.00				

Example 2: Skip the header

In the previous example, we read the header in as numeric data and got missing values. In this example, we will skip the first row to avoid reading the header as numeric data.

```
//Create file name with full path
file = getGAUSSHome() %+ "examples/housing.csv";

//Row range will be from line 2 to the end of the file
//If no end to the range is specified, GAUSS will read to
the end of the file
row_range = 2;

//Load the data from row 2 to the end of the file into
'housing'
housing = csvReadM(file, row_range);

//Print the first 5 rows of all columns
print housing[1:5,.];
```

After the code above, *housing* should equal:

3104.00	4.00	2.00	0.00	279.90
2048.00				
1173.00	2.00	1.00	0.00	146.50
912.00				
3076.00	4.00	2.00	0.00	237.70

csvReadM

1654.00				
1608.00	3.00	2.00	0.00	200.00
2068.00				
1454.00	3.00	3.00	0.00	159.90
1477.00				

Example 3: Read from a row range

```
//Create file name with full path
file = getGAUSSHome() %+ "examples/housing.csv";

//Row range will be from line 3 to line 5
row_range = { 3, 5 };

//Load the data from rows 3 to 5 of the file into 'housing'
housing = csvReadM(file, row_range);

//Print the entire contents of the variable 'housing'
print housing;
```

After the code above, *housing* should equal:

1173.00	2.00	1.00	0.00	146.50
912.00				
3076.00	4.00	2.00	0.00	237.70
1654.00				
1608.00	3.00	2.00	0.00	200.00
2068.00				

Example 4: Read all rows of a range of columns

```
//Create file name with full path
```



```

file = getGAUSSHome() $+ "examples/housing.csv";

//Row range from the first line to the end of the file
row_range = 1;

//Read only columns 2 through 4
col_range = { 2, 4 };

//Load the data from columns 2 through 4 into 'x'
x = csvReadM(file, row_range, col_range);

```

Example 5: Read all rows of one specific column

```

//Create file name with full path
file = getGAUSSHome() $+ "examples/housing.csv";

//Row range from the first line to the end of the file
row_range = 1;

//Read only the 3rd column
col_range = { 3, 3 };

//Load the data from the 3rd column into 'x'
x = csvReadM(file, row_range, col_range);

```

Example 6: Read all rows and all cols, with specified delimiter

Enter 1 for the *row_range* and *col_range* if you want to read all contents of a file, but need to specify the field delimiter.

```

x = csvReadM("myfile.csv", 1, 1, ";");

```

Example 7: Specify full path to file

Windows: Notice that double backslashes are needed inside of a string.

csvReadSA

```
x = csvReadM("C:\\mydata\\myfile.csv");
```

Mac

```
x = csvReadM("/Users/MyUserName/myfile.csv");
```

Linux

```
x = csvReadM("/home/my_user/myfile.csv");
```

Remarks

The standard input stream (stdin) can be read with **csvReadM** by passing in `__STDIN` as the filename input. Note that `__STDIN` should not be passed as a string, surrounded by quotes. Correct usage is shown below:

```
x = csvReadM(__STDIN);
```

See Also

[csvReadSA](#), [xlsWrite](#), [xlsWriteM](#), [xlsWriteSA](#), [xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

csvReadSA

Purpose

Reads data from CSV file into a **GAUSS** string array.

Format

```
s = csvReadSA(file);  
s = csvReadSA(file, row_range);  
s = csvReadSA(file, row_range, col_range);  
s = csvReadSA(file, row_range, col_range, delimiter);
```

Input

<i>file</i>	string, name of CSV file.
<i>row_range</i>	Optional input: scalar, or 2x1 matrix. The first element of <i>row_range</i> will specify the first row of the file to read. If there is a second element in <i>row_range</i> , it will specify the last row to read from the file. If there is no second element in <i>row_range</i> , GAUSS will read to the end of the file. If <i>row_range</i> is not passed in, all rows will be read from the file. Default = 1.
<i>col_range</i>	Optional input: scalar, or 2x1 matrix. The first element of <i>col_range</i> will specify the first column of the file to read. If there is a second element in <i>col_range</i> , it will specify the last column to read from the file. If there is no second element in <i>col_range</i> , GAUSS will read to the end of the file. If <i>col_range</i> is not passed in, all columns will be read from the file. Default = 1.
<i>delimiter</i>	Optional input: string. The character used to separate elements in the file. Examples include: <ul style="list-style-type: none">• space " "• tab "\t"• semi-colon ";• comma "," (Default)

Output

<i>s</i>	string array, data read from the CSV file.
<i>row_range</i>	Optional input: scalar, or 2x1 matrix. The first element of <i>row_range</i> will specify the first row of the file to read. If there is a second element in <i>row_range</i> , it will specify the last row to read from the file. If there is no second element in <i>row_range</i> , GAUSS will read to the end of the file. If <i>row_range</i> is not passed in, all rows will be read from the file. Default = 1.
<i>col_range</i>	Optional input: scalar, or 2x1 matrix. The first element of <i>col_range</i> will specify the first column of the file to read. If there is a second element in <i>col_range</i> , it will specify the last column to read from the file. If there is no second element in <i>col_range</i> , GAUSS will read to the end of the file. If <i>col_range</i> is not passed in, all columns will be read from the file. Default = 1.
<i>delimiter</i>	Optional input: string. The character used to separate elements in the file. Examples include: <ul style="list-style-type: none">• space "• tab "\t"• semi-colon ";• comma "," (Default)

Examples

Example 1: Basic Example

Read all contents from the file `myfile.csv` located in your current GAUSS working directory.

```
s = csvReadSA("myfile.csv");
```

Example 2: Read From a Row Range

```
//Row range will be from line 1 to line 25
row_range = { 1, 25 };

//Load the data from rows 1 to 25 into 's'
s = csvReadSA("myfile.csv", row_range);
```

Example 3: Read all rows from a range of columns

```
//Row range from the first line to the end of the file
row_range = 1;

//Read only columns 2 through 7
col_range = { 2, 7 };

//Load the data from columns 2 through 7 into 's'
s = csvReadSA("myfile.csv", row_range, col_range);
```

Example 4: Read all rows from one column

```
//Row range from the first line to the end of the file
row_range = 1;

//Read only column 4
col_range = { 4, 4 };
```

csvWriteM

```
//Load the data from column 4 into 's'  
s = csvReadSA("myfile.csv", row_range, col_range);
```

Example 5: Specify full path to file

Windows: Notice that double backslashes are needed inside of a string.

```
s = csvReadSA("C:\\mydata\\myfile.csv");
```

Mac

```
s = csvReadSA("/Users/MyUserName/myfile.csv");
```

Linux

```
s = csvReadSA("/home/my_user/myfile.csv");
```

Remarks

The standard input stream (stdin) can be read with **csvReadSA** by passing in `__STDIN` as the filename input. Note that `__STDIN` should not be passed in as a string. Correct usage is shown below:

```
x = csvReadSA(__STDIN);
```

See Also

[csvReadM](#), [xlsReadM](#), [xlsWrite](#), [xlsReadSA](#)

csvWriteM

Purpose

Write the contents of a **GAUSS** matrix to a CSV file.

Format

```
ret = csvWriteM(data, filename);
ret = csvWriteM(data, filename, sep);
ret = csvWriteM(data, filename, sep, prec);
ret = csvWriteM(data, filename, sep, prec, append);
ret = csvWriteM(data, filename, sep, prec, append, newline);
```

Input

<i>data</i>	Matrix, containing the data to be written
<i>filename</i>	string, valid filespec, name of CSV file to write.
<i>sep</i>	optional string, the character to separate the data. Default = ",".
<i>prec</i>	optional scalar, the number of digits of precision to retain. Default = 15.
<i>append</i>	optional scalar, 0 to overwrite entire file or 1 to append to file. Default = 0.
<i>newline</i>	optional string specifying the character(s) to end a line in the file. Default = "\n".

Output

ret	Scalar return code. 0 for success, or non-zero if an error occurred.
-----	--

Examples

Example 1: Basic Example

Write the contents from a matrix to a new file named `mydata.csv` located in your current working directory.

csvWriteM

```
//Create a simple matrix
x = { 1 2,
      3 4,
      5 6 };

//Write the contents of 'x' to a file named 'myfile.csv'
ret = csvWriteM(x, "myfile.csv");
```

Example 2: Create a tab separated text file

```
//Create a simple matrix
x = { 1 2,
      3 4,
      5 6 };

//Specify the optional separator input to be a tab character
sep = "\t";

//Write the data to the file 'mytabdata.csv'
ret = csvWriteM(x, "mytabdata.csv", sep);
```

Example 3: Specify the precision with which to write the data

```
//Create a simple matrix
x = { 1.102 2.001,
      3.041 4.232,
      5.113 6.523 };

//Specify the optional separator input to be a comma
sep = ",";

//Specify the number of significant digits to print
```



```
prec = 2;

//Write the data to the file 'mydata.csv'
ret = csvWriteM(x, "mydata.csv", sep, prec);
```

Example 4: Append to an existing file

```
//Create a simple matrix
x = { 9.008  1.005,
      1.445  4.247,
      2.913  1.020 };

//1 for append
append_flag = 1;

//Append the data to the file 'mydata.csv'
ret = csvWriteM(x, "mydata.csv", ",", 2, append_flag);
```

Example 5: Specify Windows style CRLF line endings

```
//Create a simple matrix
x = { 9.008  1.005,
      1.445  4.247,
      2.913  1.020 };

//'\c\r' indicates carriage return followed by a line feed
line_feed= "\c\r";

//Append the data to the file 'mydata.csv'
ret = csvWriteM(x, "mydata.csv", ",", 2, 0, line_feed);
```

Example 6: Specify full path to file

Windows: Notice that double backslashes are needed inside of a string.

cumprodc

```
ret = csvWriteM(x, "C:\\mydata\\myfile.csv");
```

Mac

```
ret = csvWriteM(x, "/Users/MyUserName/myfile.csv");
```

Linux

```
ret = csvWriteM(x, "/home/my_user/myfile.csv");
```

Remarks

The standard output and standard error streams (stdin, stderr) can be written to with **csvWriteM** by passing in the variable `__STDOUT`, or `__STDERR` as the filename input. Note that `__STDOUT`, or `__STDERR` should not be passed in as a string. The following example shows correct usage:

```
x = csvWriteM(__STDOUT);
```

See Also

[csvReadSA](#), [xlsWrite](#), [xlsWriteM](#), [xlsWriteSA](#), [xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

cumprodc

Purpose

Computes the cumulative products of the columns of a matrix.

Format

```
y = cumprodc(x);
```

Input

x	$N \times K$ matrix.
-----	----------------------

Output

y	$N \times K$ matrix containing the cumulative products of the columns of x .
-----	--

Remarks

This is based on the recursive series **recsercp**. **recsercp** could be called directly as follows:

```
recsercp(x, zeros(1,cols(x)));
```

to accomplish the same thing.

Example

```
x = { 1 -3,  
      2  2,  
      3 -1 };  
y = cumprodc(x);
```

Now if you view y , you will see:

```
1.000 -3.000  
y = 2.000 -6.000  
6.000  6.000
```

Source

cumprodc.src

cumsumc

See Also

[cumsumc](#), [recsercp](#), [recserar](#)

cumsumc

Purpose

Computes the cumulative sums of the columns of a matrix.

Format

$y = \text{cumsumc}(x);$

Input

x NxK matrix.

Output

y NxK matrix containing the cumulative sums of the columns of x .

Remarks

This is based on the recursive series function **recserar**. **recserar** could be called directly as follows:

`recserar(x, x[1, .], ones(1, cols(x)))`

to accomplish the same thing.

Example

```
x = { 1 -3,
      2  2,
      3 -1 };

y = cumsumc(x);
```

Now if you view *y*, you will see:

```
      1.000 -3.000
y =  3.000 -1.000
     6.000 -2.000
```

Source

cumsumc.src

See Also

[cumprodc](#), [recsercp](#), [recserar](#)

curve

Purpose

Computes a one-dimensional smoothing curve.

Format

```
{ u, v } = curve(x, y, d, s, sigma, G);
```

curve

Input

x	Kx1 vector, x-abscissae (X-axis values).
y	Kx1 vector, y-ordinates (Y-axis values).
d	Kx1 vector or scalar, observation weights.
s	scalar, smoothing parameter. If $s = 0$, curve performs an interpolation. If d contains standard deviation estimates, a reasonable value for s is K.
σ	scalar, tension factor.
G	scalar, grid size factor.

Output

u	(K*G)x1 vector, x-abscissae, regularly spaced.
v	(K*G)x1 vector, y-ordinates, regularly spaced.

Remarks

σ contains the tension factor. This value indicates the curviness desired. If σ is nearly zero (e.g. .001), the resulting curve is approximately the tensor product of cubic curves. If σ is large, (e.g. 50.0) the resulting curve is approximately bi-linear. If σ equals zero, tensor products of cubic curves result. A standard value for σ is approximately 1.

G is the grid size factor. It determines the fineness of the output grid. For $G = 1$, the input and output vectors will be the same size. For $G = 2$, the output grid is twice as fine as the input grid, i.e., u and v will have twice as many rows as x and y .

Source

spline.src

cvtos

Purpose

Converts a character vector to a string.

Format

```
s = cvtos(v);
```

Input

v

Nx1 character vector, to be converted to a string.

Output

s

string, contains the contents of *v*.

Remarks

cvtos in effect appends the elements of *v* together into a single string.

cvtos was written to operate in conjunction with **stocv**. If you pass it a character vector that does not conform to the output of **stocv**, you may get unexpected results. For example, **cvtos** does NOT look for 0 terminating bytes in the elements of *v*; it assumes every element except the last is 8 characters long. If this is not true, there will be 0's in the middle of *s*.

If the last element of *v* does not have a terminating 0 byte, **cvtos** supplies one for *s*.

Example

```
let v = { "Now is t" "he time " "for all " "good men" };
```

cvtosa

```
s = cvtos(v);
```

Now the variable *s* is a string with the following contents.

```
s = "Now is the time for all good men"
```

See Also

[stocv](#), [vget](#), [vlist](#), [vput](#), [vread](#)

cvtosa

Purpose

Converts an NxK character vector to an NxK string array.

Format

```
sa = cvtosa(cv);
```

Input

<i>cv</i>	NxK character vector, to be converted to a string array.
-----------	--

Output

<i>sa</i>	NxK string array, contains the contents of <i>cv</i> .
-----------	--

Example

```
cv = { MEAN MEDIAN MODE, MAX MIN QUARTILE };  
sa = cvtosa(cv);  
print sa;
```

C

Now the variable `sa` is a string array with the same contents as `cv` as we can see from the output below:.

MEAN	MEDIAN	MODE
MAX	MIN	QUARTILE

See Also

[stocv](#), [vget](#), [vlist](#), [vput](#), [vread](#)

datacreate

d

datacreate

Purpose

Creates a real data set.

Format

```
fh = datacreate(filename, vnames, col, dtyp, vtyp);
```

Input

<i>filename</i>	string, name of data file.
<i>vnames</i>	string or Nx1 string array, names of variables.
<i>col</i>	scalar, number of variables.
<i>dtyp</i>	scalar, data precision, one of the following: 2 2-byte, signed integer. 4 4-byte, single precision. 8 8-byte, double precision.
<i>vtyp</i>	scalar or Nx1 vector, types of variables, may contain one or both of the following: 0 character variable. 1 numeric variable.

Output

<i>fh</i>	scalar, file handle.
-----------	----------------------

Remarks

The file handle returned by **datacreate** is a scalar containing a positive integer value that uniquely identifies each file. This value is assigned by **GAUSS** when the **create**, **datacreate**, **datacreatecomplex**, **open** or **dataopen** commands are executed. The file handle is used to reference the file in the commands **readr** and **writer**. If **datacreate** fails, it returns a -1.

If *filename* does not include a path, then the file is placed on the current directory. The file is given a .dat extension if no extension is specified.

If *col* is set to 0, then the number of columns in the data set is controlled by the contents of *vnames*. If *col* is positive, then the file will contain *col* columns.

If *vnames* contains *col* elements, then each column is given the name contained in the corresponding row of *vnames*. If *col* is positive and *vnames* is a string, then the columns are given the names *vnames1*, *vnames2*, ..., *vnamesN* (or *vnames01*, *vnames02*, ..., *vnamesN*), where $N = col$. The numbers appended to *vnames* are padded on the left with zeros to the same length as *N*.

The *dtyp* argument allows you to specify the precision to use when storing your data. Keep in mind the following range restrictions when selecting a value for *dtyp*:

Data Type	Digits	Range
integer	4	$-32768 < X < 32767$
single	6-7	$8.43 \times 10^{-37} < X \leq 3.37 \times 10^{+38}$
double	15-16	$4.19 \times 10^{-307} < X < 1.67 \times 10^{+308}$

If the integer type is specified, numbers are rounded to the nearest integer as they are written to the data set. If the data to be written to the file contains character data, the precision must be 8 or the character information will be lost.

datacreatecomplex

If `vtyp` is a scalar, then the value in `vtyp` controls the types of all of the columns in the data set. If it is an $N \times 1$ vector, then the type of each column is controlled by the value in the corresponding row of `vtyp`.

Example

```
fh = datacreate("myfile.dat", "v", 100, 8, 1);  
x = rndn(500, 100);  
r = writer(fh, x);  
ret = close(fh);
```

This example creates a double precision data file called `myfile.dat`, which is placed in the current directory. The file contains 100 columns with 500 observations (rows), and the columns are given the names 'V001', 'V002', ..., 'V100'.

Source

`datafile.src`

See Also

[datacreatecomplex](#), [create](#), [dataopen](#), [writer](#)

datacreatecomplex

Purpose

Creates a complex data set.

Format

```
fh = datacreatecomplex(filename, vnames, col, dtyp,  
vtyp);
```

Input

<i>filename</i>	string, name of data file.
<i>vnames</i>	string or Nx1 string array, names of variables.
<i>col</i>	scalar, number of variables.
<i>dtyp</i>	scalar, data precision, one of the following: 2 2-byte, signed integer. 4 4-byte, single precision. 8 8-byte, double precision.
<i>vtyp</i>	scalar or Nx1 vector, types of variables, may contain one or both of the following: 0 character variable. 1 numeric variable.

Output

<i>fh</i>	scalar, file handle.
-----------	----------------------

Remarks

The file handle returned by **datacreatecomplex** is a scalar containing a positive integer value that uniquely identifies each file. This value is assigned by **GAUSS** when the **create**, **datacreate**, **datacreatecomplex**, **open** or **dataopen** commands are executed. The file handle is used to reference the file in the commands **readr** and **writer**. If **datacreatecomplex** fails, it returns a -1.

Complex data is stored a row at a time, with the real and imaginary halves interleaved, element by element. For columns containing character data, the imaginary parts are zeroed out.

datacreatecomplex

If *filename* does not include a path, then the file is placed on the current directory. The file is given a *.dat* extension if no extension is specified.

If *col* is set to 0, then the number of columns in the data set is controlled by the contents of *vnames*. If *col* is positive, then the file will contain *col* columns.

If *vnames* contains *col* elements, then each column is given the name contained in the corresponding row of *vnames*. If *col* is positive and *vnames* is a string, then the columns are given the names *vnames1*, *vnames2*, ..., *vnamesN* (or *vnames01*, *vnames02*, ..., *vnamesN*), where $N = col$. The numbers appended to *vnames* are padded on the left with zeros to the same length as *N*.

The *dtyp* argument allows you to specify the precision to use when storing your data. Keep in mind the following range restrictions when selecting a value for *dtyp*:

Data Type	Digits	Range
integer	4	$-32768 < X < 32767$
single	6-7	$8.43 \times 10^{-37} < X \leq 3.37 \times 10^{+38}$
double	15-16	$4.19 \times 10^{-307} < X < 1.67 \times 10^{+308}$

If the integer type is specified, numbers are rounded to the nearest integer as they are written to the data set. If the data to be written to the file contains character data, the precision must be 8 or the character information will be lost.

If *vtyp* is a scalar, then the value in *vtyp* controls the types of all of the columns in the data set. If it is an *N* x 1 vector, then the type of each column is controlled by the value in the corresponding row of *vtyp*.

Example

```
string vnames = { "random1", "random2" };
fh = datacreatecomplex("myfilecplx.dat",vnames,2,8,1);
x = complex(rndn(1000,2),rndn(1000,2));
r = writer(fh,x);
```

```
ret = close(fh);
```

This example creates a complex double precision data file called `myfilecplx.dat`, which is placed in the current directory. The file contains 2 columns with 1000 observations (rows), and the columns are given the names 'random1' and 'random2'.

Source

`datafile.src`

See Also

[datacreate](#), [create](#), [dataopen](#), [writer](#)

datalist

Purpose

List selected variables from a data set.

Format

```
datalist dataset [[var 1 [[var 2 ...]]]];
```

Input

<i>dataset</i>	literal, name of the data set.
<i>var#</i>	literal, the names of the variables to list.

Global Input

<u> </u> <i>range</i>	scalar, the range of rows to list. The default is all rows.
<u> </u> <i>miss</i>	scalar, controls handling of missing values.

dataload

	0	display rows with missing values.
	1	do not display rows with missing values.
The default is 0.		
<code>__prec</code>	scalar, the number of digits to the right of the decimal point to display. The default is 3.	

Remarks

The variables are listed in an interactive mode. As many rows and columns as will fit on the screen are displayed. You can use the cursor keys to pan and scroll around in the listing.

Example

```
datalist freq age sex pay;
```

This command will display the variables *age*, *sex*, and *pay* from the data set *freq.dat*.

Source

`datalist.src`

dataload

Purpose

Loads matrices, N-dimensional arrays, strings and string arrays from a disk file.

Format

```
y = dataload(filename);
```


Input

<code>filename</code>	string, name of data file.
-----------------------	----------------------------

Output

<code>y</code>	matrix, array, string or string array, data retrieved from the file.
----------------	--

Remarks

The proper extension must be included in the file name. Valid extensions are as follows:

<code>.fmt</code>	matrix file
	array file
<code>.fst</code>	string file
	string array file

See **FOREIGN LANGUAGE INTERFACE, CHAPTER 1**, for details on these file types.

Example

```
y = dataload("myfile.fmt");
```

See Also

[load](#), [datasave](#)

dataloop (dataloop)

dataloop (dataloop)

Purpose

Specifies the beginning of a data loop.

Format

```
dataloop infile outfile;
```

Input

infile string variable or literal, the name of the source data set.

Output

outfile string variable or literal, the name of the output data set.

Remarks

The statements between the **dataloop... endata** commands are assumed to be metacode to be translated at compile time. The data from *infile* is manipulated by the specified statements, and stored to the data set *outfile*. Case is not significant within the **dataloop... endata** section, except for within quoted strings. Comments can be used as in any **GAUSS** code.

Example

```
src = "source";  
dataloop ^src dest;  
make newvar = x1 + x2 + log(x3);  
x6 = sqrt(x4);  
keep x6, x5, newvar;
```

```
endata;
```

Here, *src* is a string variable requiring the caret (^) operator, while *dest* is a string literal.

dataopen

Purpose

Opens a data set.

Format

```
fh = dataopen(filename, mode);
```

Input

<i>filename</i>	string, name of data file.
<i>mode</i>	string containing one of the following:
"read"	Open file for read only.
"append"	Open file for append. The file pointer will start at the end of the file to add new rows.
"update"	Open file for update. Allows reading and writing. The file pointer will start at the first row.

Output

<i>fh</i>	scalar, file handle.
-----------	----------------------

Examples

Example 1: Read from a GAUSS dataset

This example opens the data file `credit.dat` from the **GAUSS** examples directory and reads 100 observations (rows) from the file into the global variable `y`.

```
//Create a file name with full path
file_name = getGAUSSHome() $+ "examples/credit.dat";

//Open file handle to dataset and assign it to 'fh'
fh = dataopen(file_name, "read");

//Read 100 rows from the dataset into the variable 'y'
y = readr(fh, 100);

//Close file handle
ret = close(fh);
```

Example 2: Write to a GAUSS dataset

```
//Create variable names for dataset
var_names = "alpha" $| "beta";

//Create dataset containing 2 variables with 5 observations
all equal to 1
x = ones(5, 2);
call saved(x, "my_ones.dat", var_names);

//Open file handle to dataset and assign it to 'fh'
fh = dataopen("my_ones.dat", "update");

//Write to the first row
y = { 17 21 };
call writer(fh, y);
```

```
//Close file handle
ret = close(fh);

//Load all contents of dataset
new_x = load("my_ones.dat");
```

After the code above, *new_x* should be equal to:

```
17 21
 1  1
 1  1
 1  1
 1  1
```

Remarks

1. The **file must exist before it can be opened** with the **dataopen** command (to create a new file, see **datacreate** or **datasave**).
2. The file handle returned by **dataopen** is a scalar containing a positive integer value that uniquely identifies each file. This value is assigned by **GAUSS** when the **create**, **datacreate**, **datacreatecomplex**, **open** or **dataopen** commands are executed. The file handle is used to reference the file in the commands **readr** and **writer**. If **dataopen** fails, it returns a -1.
3. A file can be opened simultaneously under more than one handle. If the value that is in the file handle when the **dataopen** command begins to execute matches that of an already open file, the process will be aborted and a File already open error message will be given. This gives you some protection against opening a second file with the same handle as a currently open file. If this happens, you would no longer be able to access the first file.
4. It is important to set unused file handles to zero because both **dataopen** and **datacreate** check the value that is in a file handle to see if it matches that of an

dataopen

open file before they proceed with the process of opening a file. You may set unused file handles to zero with the **close** or **closeall** commands.

5. If *filename* does not have an extension, **dataopen** appends a `.dat` extension before searching for the file. If the file is an `.fmt` matrix file, the extension must be explicitly given. If no path information is included, then **dataopen** searches for the file in the current directory.

6. Files opened in *read* mode cannot be written to. The pointer is set to the beginning of the file and the **writer** function is disabled for files opened in this way. This is the only mode available for matrix files (`.fmt`), which are always written in one piece with the **save** command.

7. Files opened in *append* mode cannot be read. The pointer is set to the end of the file so that a subsequent write to the file with the **writer** function will add data to the end of the file without overwriting any of the existing data in the file. The **readr** function is disabled for files opened in this way. This mode is used to add additional rows to the end of a file.

8. Files opened in *update* mode can be read from and written to. The pointer is set to the beginning of the file. This mode is used to make changes in a file.

9. The supported data set types are `"*.dat"`, `"*.h5"`, `"*.fmt"`.

For HDF5 file, the *dataset* must include schema and both file name and data set name must be provided, e.g. **glm**("h5://C:/gauss17/examples/testdata.h5/mydata").

Source

`datafile.src`

See Also

[open](#), [datacreate](#), [writer](#), [readr](#)

datasave

Purpose

Saves matrices, N-dimensional arrays, strings and string arrays to a disk file.

Format

```
ret = datasave(filename, x);
```

Input

<i>filename</i>	string, name of data file.
<i>x</i>	matrix, array, string or string array, data to write to disk.

Output

<i>ret</i>	scalar, return code, 0 if successful, or -1 if it is unable to write the file.
------------	--

Remarks

datasave can be used to save matrices, N-dimensional arrays, strings and string arrays. The following extensions are given to files that are saved with **datasave**:

matrix	.fmt
array	.fmt
string	.fst
string array	.fst

See **FOREIGN LANGUAGE INTERFACE**, CHAPTER 1, for details on these file types.

date

Use **dataload** to load a data file created with **datasave**.

Example

```
x = rndn(1000,100);  
ret = datasave("myfile.fmt",x);
```

See Also

[save](#), [dataload](#)

date

Purpose

Returns the current date in a 4-element column vector, in the order: year, month, day, and hundredths of a second since midnight.

Format

```
y = date;
```

Remarks

The hundredths of a second since midnight can be accessed using **hsec**.

Example

```
print date;
```

```
2015.0  
7.0
```



```
16.0
4571524.7
```

See Also

[time](#), [timestr](#), [ethsec](#), [hsec](#), [etstr](#)

datestr

Purpose

Returns a date in a string.

Format

```
str = datestr(d);
```

Input

<i>d</i>	4x1 vector, like the date function returns. If this is 0, the date function will be called for the current system date.
----------	---

Output

<i>str</i>	8 character string containing current date in the form: <i>mo/dy/yr</i>
------------	--

Example

```
d = { 2015, 10, 09, 0 };
y = datestr(d);
print y;
```

datestring

```


```

produces the following output:

```

    10/09/15

```

Source

time.src

See Also

[date](#), [datestring](#), [datestrymd](#), [time](#), [timestr](#), [ethsec](#)

datestring

Purpose

Returns a date in a string with a 4-digit year.

Format

```
str = datestring(d);
```

Input

<i>d</i>	4x1 vector, like the date function returns. If this is 0, the date function will be called for the current system date.
----------	---

Output

<i>str</i>	10 character string containing current date in the form: <i>mm/dd/yyyy</i>
------------	---

Example

```
dt = { 2015, 12, 18, 0 };  
y = datestring(dt);  
print y;
```

produces the following output:

```
12/18/2015
```

Source

time.src

See Also

[date](#), [datestr](#), [datestrymd](#), [time](#), [timestr](#), [ethsec](#)

datestrymd

Purpose

Returns a date in a string in the form *yyyymmdd*.

Format

```
str = datestrymd(d);
```

Input

<i>d</i>	4x1 vector, like the date function returns. If this is 0, the date function will be called for the current system date.
----------	---

dayinyr

Output

<code>str</code>	8 character string containing current date in the form: <i>yyyymmdd</i>
------------------	--

Example

```
d = { 2015, 10, 16, 0 };
y = datestrymd(d);
print y;
```

returns:

```
20151016
```

Source

`time.src`

See Also

[date](#), [datestr](#), [datestring](#), [time](#), [timestr](#), [ethsec](#)

dayinyr

Purpose

Returns day number in the year of a given date.

Format

```
daynum = dayinyr(dt);
```

Input

<i>dt</i>	3x1 or 4x1 vector, date to check. The date should be in the form returned by date .
-----------	--

Output

<i>daynum</i>	scalar, the day number of that date in that year.
---------------	---

Example

```
x = { 1973, 8, 31, 0 };  
y = dayinyr(x);  
print y;
```

produces:

```
y = 243.00000
```

Source

time.src

Globals

_isleap

dayofweek

Purpose

Returns day of week.

dayofweek

Format

```
d = dayofweek(a);
```

Input

a	Nx1 vector, dates in DT format.
---	---------------------------------

Output

d	Nx1 vector, integers indicating day of week of each date:
1	Sunday
2	Monday
3	Tuesday
4	Wednesday
5	Thursday
6	Friday
7	Saturday

Remarks

The DT scalar format is a double precision representation of the date and time. In the DT scalar format, the number

```
a = 20150415183207;
```

represents 18:32:07 or 6:32:07 PM on April 4, 2015.

```
d = dayofweek(a);
```

After running above code, d is 4 which means Wednesday.

Source

`time.src`

See Also

[dtday](#), [dttime](#), [dtdate](#), [dttostr](#)

dbAddDatabase

Purpose

Adds a database to the list of database connections using the driver type or a connection URL.

Format

```
db_id = dbAddDatabase(driver_type);  
db_id = dbAddDatabase(connection_url);
```

Input

<i>driver_type</i>	string, supported options include:
	DB2,
	IBASE,
	MYSQL,
	OCI,
	ODBC,
	PSQL,
	SQLITE,
	SQLITE2,

dbAddDatabase

<i>connection_</i> <i>url</i>	TDS. string, with the following format: driver://username:password@hostname:port/database_ name
----------------------------------	--

Output

<i>db_id</i>	scalar, index into a table of all opened database connections, or 0 on failure.
--------------	---

Example

```
db_id = dbAddDatabase ("MYSQL");  
  
url = "mysql://webuser:pswd@localhost:3306/dev";  
db_id = dbAddDatabase (url);
```

Remarks

Before using the connection, it must be initialized. e.g., call some or all of **dbSetDatabaseName()**, **dbSetUserName()**, **dbSetPassword()**, **dbSetHostName()**, **dbSetPort()**, and **dbSetConnectOptions()**, and, finally, **dbOpen()**.

The exception to this is using a connection URL, since this performs the above mentioned steps. Omitting portions of the connection URL is allowed, but the syntax must remain the same. For example:

```
id = dbAddDatabase ("oci://root:@localhost:/testing");
```

is a valid connection URL, but will not set the password or port number fields.

dbClose

Purpose

Closes a database connection and destroys any remaining queries.

Format

```
dbClose(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Remarks

dbClose() does not remove the database connection from the list of available database connections. The connection can be opened again without repeating the database initialization and setup steps.

dbCommit

Purpose

Commits a transaction to the database if the driver supports transactions and a **dbTransaction()** has been started.

Format

```
ret = dbCommit(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

dbCreateQuery

Output

<code>ret</code>	scalar, 1 for success or 0 for failure.
------------------	---

Example

```
db_id = dbAddDatabase("SQLITE");  
dbExecQuery(db_id, "INSERT INTO PEOPLE  
    (first, last) VALUES ('John', 'Doe');");  
dbCommit(db_id);  
dbClose(db_id);
```

Remarks

Note: For some databases, the commit will fail and return 0 if there is an active query using the database for a SELECT statement. Make the query inactive before doing the commit to resolve this problem.

Call `dbGetLastError()` to get information about errors.

dbCreateQuery

Purpose

Process an SQL statement and prepare a query. If *placeholders* is present, these values are bound sequentially to ODBC style parameters.

Format

```
qid = dbCreateQuery(db_id);  
qid = dbCreateQuery(db_id, query);  
qid = dbCreateQuery(db_id, query, placeholders);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>query</i>	string, database query to construct.
<i>placeholders</i>	string, or string array containing bind value(s).

Output

<i>qid</i>	scalar, query id to be used for result retrieval.
------------	---

Example

Example 1

```
qid = dbCreateQuery("SELECT * FROM GDP
                    WHERE COUNTRY = ?", "USA");
dbQueryExecPrepared(qid);

// Results as a matrix
results = dbQueryFetchAllM(qid);
```

Example 2

```
qid = dbCreateQuery("INSERT INTO
                    PEOPLE(id, fname, lname) VALUES
                    (NULL, ?, ?);");
dbQueryBindValue(qid, "Joe");
dbQueryBindValue(qid, "Smith");
dbQueryExecPrepared(qid);
```

Remarks

If the *placeholders* parameter is passed in, the values are bound sequentially to ODBC style parameters.

See also

[dbQueryPrepare](#)

dbExecQuery

dbExecQuery

Purpose

Executes an SQL statement and creates a query.

Format

```
qid = dbExecQuery(db_id, sql_statement);  
qid = dbExecQuery(db_id, sql_statement, placeholders);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>sql_statement</i>	string containing a valid SQL statement.
<i>placeholders</i>	string (array) containing bind value(s).

Output

<i>qid</i>	scalar, query id to be used for result retrieval.
------------	---

Example

In the examples below, *db_id* is a previously created database id.

Example 1

```
qid = dbExecQuery(db_id, "SELECT * FROM GDP  
WHERE COUNTRY = ?", "USA");  
  
// Results as a matrix  
results = dbQueryFetchAllM(qid);
```

Example 2

```
bd_vals = "Joe"|"Smith";
qid = dbExecQuery(db_id, "INSERT INTO PEOPLE(id,
      fname, lname); VALUES (NULL, ?, ?);", bd_vals);
```

Example 3

```
qid = dbExecQuery("SELECT * FROM PEOPLE
      p WHERE p.FNAME = ?", "Joe");

// Results as a string array
results = dbQueryFetchAllSA(qid);
```

dbGetConnectOptions

Purpose

Returns the connection options string used for a database connection.

Format

```
options = dbGetConnectOptions(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>options</i>	string, containing the connection options for the specified database connection.
----------------	--

See Also

[dbSetConnectOptions](#)

dbGetDatabaseName

Remarks

If you have not set any connection options with **dbSetConnectOptions**, then this function will return an empty string. For a full list of options see **dbSetConnectOptions**.

dbGetDatabaseName

Purpose

Returns the name of the database.

Format

```
db_name = dbGetDatabaseName(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>db_name</i>	string, name of the database.
----------------	-------------------------------

dbGetDriverName

Purpose

Returns the name of the connection's database driver.

Format

```
driver_name = dbGetDriverName(db_id);
```

dbGetDrivers();

Input

db_id scalar, database connection index number.

Output

driver_name string, name of the database driver.

Example

```
db_id = dbAddDatabase("SQLITE");  
print "Driver = " dbGetDriverName(db_id);
```

will print the following output

```
Driver = SQLITE
```

dbGetDrivers();

Purpose

Returns a list of available database drivers.

Format

```
drivers = dbGetDrivers();
```

Output

drivers Nx1 string array, list of available database drivers.

dbGetHostName

Example

```
print  
dbGetDrivers () ;
```

```
DB2  
MYSQL  
OCI  
ODBC  
PSQL  
SQLITE
```

dbGetHostName

Purpose

Returns the database connection's host name

Format

```
host_name = dbGetHostName(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>host_name</i>	string, name of database connection.
------------------	--------------------------------------

dbGetLastErrorNum

Purpose

Returns information about the last error that occurred on the database.

Format

```
last_error = dbGetLastErrorNum(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>last_error</i>	scalar, number of last error on the specified database.
-------------------	---

dbGetLastErrorText

Purpose

Returns information about the last error that occurred on the database.

Format

```
last_error = dbGetLastErrorText(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

dbGetNumericalPrecPolicy

Output

<i>last_error</i>	string, details of last error on the specified database.
-------------------	--

dbGetNumericalPrecPolicy

Purpose

Returns the default numerical precision policy for a specified database connection.

Format

```
prec_policy = dbGetNumericalPrecPolicy(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>prec_policy</i>	scalar:	
	DB_HIGH_PRECISION	strings will be used to preserve precision
	DB_LOW_PRECISION_INT32	Force 32-bit integer values
	DB_LOW_PRECISION_INT64	Force 64-bit integer values

DB_LOW_ PRECISION_ DOUBLE	Force double values. This is the default policy.
---------------------------------	--

dbGetPassword

Purpose

Returns a connection's password.

Format

db_password = **dbGetPassword**(*db_id*);

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>db_ password</i>	string containing the password for the specified database connection or a null string.
-------------------------	--

Remarks

dbGetPassword() will only return passwords set with **dbSetPassword()**.

dbGetPort

Purpose

Returns the database connection's port number if it has been set.

dbGetPrimaryIndex

Format

```
db_port = dbGetPort(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>db_port</i>	scalar, the port number of the specified database connection.
----------------	---

Remarks

dbGetPort() will only return the port number if it was previously set with **dbSetPort()**.

dbGetPrimaryIndex

Purpose

Returns the primary index for the specified table.

Format

```
primary_index = dbGetPrimaryIndex(db_id, table_name);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>table_name</i>	string, name of the table to reference.

Output

<i>primary_index</i>	2x1 string array; the [1,1] element is the cursor name and the [2,1] element is the index name
----------------------	--

dbGetTableHeaders

Purpose

Returns a string array populated with the names of all the fields in a specified table (or view).

Format

```
field_names = dbGetTableHeaders(db_id, table_name);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>table_name</i>	string, name of table or view.

Output

<i>field_names</i>	string array containing the column names for the specified table or view.
--------------------	---

Remarks

The order in which the fields appear in the record is undefined.

dbGetTables

dbGetTables

Purpose

Returns the database's tables, system tables and views.

Format

```
tables = dbGetTables(db_id, type);  
tables = dbGetTables(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.	
<i>type</i>	string:	
	"Tables"	All tables visible to the user. This is the default value.
	"System Tables"	Internal tables used by the database.
	"Views"	All views visible to the user.
	"All"	All of the above.

Output

<i>tables</i>	Nx1 string array containing the information specified by the 'type' parameter.
---------------	--

dbGetUserName

Purpose

Returns the database connection's user name.

Format

```
user_name = dbGetUserName(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>user_name</i>	string containing the user name associated with the specified database connection.
------------------	--

See Also

[dbSetUserName](#)

dbHasFeature

Purpose

Returns a 1 if the database supports the specified feature.

Format

```
ret = dbHasFeature(db_id, feature);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>feature</i>	string: DB_TRANSACTIONS DB_QUERY_SIZE

dbHasFeature

DB_BLOB
DB_UNICODE
DB_PREPARED_QUERIES
DB_NAMED_PLACEHOLDERS
DB_POSITIONAL_PLACEHOLDERS
DB_LAST_INSERT_ID
DB_BATCH_OPERATIONS
DB_SIMPLE_LOCKING
DB_LOW_PRECISION_NUMBERS
DB_EVENT_NOTIFICATIONS
DB_FINISH_QUERY
DB_MULTIPLE_RESULT_SETS

Output

<i>ret</i>	scalar, 1 if the database supports the specified feature, or 0 if not.
------------	--

Example

```
db_id = dbAddDatabase("MYSQL");  
  
// Create empty query  
qid = dbCreateQuery(db_id);  
  
if dbHasFeature(db_id, "NamedPlaceholders");  
    dbQueryPrepare(qid, "SELECT * FROM GDP  
    WHERE COUNTRY = :country");  
    dbQueryBindValue(qid, ":country", "USA");  
else;  
    dbQueryPrepare(qid, "SELECT * FROM GDP  
    WHERE COUNTRY = ?");
```



```
        dbQueryAddBindValue (qid, "USA");  
    endif;  
  
    dbQueryExecPrepared (qid);
```

Remarks

Note that some databases need to be opened with **dbOpen ()** before this can be determined.

dbIsDriverAvailable

Purpose

Returns 1 if a specified database driver is available.

Format

```
ret = dbIsDriverAvailable(name);
```

Input

<i>name</i>	string, name of driver to check
-------------	---------------------------------

Output

<i>ret</i>	scalar, 1 if the specified driver is available, or 0 if not.
------------	--

dbIsOpen

dbIsOpen

Purpose

Reports whether a specified database connection is open.

Format

```
ret = dbIsOpen(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>ret</i>	scalar, 1 if the connection is open or 0 if it is closed.
------------	---

dbIsOpenError

Purpose

Reports whether an error occurred while attempting to open the database connection.

Format

```
ret = dbIsOpenError(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>ret</i>	scalar, 1 if there was an error or 0 if not.
------------	--

dbIsValid

Purpose

Reports whether a specified database connection has a valid driver.

Format

```
ret = dbIsValid(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>ret</i>	scalar, 1 if the database connection has a valid driver or 0 if not.
------------	--

Example

```
// Use default connection
db_id = dbAddDatabase("SQLITE");
ret = dbIsValid(db_id);           // Returns 1 for 'true'

db_id = dbAddDatabase("BAD_DRIVER_NAME");
ret = dbIsValid(db_id);          // Returns 0 for 'false'
```

dbNumericalPrecPolicy

dbNumericalPrecPolicy

Purpose

Returns the default numerical precision policy for a specified database connection.

Format

```
prec_policy = dbNumericalPrecPolicy(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>prec_policy</i>	scalar:
0	strings will be used to preserve precision
1	Force 32-bit integer values
2	Force 64-bit integer values
4	Force double values. This is the default policy.

dbOpen

Purpose

Opens a specified database connection using the current connection values.

Format

```
ret = dbOpen(db_id, user_name, password);  
ret = dbOpen(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>user_name</i>	string, user name for the database being connected to.
<i>password</i>	string, password associated with the specified user name for this database.

Output

<i>ret</i>	scalar, 1 for success.
------------	------------------------

Example

Set driver and host

```
db_id = dbAddDatabase("MYSQL");  
dbSetHostName(db_id, "localhost");
```

then, either

```
dbSetUserName(db_id, "test");  
dbSetPassword(db_id, "secret_passw0rd");  
ret = dbOpen(db_id);
```

or

```
ret = dbOpen(db_id, "test",  
            "secret_passw0rd");
```

dbQueryBindValue

dbQueryBindValue

Purpose

Set the placeholder placeholder to be bound to value val in the prepared statement. Note that the placeholder mark (e.g :) must be included when specifying the placeholder name.

Format

```
dbQueryBindValue(qid, placeholder, val);
```

Input

<i>qid</i>	scalar, query number.
<i>placeholder</i>	string, Oracle style (:value_name) or index of ODBC style (?) placeholder.
<i>val</i>	valid type, the value to be bound.

Remarks

Values cannot be bound to multiple locations in the query.

Example

```
db_id = dbAddDatabase("MYSQL");
qid = dbCreateQuery(db_id);
dbQueryPrepare(qid, "SELECT * FROM
    PEOPLE WHERE FIRST = :fname AND
    LAST = :lname");
dbQueryBindValue(qid, ":fname", "John");
dbQueryBindValue(qid, ":lname", "Doe");
dbQueryExecPrepared(qid);
```

dbQueryClear

Purpose

Clears the result set and releases any resources held by the query. Sets the query state to inactive.

Format

```
dbQueryClear(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Remarks

You should rarely if ever need to call this function.

dbQueryCols

Purpose

Returns the number of fields in the record.

Format

```
num_fields = dbQueryCols(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

dbQueryExecPrepared

Output

<i>num_ fields</i>	scalar, number of fields.
------------------------	---------------------------

dbQueryExecPrepared

Purpose

Executes a previously created and prepared query.

Format

```
ret = dbQueryExecPrepared(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>ret</i>	scalar, 1 for success and 0 for failure.
------------	--

Example

```
qid = dbCreateQuery(db_id);  
dbQueryPrepare(qid, "SELECT * FROM  
    USERS WHERE ID = :id");  
dbQueryBindValue(qid, ":id", "5");  
dbQueryExecPrepared(qid);  
  
results = dbQueryFetchAllSA(qid);
```



```
dbQueryBindValue(qid, ":id", "10");

// Re-execute the query with new value
dbQueryExecPrepared(qid);

results = dbQueryFetchAllSA(qid);
```

dbQueryFetchAllM

Purpose

Returns the result set for the current query as a matrix.

Format

```
result = dbQueryFetchAllM(qid);
result = dbQueryFetchAllM(qid, columns);
```

Input

<i>qid</i>	scalar, query number.
<i>columns</i>	string or string array, specific columns to pull out from result matrix. Must be a subset of fields from SELECT statement.

Output

<i>result</i>	matrix, the result set; or if the result set is empty, a scalar error code.
---------------	---

dbQueryFetchAllM

Remarks

For string results, or to treat numerical results as strings, use **dbQueryFetchAllSA()** to return a string array.

This function retrieves all rows at once. You can process rows in an iterative manner by using the **dbQueryFetchOneM()** and **dbQueryFetchOneSA()** functions.

Example

Example 1

```
qid = dbExecQuery(db_id, "SELECT * FROM GDP");

gdp = dbQueryFetchAllM(qid);

// If 'gdp' is a scalar error code
if scalmiss(gdp);
    print "No results";
else;
    // do something with gdp
endif;
```

Example 2

```
qid = dbExecQuery(db_id, "SELECT * FROM
    PEOPLE WHERE COUNTRY = ?", "USA");

// specify zipcode as column of interest
zipcodes = dbQueryFetchAllM(qid, "ZIPCODE");

if not scalmiss(zipcodes);
    print "zip codes = " zipcodes;
endif;
```

See Also

[dbQueryFetchAllSA](#), [dbQueryFetchOneM](#), [dbQueryFetchOneSA](#)

dbQueryFetchAllSA

Purpose

Returns the result set for the current query as a string array.

Format

```
result = dbQueryFetchAllSA(qid);  
result = dbQueryFetchAllSA(qid, columns);
```

Input

<i>qid</i>	scalar, query number.
<i>columns</i>	string or string array, specific columns to pull out from result matrix. Must be a subset of fields from SELECT statement.

Output

<i>result</i>	string array, containing the result set for the current query. If the result set is empty, a scalar error code is returned.
---------------	---

Remarks

For numerical only results, use **dbQueryFetchAllM()** to return a matrix.
This function retrieves all rows at once. You can process rows in an iterative manner by using the **dbQueryFetchNextM()** and **dbQueryFetchNextSA()** functions.

Example

```
qid = dbExecQuery(db_id, "SELECT * FROM
```

dbQueryFetchOneM

```
        PEOPLE WHERE COUNTRY = ?", "USA");

// specify names as columns of interest
names = dbQueryFetchAllSA(qid,
    "FIRST_NAME"|$|"LAST_NAME");

// If 'names' is not a scalar error code
if not scalmiss(names);
    print "People in the USA = " names;
endif;
```

See Also

[dbQueryFetchAllM](#), [dbQueryFetchNextSA](#), [dbQueryFetchNextM](#)

dbQueryFetchOneM

Purpose

Returns a single row as an Nx1 matrix where N is the column count of the SELECT statement containing the field information for the current query.

Format

```
record = dbQueryFetchOneM(qid);
record = dbQueryFetchOneM(qid, columns);
```

Input

<i>qid</i>	scalar, query number.
<i>columns</i>	string or string array, specific columns to pull from the result matrix. Must be a subset of fields from the SELECT statement.

Output

record matrix, if the query points to a valid row (**dbQueryIsValid()** returns true), the record is populated with the row's values. An empty record (**scalmiss** (*record*) is true) is returned when there is no active query (**dbQueryIsActive()** returns false).

Remarks

This function is only useful in an iterative context. You can easily retrieve all the results at once by using the **dbQueryFetchAllM()** and **dbQueryFetchAllSA()** functions.

For string results, or to treat numerical results as a string, using **dbQueryFetchOneSA()** will return a string array.

Example

```
qid = dbExecQuery(db_id, "SELECT YTD,  
    TOTAL FROM GDP");  
  
do while dbQuerySeekNext(qid);  
    record = dbQueryFetchOneM(qid);  
    ytd = record[1];  
    total = record[2];  
endo;
```

See Also

[dbQueryFetchOneSA](#), [dbQueryFetchAllM](#), [dbQueryFetchAllSA](#), [dbQueryGetField](#)

dbQueryFetchOneSA

dbQueryFetchOneSA

Purpose

Returns a single row as a string vector containing the field information for the current query.

Format

```
record = dbQueryFetchOneSA(qid);  
record = dbQueryFetchOneSA(qid, columns);
```

Input

<i>qid</i>	scalar, query number.
<i>columns</i>	string or string array, specific columns to pull from the result matrix. Must be a subset of fields from the SELECT statement.

Output

<i>record</i>	string array, if the query points to a valid row (dbQueryIsValid() returns true), the record is populated with the row's values. An empty record (scalmiss (<i>record</i>) is true) is returned when there is no active query dbQueryIsActive() returns false).
---------------	--

Remarks

This function is only useful in an iterative context. You can easily retrieve all the results at once by using the **dbQueryFetchAllM()** and **dbQueryFetchAllSA**

() functions.

For numerical only results, using **dbQueryFetchOneM()** will return a matrix instead of a string array.

Example

```
qid = dbExecQuery(db_id, "SELECT COUNTRY,
    TOTAL FROM GDP");
do while dbQuerySeekNext(qid);
    record = dbQueryFetchOneSA(qid);
    country = record[1];
    total = record[2];
endo;
```

See Also

[dbQueryFetchOneM](#), [dbQueryFetchAllM](#), [dbQueryFetchAllSA](#), [dbQueryGetField](#)

dbQueryFinish

Purpose

Instructs the database driver that no more data will be fetched from this query until it is re-executed.

Format

```
dbQueryFinish(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

dbQueryGetBoundValue

Remarks

There is normally no need to call this function, but it may be helpful in order to free resources such as locks or cursors if you intend to re-use the query at a later time.

Sets the query to inactive. Bound values retain their values.

dbQueryGetBoundValue

Purpose

Returns the value for a placeholder in a query.

Format

```
val = dbQueryGetBoundValue(qid, placeholder);
```

Input

<i>qid</i>	scalar, query number.
<i>placeholder</i>	string, Oracle style (:value_name) or index of ODBC style (?) placeholder.

Output

<i>val</i>	string, bound value if previously set.
------------	--

Example

```
db_id = dbAddDatabase ("MYSQL");  
qid = dbCreateQuery (db_id);  
dbQueryPrepare (qid, "SELECT * FROM
```



```
        PEOPLE WHERE FIRST = :fname AND
        LAST = :lname");
dbQueryBindValue(qid, ":fname", "John");
dbQueryBindValue(qid, ":lname", "Doe");

print "Name = ";
print dbQueryGetBoundValue(qid, ":fname");
print dbQueryGetBoundValue(qid, ":lname");
```

or

```
db_id = dbAddDatabase("MYSQL");
string args = { "John", "Doe" };
qid = dbCreateQuery(db_id, "SELECT * FROM
        PEOPLE WHERE FIRST = ? AND LAST = ?", args);

print "Name = ";
print dbQueryGetBoundValue(qid, 1);
print dbQueryGetBoundValue(qid, 2);
```

results in

```
Name = John Doe
```

dbQueryGetBoundValues

Purpose

Returns an Nx2 string array containing the placeholders and their corresponding values in a query.

Format

```
bound_values = dbQueryGetBoundValues(qid);
```

dbQueryGetField

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>bound_values</i>	Nx2 string array. The first column contains the placeholders and the second column contains the corresponding values.
---------------------	---

Example

```
db_id = dbAddDatabase("MYSQL");
qid = dbCreateQuery(db_id);
dbQueryPrepare(qid, "SELECT * FROM
    PEOPLE WHERE FIRST = :fname AND
    LAST = :lname");
dbQueryBindValue(qid, ":fname", "John");
dbQueryBindValue(qid, ":lname", "Doe");

print "Vars = " dbQueryGetBoundValues(qid);
```

will print

```
Vars =
    :fname    John
    :lname    Doe
```

dbQueryGetField

Purpose

Returns the value of a specified field in the current record. An overloaded version that accepts a column name as input is available, but not as efficient.

Format

```
field_value = dbQueryGetField(qid, idx);  
field_value = dbQueryGetField(qid, name);
```

Input

<i>qid</i>	scalar, query number.
<i>idx</i>	scalar, index of the field whose value should be returned.

Remarks

The fields are numbered from left to right using the text of the SELECT statement, e.g. in

```
qid = dbExecQuery("SELECT forename, surname FROM people");  
  
do while dbQuerySeekNext(qid);  
    forename = dbQueryGetField(qid, 1);  
    // Using field index  
    surname = dbQueryGetField(qid, 2);  
    // Using field index  
    forename = dbQueryGetField(qid, "forename");  
    // Using field name  
    surname = dbQueryGetField(qid, "surname");  
    // Using field name  
end;
```

Field 1 is *forename* and field 2 is *surname*. Using SELECT * is not recommended because the order of the fields in the query is undefined.

See Also

[dbQueryFetchOneM](#), [dbQueryFetchOneSA](#)

dbQueryGetLastErrorNum

dbQueryGetLastErrorNum

Purpose

Returns error information about the last error that occurred (if any) with the last executed query.

Format

```
err_num = dbQueryGetLastErrorNum();
```

Output

<i>err_num</i>	scalar, number of last error.
----------------	-------------------------------

Remarks

Because a failed query will not have a valid handle (id), this function retrieves stored error information about the last executed query.

See also

[dbQueryGetLastErrorText](#)

dbQueryGetLastErrorText

Purpose

Returns error information about the last error that occurred (if any) with the last executed query.

Format

```
err_txt = dbQueryGetLastErrorText();
```

Output

<i>err_txt</i>	2x1 string array, database and driver text of last error.
----------------	---

Remarks

Because a failed query will not have a valid handle (id), this function retrieves stored error information about the last executed query.

See also

[dbQueryGetLastErrorNum](#)

dbQueryGetLastInsertID

Purpose

Returns the object ID of the most recent inserted row if supported by the database.

Format

```
last_insert = dbQueryGetLastInsertID(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>last_insert</i>	scalar, object id
--------------------	-------------------

dbQueryGetLastQuery

Remarks

If more than one row was touched by the insert, the behavior is undefined.

For MySQL databases the row's auto-increment field will be returned.

With a PSQL database, the table must contain OID's which were not created by default. Check the *default_with_oids* configuration variable to be sure.

Example

```
// Given NAMES is an empty MySQL
// table with the *id* column
// auto-incrementing.
db_id = dbAddDatabase("MYSQL");
qid = dbCreateQuery(db_id, "INSERT
    INTO NAMES (first, last) VALUES
    ('John', 'Doe');");

if dbHasFeature(db_id, "LastInsertId");
    last_id = dbQueryGetLastInsertID(qid);
endif;
```

See Also

[dbHasFeature](#)

dbQueryGetLastQuery

Purpose

Returns the text of the current query being used.

Format

```
query_string = dbQueryGetLastQuery(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>query_string</i>	string, text of the current query, or empty string if there is no current query.
---------------------	--

dbQueryGetNumRowsAffected**Purpose**

Reports the number of rows affected by the result's SQL statement.

Format

```
num_rows = dbQueryGetNumRowsAffected(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>num_rows</i>	scalar, the number of rows affected by the result's SQL statement, or a -1 if it cannot be determined or the query is not active.
-----------------	---

dbQueryGetPosition

Remarks

Note that for SELECT statements, the value is undefined; use `dbQueryRows()` instead.

Example

```
qid = dbCreateQuery(db_id, "INSERT INTO
    PEOPLE (fname, lname) VALUES
    ('John', 'Doe');");
print dbQueryGetNumRowsAffected(qid) " row(s) were
affected";
```

results in

```
1 row(s) were affected
```

See Also

[dbQueryRows](#), [dbHasFeature](#)

dbQueryGetPosition

Purpose

Returns the current internal position of the query.

Format

```
index = dbQueryGetPosition(qid)
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>index</i>	scalar, query position
--------------	------------------------

Remarks

The first record is at position zero. If the position is invalid, the function returns QSql::BeforeFirstRow or QSql::AfterLastRow, which are special negative values.

Example

```
qid = dbCreateQuery(db_id, "SELECT *  
    FROM PEOPLE");  
do while dbQuerySeekNext(qid);  
    print "Current index = "  
        dbQueryGetPosition(qid);  
endo;
```

dbQueryIsActive

Purpose

Returns 1 if the query is active.

Format

```
ret = dbQueryIsActive(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

dbQueryIsForwardOnly

Output

<i>ret</i>	scalar, 1 if the query is active or 0 if not.
------------	---

Remarks

An active query is one that has been **dbQueryExecPrepared()**'d successfully, but not yet finished with. When you are finished with an active query, you can make the query inactive by calling **dbQueryFinish()** or **dbQueryClear()**.

Note: Of particular interest is an active query that is a SELECT statement. For some databases that support transactions, an active query that is a SELECT statement can cause a **dbCommit()** or a **dbRollback()** to fail, so before committing or rolling back, you should make your active SELECT statement query inactive using one of the methods listed above.

Example

```
qid = dbCreateQuery(db_id);

dbQueryIsActive(qid); // False dbQueryPrepare(qid, "INSERT
INTO TEST
    (foo, bar) VALUES (1, 2);");

dbQueryIsActive(qid); // False dbQueryExecPrepared(qid);

dbQueryIsActive(qid); // True dbQueryFinish(qid);

dbQueryIsActive(qid); // False
```

dbQueryIsForwardOnly

Purpose

Reports whether you can only scroll forward through a result set.

Format

```
ret = dbQueryIsForwardOnly(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>ret</i>	scalar, 1 if the result set can only be scrolled through forward, otherwise a 0.
------------	--

Remarks

Setting a query to "forward only" will usually improve performance. By default, queries are created with "forward only" off.

See Also

[dbQuerySetForwardOnly](#), [dbQuerySeekNext](#)

dbQueryIsNull

Purpose

Returns 1 if the query is active, positioned on a valid record and the field is NULL; otherwise returns 0. Reports whether the current field pointed at by an active query positioned on a valid record is NULL.

Format

```
ret = dbQueryIsNull(qid, field);
```

dbQueryIsSelect

Input

<i>qid</i>	scalar, query number.
<i>field</i>	scalar, index into result set.

Output

<i>ret</i>	scalar, 1 if the field is NULL or 0 otherwise.
------------	--

Remarks

Note that for some drivers, **dbQueryIsNull()** will not return accurate information until after an attempt is made to retrieve data.

See Also

[dbQueryIsActive](#), [dbQueryIsValid](#)

dbQueryIsSelect

Purpose

Reports whether the specified query is a SELECT statement.

Format

```
ret = dbQueryIsSelect(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>ret</i>	scalar, 1 if the query is a SELECT statement or 0 otherwise.
------------	--

Example

```
qid = dbExecQuery(db_id, "SELECT *  
    FROM PEOPLE");  
  
dbQueryIsSelect(qid); // True  
  
qid = dbExecQuery(db_id, "INSERT INTO  
    PEOPLE (fname, lname) VALUES  
    ('John', 'Doe');");  
  
dbQueryIsSelect(qid); // False
```

dbQueryIsValid

Purpose

Reports whether the specified query is positioned on a valid record.

Format

```
ret = dbQueryIsValid(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

dbQueryPrepare

Output

<i>ret</i>	scalar, 1 if the query is positioned on a valid record or 0 otherwise.
------------	--

Example

```
qid = dbExecQuery(db_id, "SELECT * FROM
    PEOPLE");

dbQueryIsValid(qid); // False
// Give it a valid position
dbQuerySeekFirst(qid);

// Iterate until no longer valid.
do while dbQueryIsValid(qid);
    // dbQueryIsValid = True
    dbQuerySeekNext(qid);
endo;

dbQueryIsValid(qid); // False
```

dbQueryPrepare

Purpose

Prepares a SQL query for execution.

Format

```
ret = dbQueryPrepare(qid, query);
```

Input

<i>qid</i>	scalar, query index number.
<i>query</i>	string, database query to prepare.

Output

<i>ret</i>	scalar, 1 for success and 0 for failure.
------------	--

Example

```
db_id = dbAddDatabase("MYSQL");
qid = dbCreateQuery(db_id);
ret = dbQueryPrepare(qid, "SELECT *
    FROM STOCKS WHERE SYMBOL = :sym");
dbQueryBindValue(qid, ":sym", "GOOG");
ret = dbQueryExecPrepared(qid);
```

Remarks

The query may contain placeholders for binding values. Both Oracle style colon-name (e.g., :surname), and ODBC style (?) placeholders are supported; but they cannot be mixed in the same query.

Portability note: Some databases choose to delay preparing a query until it is executed the first time. In this case, preparing a syntactically incorrect query succeeds, but every consecutive **dbQueryExecPrepared()** will fail.

For SQLite, the query string can contain only one statement at a time. If more than one statement is given, the function returns 0.

See also

[dbQueryBindValue](#)

dbQueryRows

dbQueryRows

Purpose

Returns the size of the result (number of rows returned), or -1 if the size cannot be determined or if the database does not support reporting information about query sizes.

Format

```
result_size = dbQueryRows(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>result_size</i>	scalar, number of rows in the current result set of the active query. If the number of rows cannot be determined a -1 is returned.
--------------------	--

Remarks

Note that if the query is not active or if the query is not a SELECT statement, a -1 is returned. These properties can be checked with **dbQueryIsActive()** or **dbQueryIsSelect()**.

Example

```
// Given a table with US States.  
qid = dbCreateQuery(db_id, "SELECT *  
    FROM STATES");
```



```
count = dbQueryRows(qid); // count = 50
```

dbQuerySeek

Purpose

Retrieves the record at a specified position, if available, and positions the query on the retrieved record.

Format

```
ret = dbQuerySeek(qid, idx, idx_type);  
ret = dbQuerySeek(qid, idx);
```

Input

<i>qid</i>	scalar, query number.
<i>idx</i>	scalar, the index at which to place the cursor.
<i>idx_type</i>	scalar, 1 for relative position or 0 for absolute positioning. If not specified, absolute positioning is used.

Output

<i>ret</i>	scalar, 1 if successful.
------------	--------------------------

Remarks

The first record is at position 1. Note that the query must be in an active state before calling this function. The state of the query may be verified with the function

dbQuerySeekFirst

dbQueryIsSelect()

If *idx_type* is 0 (the default), the following rules apply:

If *idx* is negative, the result is positioned before the first record and 0 is returned. Otherwise, an attempt is made to move to the record at position *idx*. If the record at position *idx* could not be retrieved, the result is positioned after the last record and 0 is returned. If the record is successfully retrieved, 1 is returned.

If *idx_type* is 1, the following rules apply:

If the result is currently positioned before the first record or on the first record, and *idx* is negative, there is no change, and 0 is returned.

If the result is currently located after the last record, and *idx* is positive, there is no change, and 0 is returned. If the result is currently located somewhere in the middle, and the relative offset *idx* moves the result below zero, the result is positioned before the first record and 0 is returned.

Otherwise, an attempt is made to move to the record *idx* records ahead of the current record (or *idx* records behind the current record if *idx* is negative).

If the record at offset *idx* could not be retrieved, the result is positioned after the last record if *idx* \geq 0, (or before the first record if *idx* is negative), and 0 is returned. If the record is successfully retrieved, 1 is returned.

See Also

[dbQuerySeekFirst](#), [dbQuerySeekLast](#), [dbQuerySeekNext](#), [dbQuerySeekPrevious](#)

dbQuerySeekFirst

Purpose

Retrieves the first record in the result, if available, and positions the query on the retrieved record.

Format

```
ret = dbQuerySeekFirst(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>ret</i>	1 if successful. If unsuccessful the query position is set to an invalid position and 0 is returned.
------------	--

Remarks

Note that the result must be in the active state or it will do nothing and return. This can be verified by calling the **dbQueryIsSelect()** function.

Example

```
qid = dbCreateQuery(db_id, "SELECT *  
    FROM PEOPLE");  
  
do while dbQuerySeekNext(qid);  
    // iterate over results  
endo;  
  
// set back to start  
dbQuerySeekFirst(qid);  
  
do while dbQuerySeekNext(qid);  
    // iterate over results AGAIN  
endo;
```

dbQuerySeekLast

dbQuerySeekLast

Purpose

Retrieves the last record in the result, if available, and positions the query on the retrieved record.

Format

```
ret = dbQuerySeekLast(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>ret</i>	scalar, returns 1 if successful. If unsuccessful the query position is set to an invalid position and 0 is returned.
------------	--

Remarks

Note that the result must be in the active state and `dbQueryIsSelect()` must return 1 before calling this function or it will do nothing and return 0.

Example

```
// Given STATES is a table with all
// 50 states listed alphabetically
qid = dbExecQuery(db_id, "SELECT name
    FROM STATES");

// Move to last state
```

```
ret = dbQuerySeekLast(qid);

//If 'ret' is equal to 0
if not ret;
    print "dbQuerySeekLast failed";
else;
    // Print last state: Wyoming
print dbQueryFetchOneSA(qid);
endif;
```

See also

[dbQuerySeekNext](#), [dbQuerySeekPrevious](#), [dbQuerySeekFirst](#), [dbQuerySeek](#), [dbQueryGetPosition](#)

dbQuerySeekNext

Purpose

Retrieves the next record in the result, if available, and positions the query on the retrieved record.

Format

```
ret = dbQuerySeekNext(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>ret</i>	scalar, if the record could not be retrieved, the result is positioned after the last record and 0 is returned. If the
------------	--

dbQuerySeekNext

record is successfully retrieved, 1 is returned.

Remarks

Note that the result must be in the active state before calling this function or it will do nothing and return 0. You can verify the status of the query with **dbQueryIsSelect()**.

The following rules apply:

If the result is currently located before the first record, e.g., immediately after a query is executed, an attempt is made to retrieve the first record.

If the result is currently located after the last record, there is no change and 0 is returned.

If the result is located somewhere in the middle, an attempt is made to retrieve the next record.

Example

```
qid = dbCreateQuery(db_id, "SELECT *  
    FROM PEOPLE");  
  
do while dbQuerySeekNext(qid);  
    row = dbQueryFetchOneSA(qid);  
    // Or dbQueryFetchOneM(qid) if data  
    // is numeric  
endo;
```

See Also

[dbQuerySeekFirst](#), [dbQuerySeekLast](#), [dbQuerySeekPrevious](#), [dbQuerySeek](#),
[dbQueryGetPosition](#)

dbQuerySeekPrevious

Purpose

Retrieves the previous record in the result, if available, and positions the query on the retrieved record.

Format

```
ret = dbQuerySeekPrevious(qid);
```

Input

<i>qid</i>	scalar, query number.
------------	-----------------------

Output

<i>ret</i>	scalar, 1 if the record is successfully retrieved. If the record could not be retrieved, the result is positioned before the first record and 0 is returned.
------------	--

Remarks

Note that the result must be in the active state before calling this function or it will do nothing and return false. The state of the query can be verified with **dbQueryIsSelect()**.

The following rules apply:

If the result is currently located before the first record, there is no change and 0 is returned.

If the result is currently located after the last record, an attempt is made to retrieve the last record.

dbQuerySetForwardOnly

If the result is somewhere in the middle, an attempt is made to retrieve the previous record.

See Also

[dbQuerySeekFirst](#), [dbQuerySeekLast](#), [dbQuerySeekNext](#), [dbQuerySeek](#), [dbQueryGetPositino](#)

dbQuerySetForwardOnly

Purpose

Sets forward only mode to forward. If forward is true, only **dbQuerySeekNext()** and **dbQuerySeek()** with positive values, are allowed for navigating the results.

Format

```
dbQuerySetForwardOnly(qid, forward);
```

Input

<i>qid</i>	scalar, query number.
<i>forward</i>	scalar, 1 to set forward only or 0 to allow seeking in either direction.

Remarks

Forward only mode can be (depending on the driver) more memory efficient since results do not need to be cached. It will also improve performance on some databases. For this to be true, you must call **dbQuerySetForwardOnly()** before the query is prepared or executed.

Forward only mode is enabled by default.

Setting forward only to false is a suggestion to the database engine, which has the final say on whether a result set is forward only or scrollable.

dbQueryIsForwardOnly() will always return the correct status of the result set.

See Also

[dbQueryIsForwardOnly](#)

dbQuerySetNumericalPrecisionPolicy

Purpose

Sets the current precision policy for a query.

Format

```
dbQuerySetNumericalPrecisionPolicy(query_num, prec_  
policy);
```

Input

<i>query_num</i>	scalar, query number.
<i>prec_</i> <i>policy</i>	scalar:
0	Strings will be used to preserve precision.
1	Force 32-bit integer values.
2	Force 64-bit integer values.
4	Force double values. This is the default policy.

dbRemoveDatabase

dbRemoveDatabase

Purpose

Removes a database connection from the list of open database connections. Frees all related resources.

Format

```
dbRemoveDatabase(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

dbRollback

Purpose

Rolls back a transaction on the database.

Format

```
ret = dbRollback(db_id);
```

Input

<i>db_id</i>	scalar, database connection index number.
--------------	---

Output

<i>ret</i>	scalar, 1 to indicate success and a 0 if the rollback fails.
------------	--

Remarks

A rollback is only possible if the SQL driver supports transactions and a **dbTransaction()** has been started.

Note: For some databases, the rollback will fail and return 0 if there is an active query using the database for a **SELECT**. Make the query inactive before doing the rollback.

Call **dbGetLastError()** to get information about errors.

dbSetConnectOptions

Purpose

Sets database-specific options.

Format

```
dbSetConnectOptions(db_id, db_options);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>db_options</i>	string, a semi-colon separated list of option names or option=value pairs. Available options will depend upon the database being used.

dbSetConnectOptions

Remarks

This must be done before the connection is opened or it has no effect (or you can **dbClose()** the connection, call this function and **dbOpen()** the connection again). The format of the options string is a semicolon separated list of option names or option=value pairs. The options depend on the database client used:

```
ODBC
SQL_ATTR_ACCESS_MODE
SQL_ATTR_LOGIN_TIMEOUT
SQL_ATTR_CONNECTION_TIMEOUT
SQL_ATTR_CURRENT_CATALOG
SQL_ATTR_METADATA_ID
SQL_ATTR_PACKET_SIZE
SQL_ATTR_TRACEFILE
SQL_ATTR_TRACE
SQL_ATTR_CONNECTION_POOLING
SQL_ATTR_ODBC_VERSION
```

```
MySQL
CLIENT_COMPRESS
CLIENT_FOUND_ROWS
CLIENT_IGNORE_SPACE
CLIENT_SSL
CLIENT_ODBC
CLIENT_NO_SCHEMA
CLIENT_INTERACTIVE
UNIX_SOCKET
MYSQL_OPT_RECONNECT
```

```
PostgreSQL
connect_timeout
options
tty
requiressl
service
```

```
DB2
SQL_ATTR_ACCESS_MODE
SQL_ATTR_LOGIN_TIMEOUT

OCI
OCI_ATTR_PREFETCH_ROWS
OCI_ATTR_PREFETCH_MEMORY

TDS
none

SQLite
SQLITE_BUSY_TIMEOUT
SQLITE_OPEN_READONLY
SQLITE_ENABLE_SHARED_CACHE

Interbase
ISC_DPB_LC_CTYPE
ISC_DPB_SQL_ROLE_NAME
```

Example

```
// MySQL connection
// use an SSL connection to the server
dbSetConnectOptions(db_id, "CLIENT_SSL=1;
    CLIENT_IGNORE_SPACE=1");

if not dbOpen();
    // clears the connect option string
    dbSetConnectOptions(db_id, "");
    ...
endif;

...
```

dbSetDatabaseName

```
// PostgreSQL connection
// enable PostgreSQL SSL connections
dbSetConnectOptions(db_id, "requiressl=1");
if not dbOpen();
    // clear options
    dbSetConnectOptions(db_id, "");
    ...
endif;

...

// ODBC connection
dbSetConnectOptions(db_id, "SQL_ATTR_ACCESS_MODE=
    SQL_MODE_READ_ONLY;
    SQL_ATTR_TRACE=
    SQL_OPT_TRACE_ON");
// set ODBC options
if not dbOpen();
    // don't try to set this option
    dbSetConnectOptions(db_id, "");
    ...
endif;
```

dbSetDatabaseName

Purpose

Sets the connection's database name to name. To have effect, the database name must be set before the connection is opened. Alternatively, you can **dbClose()** the connection, set the database name, and call **dbOpen()** again.

Format

```
dbSetDatabaseName(db_id, database_name);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>database_name</i>	string, database name to apply to specified database connection.

Remarks

For the OCI (Oracle) driver, the database name is the TNS Service Name.

For the ODBC driver, the name can either be a DSN, a DSN filename (in which case the file must have a .dsn extension), or a connection string.

For example, Microsoft Access users can use the following connection string to open an .mdb file directly, instead of having to create a DSN entry in the ODBC manager:

```
...
db_id = dbAddDatabase("ODBC");
dbSetDatabaseName(db_id, "DRIVER=
    {Microsoft Access Driver (*.mdb)};
    FIL={MS Access};
    DBQ=myaccessfile.mdb");
dbOpen(db_id);
...
```

See Also

[dbGetDatabaseName](#)

dbSetHostName

Purpose

Sets the specified database connection's host name.

dbSetPassword

Format

```
dbSetHostName(db_id, host_name);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>host_name</i>	string, the name to which the specified connection's host name should be assigned.

Remarks

For this function to have an effect, it must be called before the database connection is opened with **dbOpen ()**.

dbSetPassword

Purpose

Sets the database connection's password.

Format

```
dbSetPassword(db_id, pswd);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>pswd</i>	string, password for database.

Remarks

This function must be called before the connection is opened with **dbOpen ()** to have an effect.

See Also

[dbGetPassword](#)

dbSetPort

Purpose

Sets the specified database connection's port number.

Format

```
dbSetPort(db_id, port_num);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>port_num</i>	scalar, port number for database connection to use.

Remarks

This function must be called before the connection is opened with **dbOpen ()** to have an effect.

See Also

[dbGetPort](#)

dbSetUserName

dbSetUserName

Purpose

Sets the specified database connection's user name.

Format

```
dbSetUserName(db_id, user_name);
```

Input

<i>db_id</i>	scalar, database connection index number.
<i>user_name</i>	string, user name to apply to specified database connection.

Remarks

This function must be called before the connection is opened with **dbOpen ()** to have an effect.

See Also

[dbGetUserName](#)

dbTransaction

Purpose

Begins a transaction on the database.

Format

```
ret = dbTransaction(db_id);
```

Input

<code>db_id</code>	scalar, database connection index number.
--------------------	---

Output

<code>ret</code>	scalar, 1 to indicate success and a 0 if the transaction fails.
------------------	---

Example

```
//If 'dbTransaction' succeeds
if dbTransaction(db_id);
    // All queries must succeed, or all fail.
    if not dbExecQuery(db_id,
        "INSERT INTO TEST...");
        dbRollback(db_id);
        errorlog("Query 1 failed");
    end;
endif;

if not dbExecQuery(db_id,
    "INSERT INTO TEST...");
    dbRollback(db_id);
    errorlog("Query 2 failed");
end;
endif;

dbCommit(db_id);
endif;
```

Remarks

This function can only be used with databases that support transactions.

debug

See Also

[dbCommit](#), [dbRollback](#)

debug

Purpose

Runs a program under the source level debugger.

Format

```
debug filename;
```

Input

<i>filename</i>	Literal, name of file to debug.
-----------------	---------------------------------

Remarks

See **Debugging**, Section 1.1.

declare

Purpose

Initializes global variables at compile time.

Format

```
declare [[type]] symbol [[aop clist]];
```

Input

<i>type</i>	<p>optional literal, specifying the type of the symbol.</p> <p><code>matrix</code> <code>string</code> <code>array</code> <code>sparse matrix</code> <code>struct structure_type</code></p> <p>if <i>type</i> is not specified, <code>matrix</code> is assumed. Set <i>type</i> to <code>string</code> to initialize a string or string array variable.</p>
<i>symbol</i>	the name of the symbol being declared.
<i>aop</i>	<p>the type of assignment to be made.</p> <p><code>=</code> if not initialized, initialize. If already initialized, reinitialize.</p> <p><code>!=</code> if not initialized, initialize. If already initialized, reinitialize.</p> <p><code>:=</code> if not initialized, initialize. If already initialized, redefinition error.</p> <p><code>?=</code> if not initialized, initialize. If already initialized, leave as is.</p> <p>If <i>aop</i> is specified, <i>clist</i> must be also.</p>
<i>clist</i>	<p>a list of constants to assign to <i>symbol</i>.</p> <p>If <i>aop clist</i> is not specified, <i>symbol</i> is initialized as a scalar 0 or a null string.</p>

Remarks

The `declare` syntax is similar to the `let` statement.

declare

`declare` generates no executable code. This is strictly for compile time initialization. The data on the right-hand side of the equal sign must be constants. No expressions or variables are allowed.

`declare` statements are intended for initialization of global variables that are used by procedures in a library system.

It is best to place `declare` statements in a separate file from procedure definitions. This will prevent redefinition errors when rerunning the same program without clearing your workspace.

The optional `aop` and `clist` arguments are allowed only for declaring matrices, strings, and string arrays. When you `declare` an N-dimensional array, sparse matrix, or structure, they will be initialized as follows:

Variable Type	Initializes To
N-dimensional array	1-dimensional array of 1 containing 0
sparse matrix	empty sparse matrix
structure	structure containing empty and/or zeroed out members

Complex numbers can be entered by joining the real and imaginary parts with a sign (+ or -); there should be no spaces between the numbers and the sign. Numbers with no real part can be entered by appending an 'i' to the number.

There should be only one declaration for any symbol in a program. Multiple declarations of the same symbol should be considered a programming error. When **GAUSS** is looking through the library to reconcile a reference to a matrix or a string, it will quit looking as soon as a symbol with the correct name is found. If another symbol with the same name existed in another file, it would never be found. Only the first one in the search path would be available to programs.

Here are some of the possible uses of the three forms of declaration:

`!=,` Interactive programming or any situation where a global by the

- = same name will probably be sitting in the symbol table when the file containing the `declare` statement is compiled. The symbol will be reset.
- := Redefinition is treated as an error because you have probably just outsmarted yourself. This will keep you out of trouble because it won't allow you to zap one symbol with another value that you didn't know was getting mixed up in your program. You probably need to rename one of them.
- ?= Interactive programming where some global defaults were set when you started and you don't want them reset for each successive run even if the file containing the `declare`'s gets recompiled. This can get you into trouble if you are not careful.

The `declare` statement warning level is a compile option. Call **config** in the command line version of **GAUSS** or select Preferences from the Configure menu in the graphical user interface to edit this option. If `declare` warnings are on, you will be warned whenever a `declare` statement encounters a symbol that is already initialized. Here's what happens when you declare a symbol that is already initialized when `declare` warnings are turned on:

```
declare !=      Reinitialize and warn.  
declare :=      End program with fatal error.  
declare ?=      Leave as is and warn.
```

If `declare` warnings are off, no warnings are given for the `!=` and `?=` cases.

Example

```
declare matrix x,y,z;  
  
x = 0    y = 0    z = 0  
declare string x = "This string.";
```

declare

```
x = "This string."

declare matrix x;

x = 0

//Initialize 'x' with the specified values and
//return a warning if 'x'already exists AND
//the 'Compile Options: declare warnings' is
//selected
declare matrix x != { 1 2 3, 4 5 6, 7 8 9 };

    1 2 3
x = 4 5 6
    7 8 9

declare matrix x[3,3] = 1 2 3 4 5 6 7 8 9;

    1 2 3
x = 4 5 6
    7 8 9

declare matrix x[3,3] = 1;

    1 1 1
x = 1 1 1
    1 1 1

declare matrix x[3,3];

    0 0 0
x = 0 0 0
    0 0 0

declare matrix x = 1 2 3 4 5 6 7 8 9;
```



```
1
2
3
x = 4
5
6
7
8
9

//Create a 2x1 character matrix
declare matrix x = alpha beta;

//To print character matrices, the '$' operator must
//be prepended to the variable name
print $x;
```

The code snippet directly above, produces:

```
ALPHA
BETA

//Since this is declared as a matrix, the text in
//quotes will create a character vector, rather
//than a string array
declare matrix x = "mean" "variance";

print $x;
```

produces:

```
mean variance
declare array a;
```

a is a 1-dimensional array of 1 containing 0.

delete

```
declare sparse matrix sm;
```

sm is an empty sparse matrix.

```
struct mystruct {  
    matrix m;  
    string s;  
    string array sa;  
    array a;  
    sparse matrix sm;  
};
```

```
declare struct mystruct ms;
```

ms is a **mystruct** structure, with its members set as follows:

<i>ms.m</i>	empty matrix
<i>ms.s</i>	null string
<i>ms.sa</i>	1x1 string array containing a null string
<i>ms.a</i>	1-dimensional array of 1 containing 0
<i>ms.sm</i>	empty sparse matrix

See Also

[let](#), [external](#)

delete

Purpose

Deletes global symbols from the symbol table.

Format

```
delete -flags symbol_list;
delete symbol_list;
```

Input

<i>flags</i>	specify the type(s) of symbols to be deleted
<i>p</i>	procedures
<i>k</i>	keywords
<i>f</i>	fn functions
<i>m</i>	matrices
<i>s</i>	strings
<i>g</i>	only procedures with global references
<i>l</i>	only procedures with all local references
<i>n</i>	no pause for confirmation
<i>symbol</i>	literal, name of symbol to be deleted. If symbol ends in an asterisk, all symbols matching the leading characters will be deleted.

Remarks

This completely and irrevocably deletes a symbol from **GAUSS**'s memory and work-space.

Flags must be preceded by a dash (e.g. *-pfk*). If the *n* (no pause) flag is used, you will not be asked for confirmation for each symbol.

This command is supported only from interactive level. Since the interpreter executes a compiled pseudo-code, this command would invalidate a previously compiled code image and therefore would destroy any program it was a part of. If any symbols are

delete (dataloop)

deleted, all procedures, keywords and functions with global references to those symbols will be deleted as well.

Example

```
//Create a matrix 'x'
x = { 1, 2, 3, 4 };

//'show' returns information about active symbols
show x;
```

This should return:

```
32 bytes   x           MATRIX           4,1

delete -m x;
```

At the Delete?[Yes No Previous Quit] prompt, enter y.

```
show x;
```

x no longer exists.

delete (dataloop)

Purpose

Removes specific rows in a data loop based on a logical expression.

Format

```
delete logical_expression;
```

Remarks

Deletes only those rows for which *logical_expression* is TRUE. Any variables referenced must already exist, either as elements of the source data set, as [extern](#)'s, or as the result of a previous [make](#), [vector](#), or **code** statement.

GAUSS expects *logical_expression* to return a row vector of 1's and 0's. The relational and other operators (e.g. <) are already interpreted in terms of their dot equivalents (. <), but it is up to the user to make sure that function calls within *logical_expression* result in a vector.

Example

```
delete age < 40 or sex == 'FEMALE';
```

See Also

[select \(dataloop\)](#)

deleteFile

Purpose

Deletes files.

Format

```
ret = deleteFile(name);
```

Input

<i>name</i>	string or NxK string array, name of file or files to delete.
-------------	--

delif

Output

<i>ret</i>	scalar or NxK matrix, 0 if successful.
------------	--

Remarks

The return value, *ret*, is scalar if *name* is a string. If *name* is an NxK string array, *ret* will be an NxK matrix reflecting the success or failure of each separate file deletion.

deleteFile calls the C library **unlink** function for each file. If **unlink** fails it sets the C library errno value. **deleteFile** returns the value of errno if **unlink** fails, otherwise it returns zero. If you want detailed information about the reason for failure, consult the C library **unlink** documentation for your platform for details.

delif

Purpose

Deletes rows from a matrix. The rows deleted are those for which there is a 1 in the corresponding row of *e*.

Format

y = **delif**(*x*, *e*);

Input

<i>x</i>	NxK data matrix.
<i>e</i>	Nx1 logical vector (vector of 0's and 1's).

Output

y

MxK data matrix consisting of the rows of y for which there is a 0 in the corresponding row of e . If no rows remain, **delif** will return a scalar missing.

Example

Example 1: Basic usage with column vector

```
//Create column vector
x = { 1.5,
      0.8,
      0.7,
      1.2,
      1.9,
      0.2,
      2.0 };

//Create logical vector of 1's and 0's
e = x .> 1;

//Assign 'new_x' to be equal to 'x'
//with the rows removed in which 'e' equals 1
new_x = delif(x, e);
```

After the code above, *new_x* should equal:

```
0.8
0.7
0.2
```

Example 2: Matrix case

In this example, we will remove all observations in which the value of the third column is 3.

delif

```
//Create a matrix with 3 columns
x = { 20    10    2,
      33    13    3,
      37    12    2,
      34    12    3,
      35     8    1,
      25    15    2,
      34     8    2,
      37     8    1,
      37     3    1,
      31     4    1 };

//Create logical vector of 1's and 0's
e = x[:,3] .== 3;

//Assign 'new_x' to be equal to 'x' without
//the rows in which the third column equals 3
new_x = delif(x, e);
```

After the code above, *new_x* should be equal to:

20	10	2
37	12	2
35	8	1
25	15	2
34	8	2
37	8	1
37	3	1
31	4	1

Example 3: Create new 'x' and 'y' based on 'y'

In this example, we will remove all observations from *x* and *y* in which the value of the third column is 3.

```
//Create 'y' matrix
y = { 1,
      1,
      0,
      2,
      0,
      1,
      1,
      0,
      0,
      2 };

//Create 'x' matrix
x = { 1.6841 -0.1203,
      -1.0433 0.2564,
      1.2207 -1.4388,
      0.7423 0.2133,
      0.7288 1.0434,
      0.8115 1.8166,
      -0.3230 1.4763,
      1.2944 0.7635,
      1.3839 0.6648,
      -0.6330 0.4845 };

//Create logical vector of 1's and 0's
e = y .== 2;

//Assign 'x' to be equal to 'x' without
//the rows in which 'y' equals 2
x = delif(x, e);
```

delif

```
//Remove all observations in which 'y' equals 2  
y = delif(y, e);
```

After the code above, y and x should equal:

```
y = 1    x = 1.6841  -0.1203  
    1      -1.0433   0.2564  
    0       1.2207  -1.4388  
    0       0.7288   1.0434  
    1       0.8115   1.8166  
    1      -0.3230   1.4763  
    0       1.2944   0.7635  
    0       1.3839   0.6648
```

Example 4: Logical comparison of multiple columns

```
x = { 0 10 20,  
      30 40 50,  
      60 70 80 };  
  
//Logical vector, comparing two columns  
e =(x[:,1] .gt 0) .and (x[:,3] .lt 100);  
  
y = delif(x,e);
```

After the code above:

```
y = 0 10 20
```

All rows for which the elements in column 1 are greater than 0 and the elements in column 3 are less than 100 are deleted.

Remarks

The input `e` will usually be generated by a logical expression using dot operators. For instance:

```
//Create a vector 'e' with a 1 for each row in which the
//value in the second column of 'x' is less than 100,
//otherwise a 0
e = x[:,2] .> 100;

y = delif(x, e);
```

Or the equivalent statement:

```
y = delif(x, x[:,2] .> 100);
```

will delete all rows of `x` whose second element is greater than 100. The remaining rows of `x` will be assigned to `y`.

See Also

[selif](#)

delrows

Purpose

Deletes rows from a matrix. The second argument contains the indices of the rows to be deleted.

Format

```
y = delrows(x, r);
```

Input

<code>x</code>	NxK data matrix.
<code>r</code>	Mx1 vector, indices of rows to delete.

delrows

Output

y

PxK matrix containing the remaining rows of x . If no rows remain, y will be an empty matrix.

Remarks

If r is an empty matrix, the result will be unchanged. Negative values of r are counted from the end of the matrix, therefore:

```
r = -1;  
  
y = delrows(x, r);
```

will delete last row of x . The remaining rows of x will be assigned to y .

Example

```
x = { 0 10 20,  
      30 40 50,  
      32 42 52,  
      35 45 55,  
      60 70 80 };  
  
r = { 2,  
      4 };  
  
y = delrows(x, r);
```

After the code above:

```
y =    0 10 20  
      32 42 52  
      60 70 80
```

Rows 2 and 4 are deleted.

See Also

[delif](#)

denseToSp

Purpose

Converts a dense matrix to a sparse matrix.

Format

```
y = denseToSp(x, eps);
```

Input

<i>x</i>	MxN dense matrix.
<i>eps</i>	scalar, elements of <i>x</i> whose absolute values are less than or equal to <i>eps</i> will be treated as zero.

Output

<i>y</i>	MxN sparse matrix.
----------	--------------------

Remarks

A dense matrix is just a normal format matrix.

Since sparse matrices are strongly typed in **GAUSS**, *y* must be defined as a sparse matrix before the call to **denseToSp**.

denseToSpRE

Example

```
//Declare 'y' as a sparse matrix
sparse matrix y;

x = { 0.01 0.00 0.01 1.00,
      0.00 4.00 0.02 0.00,
      0.00 0.01 0.00 0.00,
      0.02 0.00 -2 0.00 };

//Create a sparse matrix 'y' from 'x' and set all elements
//less than 0.04 equal to 0
y = denseToSp(x, 0.04);
```

After the code above, *y* is equal to:

0.00	0.00	0.00	1.00
0.00	4.00	0.00	0.00
0.00	0.00	0.00	0.00
0.00	0.00	-2.00	0.00

See Also

[spCreate](#), [spDenseSubmat](#), [spToDense](#)

denseToSpRE

Purpose

Converts a dense matrix to a sparse matrix, using a relative epsilon.

Format

```
y = denseToSpRE(x, reps);
```

Input

<i>x</i>	MxN dense matrix.
<i>reps</i>	scalar, relative epsilon. Elements of <i>x</i> will be treated as zero if their absolute values are less than or equal to <i>reps</i> multiplied by the mean of the absolute values of the non-zero values in <i>x</i> .

Output

<i>y</i>	MxN sparse matrix.
----------	--------------------

Remarks

A dense matrix is just a normal format matrix.

Since sparse matrices are strongly typed in **GAUSS**, *y* must be defined as a sparse matrix before the call to **denseToSpRE**.

Example

```
sparse matrix y;  
x = { -9   0   0   1,  
      0   4   0   0,  
      5   0   0   7,  
      0   0  -2  2.2 };  
  
y = denseToSpRE(x, 0.5);  
d = spToDense(y);
```

After the code above, *d* is equal to:

denToZero

-9.00	0.00	0.00	0.00
0.00	4.00	0.00	0.00
5.00	0.00	0.00	7.00
0.00	0.00	0.00	2.20

You can calculate the mean of the non-zero elements of x like this:

```
//Create a matrix of 1's and 0's with a 1 where the
//corresponding element in 'x' is not equal to 0
mask = x ./= 0;

//Calculate the sum of 'mask', this is the number of
//non-zeros in 'x'
nnz = sumc(sumc(mask));

//Divide the sum of the absolute value of 'x' by the number
//of non-zeros
nzmean = sumc(sumc(abs(x)))/nnz;
```

```
nnz =          7
nzmean = 4.31
```

The call to **denseToSpRE** towards the start of this example, removed all non-zeros less than $0.5 * nzmean$, or approximately 2.16.

See Also

[denseToSp](#), [spCreate](#), [spToDense](#)

denToZero

Purpose

Converts every denormal to a 0 in a matrix or array.

Format

```
y = denToZero(x);
```

Input

x	A matrix or an N-dimensional array.
-----	-------------------------------------

Output

y	A matrix or an N-dimensional array with the same orders as the input. Every denormal in the input will be converted to 0 in the output.
-----	---

Example

```
x = { 1, exp(-724.5), 3 };

//If 'x' contains any denormals set them to 0
if isden(x);
    x2 = denToZero(x);
endif;
```

After the first line above, x is equal to:

```
1.000e+000
2.902e-057
3.000e+000
```

At the end of the example, x is equal to:

```
1.000e+000
0.000e+000
3.000e+000
```

design

design

Purpose

Creates a design matrix of 0's and 1's from a column vector of numbers specifying the columns in which the 1's should be placed.

Format

```
y = design(x);
```

Input

x	$N \times 1$ vector.
-----	----------------------

Output

y	$N \times K$ matrix, where $K = \max(\mathbf{x})$; each row of y will contain a single 1, and the rest 0's. The one in the i th row will be in the $\text{round}(x[i,1])$ column.
-----	--

Remarks

Note that x does not have to contain integers: it will be rounded to nearest if necessary.

Example

This example uses **design** to interchange the rows of a matrix.

```
//Suppress printing of digits after the decimal place
format /rd 6,0;

//Set the rng seed for repeatable random numbers
```

```
rndseed 345425235;

//Create a 4x4 matrix of random integers with a standard
//deviation of 10
x = round(10*randn(4,4));
print x;
```

The code above returns:

4	12	-1	-10
5	-3	12	8
12	-2	21	-21
-7	-13	0	-1

Contintuing on with the example:

```
//The order of the rows we want
rowOrder = { 3, 1, 4, 2 };

//Create a permutation matrix from 'rowOrder'
p = design(rowOrder);
print p;
```

This section returns:

0	0	1	0
1	0	0	0
0	0	0	1
0	1	0	0

```
//Create a permuted version of 'x' with our preferred row
//order
x2 = p*x;
print x2;
```

This final section returns:

det

```
12      -2      21      -21
 4       12      -1      -10
-7      -13       0       -1
 5       -3      12        8
```

This last print statement shows us that we have indeed changed the order of the rows. In `x` the row order is 1, 2, 3, 4. However, in `x2`, the row order is 3, 1, 4, 2 (i.e. the third row is now first, the first row is now second, etc.)

Source

`design.src`

See Also

[cumprodc](#), [cumsumc](#), [recserrc](#)

det

Purpose

Returns the determinant of a square matrix.

Format

```
y = det(x);
```

Input

<code>x</code>	NxN square matrix or K-dimensional array where the last two dimensions are NxN .
----------------	--

Output

<code>y</code>	scalar or [K-2]-dimensional array, the determinant(s) of
----------------	--

x .

Remarks

x may be any valid expression that returns a square matrix (number of rows equals number of columns) or a K-dimensional array where the last two dimensions are of equal size.

If x is a K-dimensional array, the result will be a [K-2]-dimensional array containing the determinants of each 2-dimensional array described by the two trailing dimensions of x . In other words, for a 10x4x4 array, the result will be a 1-dimensional array of 10 elements containing the determinants of each of the 10 4x4 arrays contained in x .

det computes an LU decomposition.

detl can be much faster in many applications.

Example

```
x = { 3 2 1,  
      0 1 -2,  
      1 3 4 };  
y = det(x);  
  
format /rd 3,0;  
print "The determinant of y =" y;
```

The code above, produces:

```
The determinant of y = 25
```

See Also

[detl](#)

detl

detl

Purpose

Returns the determinant of the last matrix that was passed to one of the intrinsic matrix decomposition routines.

Format

```
y = detl;
```

Remarks

Whenever one of the intrinsic matrix decomposition routines is executed, the determinant of the matrix is also computed and stored in a system variable. This function will return the value of that determinant and, because the value has been computed in a previous instruction, this will require no computation.

The following functions will set the system variable used by **detl**:

```
chol(x)  
crout(x)  
croutp(x)  
det(x)  
inv(x)  
invpd(x)  
solpd(y, x)      determinant of x
```

Example

If both the inverse and the determinant of the matrix are needed, the following two commands will return both with the minimum amount of computation:

```
xi = inv(x);
xd = det1;
```

The function **det**(*x*) returns the determinant of a matrix using the Crout decomposition. If you only want the determinant of a positive definite matrix, the following code will be the fastest for matrices larger than 10x10:

```
//The 'call' keyword tells GAUSS to ignore the values
//returned from chol
call chol(x);
xd = det1;
```

The Cholesky decomposition is computed and the result from that is discarded. The determinant saved during that instruction is retrieved using **det1**. This can execute up to 2.5 times faster than **det**(*x*) for large positive definite matrices.

See Also

[det](#)

dfft

Purpose

Computes a discrete Fourier transform.

Format

```
y = dfft(x);
```

Input

<i>x</i>	Nx1 vector.
----------	-------------

dffti

Output

y	Nx1 vector.
-----	-------------

Remarks

The transform is divided by N.

This uses a second-order Goertzel algorithm. It is considerably slower than **fft**, but it may have some advantages in some circumstances. For one thing, N does not have to be an even power of 2.

Source

`dffti.src`

See Also

[dffti](#), [fft](#), [ffti](#)

dffti

Purpose

Computes inverse discrete Fourier transform.

Format

$y = \text{dffti}(x);$

Input

x	Nx1 vector.
-----	-------------

Output

y	Nx1 vector.
-----	-------------

Remarks

The transform is divided by N.

This uses a second-order Goertzel algorithm. It is considerably slower than **ffti**, but it may have some advantages in some circumstances. For one thing, N does not have to be an even power of 2.

Source

`dffti.src`

See Also

[fft](#), [dffti](#), [ffti](#)

diag

Purpose

Creates a column vector from the diagonal of a matrix.

Format

$y = \text{diag}(x);$

Input

x	NxK matrix or L-dimensional array where the last two dimensions are NxK.
-----	--

diag

Output

y

$\min(N,K) \times 1$ vector or L-dimensional array where the last two dimensions are $\min(N,K) \times 1$.

Remarks

If x is a matrix, it need not be square. Otherwise, if x is an array, the last two dimensions need not be equal.

If x is an array, the result will be an array containing the diagonals of each 2-dimensional array described by the two trailing dimensions of x . In other words, for a $10 \times 4 \times 4$ array, the result will be a $10 \times 4 \times 1$ array containing the diagonals of each of the $10 \times 4 \times 4$ arrays contained in x .

diagrv reverses the procedure and puts a vector into the diagonal of a matrix.

Example

Get the diagonal from a matrix.

```
rndseed 458716;  
x = randu(3,3);  
y = diag(x);  
print "x = " x;  
print "y = " y;
```

After the above code,

```
x =  
0.96748215 0.31791692 0.46520760  
0.04558545 0.78613263 0.20528802  
0.73825699 0.30528745 0.73350290  
y =
```

```
0.96748215
0.78613263
0.73350290
```

Using **diag** function for a 3x4x4 dimensional array.

```
x = randn(48,1);

//Reshape the 48x1 vector into a 3x4x4 dimensional array
x = areshape(x, 3|4|4);
d = diag(x);
```

Now x is equal to:

Plane [1,...]

0.082720153	-0.49502230	-0.40613944	1.9283280
0.23583965	-0.24230946	-0.66047073	-0.73098141
-1.1187279	-0.27867822	-1.7846293	-0.44603382
0.030071777	-1.0387861	0.23768949	0.019151917

Plane [2,...]

-1.7238416	0.17660645	-0.14798006	0.072065419
1.3685721	-0.11216325	-0.12985589	1.1816008
0.63154571	-1.4945397	-1.7276380	-0.28275797
-0.71832623	-1.3193506	-0.53934998	-0.78348484

Plane [3,...]

-0.71111209	-0.30818842	-0.38982318	-2.7205066
-1.5455077	-0.27131853	0.98686691	0.10870999
0.57916876	1.8180884	0.76104693	1.1237605
1.0727710	-1.1071168	1.7443178	-1.0684433

and d is a 3x4x1 array containing the diagonals from x above.

diagrv

```
Plane [1,...]

    0.082720153
   -0.24230946
   -1.7846293
    0.019151917

Plane [2,...]

   -1.7238416
  -0.11216325
   -1.7276380
  -0.78348484

Plane [3,...]

   -0.71111209
  -0.27131853
    0.76104693
   -1.0684433
```

See Also

[diagrv](#)

diagrv

Purpose

Inserts a vector into the diagonal of a matrix.

Format

```
y = diagrv(x, v);
```

Input

x	$N \times K$ matrix.
v	$\min(N,K) \times 1$ vector.

Output

y	$N \times K$ matrix equal to x with its principal diagonal elements equal to those of v .
-----	---

Remarks

diag reverses the procedure and pulls the diagonal out of a matrix.

Example

```
x = randu(3,3);  
v = ones(3,1);  
y = diagrv(x,v);
```

After the code above:

```
0.614 0.686 0.633    1.000    1.000 0.686 0.633  
x = 0.802 0.185 0.707 v = 1.000 y = 0.802 1.000 0.707  
0.551 0.761 0.418    1.000    0.551 0.761 1.000
```

See Also

[diag](#)

digamma

digamma

Purpose

Computes the digamma function.

Format

```
 $y = \text{digamma}(x);$ 
```

Input

x	MxN matrix or N-dimensional array.
-----	------------------------------------

Output

y	MxN matrix or N-dimensional array, digamma.
-----	---

Remarks

The digamma function is the first derivative of the log of the gamma function with respect to its argument.

dlibrary

Purpose

Dynamically links and unlinks shared libraries.

Format

```
dlibrary lib1 [[lib2]]...;  
dlibrary -a lib1 [[lib2]]...;  
dlibrary -d;  
dlibrary;
```

Input

<i>lib1</i>	literal, the base name of the library or the pathed name
<i>lib2...</i>	of the library.
	<p><code>dlibrary</code> takes two types of arguments, "base" names and file names. Arguments without any "/" path separators are assumed to be library base names, and are expanded by adding the suffix <code>.so</code>, <code>.dll</code> or <code>.dylib</code>, depending on the platform. They are searched for in the default dynamic library directory. Arguments that include "/" path separators are assumed to be file names, and are not expanded. Relatively pathed file names are assumed to be specified relative to the current working directory, not relative to the dynamic library directory.</p>
<i>-a</i>	append flag, the shared libraries listed are added to the current set of shared libraries rather than replacing them. For search purposes, the new shared libraries follow the already active ones. Without the <i>-a</i> flag, any previously linked libraries are dumped.
<i>-d</i>	dump flag, ALL shared libraries are unlinked and the functions they contain are no longer available to your programs. If you use <code>dllcall</code> to call one of your functions after executing a

```
dlibrary -d
```

dlibrary

your program will terminate with an error.

Remarks

- If no flags are used, the shared libraries listed are linked into **GAUSS** and any previously linked libraries are dumped. When you call `dllcall`, the shared libraries will be searched in the order listed for the specified function. The first instance of the function found will be called.
- `dlibrary` with no arguments prints out a list of the currently linked shared libraries. The order in which they are listed is the order in which they are searched for functions.
- `dlibrary` recognizes a default directory in which to look for dynamic libraries. You can specify this by setting the variable `dlib_path` in `gauss.cfg`. Set it to point to a single directory, not a sequence of directories. **sysstate**, case 24, may also be used to get and set this default.
- **GAUSS** maintains its own shared libraries which are listed when you execute `dlibrary` with no arguments, and searched when you call `dllcall`. The default shared library or libraries are searched last. You can force them to be searched earlier by listing them explicitly in a `dlibrary` statement. They are always active and are not unlinked when you execute

```
dlibrary -d
```

For more information, see **FOREIGN LANGUAGE INTERFACE, CHAPTER 1**.

Examples

Example 1: Loading a shared library and unloading previously loaded shared libraries

```
dlibrary mylib;
```


The above command passes the base name of the shared library to load. **GAUSS** will expand this base name to a platform specific shared library name. The expanded name on Windows is **mylib.dll**. On Linux it is **libmylib.so** and on Mac, **libmylib.dylib**. Since we did not pass the `-a` flag, **GAUSS**, will unload any shared libraries that were previously loaded with the `dlibrary` command.

Example 2: Loading a shared library and keeping previously loaded shared libraries

```
dlibrary -a mylib;
```

Since we passed the `-a` flag, **GAUSS** will not unload any libraries when it loads **mylib**.

See Also

[dllcall](#), [sysstate](#)

dllcall

Purpose

Calls functions located in dynamic libraries.

Format

```
dllcall [-r] [-v] func(arg1...argN);
```

`dllcall` works in conjunction with `dlibrary`. `dlibrary` is used to link shared libraries into **GAUSS**; `dllcall` is used to access the functions contained in those shared libraries. `dllcall` searches the shared libraries (see `dlibrary` for an explanation of the search order) for a function named `func`, and calls the first instance it finds. The default shared libraries are searched last.

dllcall

Input

<i>func</i>	the name of a function contained in a shared library (linked into GAUSS with dlibrary). If <i>func</i> is not specified or cannot be located in a shared library, dllcall will fail.
<i>arg#</i>	arguments to be passed to <i>func</i> , optional. These must be simple variable references; they cannot be expressions.
<i>-r</i>	optional flag. If <i>-r</i> is specified, dllcall examines the value returned by <i>func</i> , and fails if it is nonzero.
<i>-v</i>	optional flag. Normally, dllcall passes parameters to <i>func</i> in a list. If <i>-v</i> is specified, dllcall passes them in a vector. See below for more details.

Remarks

func should be written to:

1. Take 0 or more pointers to doubles as arguments.
2. Take arguments either in a list or a vector.
3. Return an integer.

In C syntax, *func* should take one of the following forms:

1. `int func(void);`
2. `int func(double *arg1 [, arg2...argN]);`
3. `int func(double *arg[]);`

[dllcall](#) can pass a list of up to 100 arguments to **func**; if it requires more arguments than that, you **MUST** write it to take a vector of arguments, and you **MUST**

specify the `-v` flag when calling it. `dllcall` can pass up to 1000 arguments in vector format. In addition, in vector format `dllcall` appends a null pointer to the vector, so you can write `func` to take a variable number of arguments and just test for the null pointer.

Arguments are passed to `func` by reference. This means you can send back more than just the return value, which is usually just a success/failure code. (It also means that you need to be careful not to overwrite the contents of matrices or strings you want to preserve.) To return data from **func**, simply set up one or more of its arguments as return matrices (basically, by making them the size of what you intend to return), and inside **func** assign the results to them before returning.

For more information, see **FOREIGN LANGUAGE INTERFACE, CHAPTER 1**.

See Also

[dlibrary](#), [sysstate](#)

do while,do until

Purpose

Executes a series of statements in a loop as long as a given expression is true (or false).

do while,do until

Format

```
do while expression;
or
do until expression;
.
.
.
    statements in loop
.
.
.
endo;
```

Remarks

expression is any expression that returns a scalar. It is TRUE if it is nonzero and FALSE if it is zero.

In a `do while` loop, execution of the loop will continue as long as the expression is TRUE.

In a `do until` loop, execution of the loop will continue as long as the expression is FALSE.

The condition is checked at the top of the loop. If execution can continue, the statements of the loop are executed until the `endo` is encountered. Then **GAUSS** returns to the top of the loop and checks the condition again.

The `do` loop does not automatically increment a counter. See the first example below.

`do` loops may be nested.

It is often possible to avoid using loops in **GAUSS** by using the appropriate matrix operator or function. It is almost always preferable to avoid loops when possible, since the corresponding matrix operations can be much faster.

Example

```
format /rdn 1,0;
space = " ";
comma = ",";
i = 1;
do while i <= 4;
    j = 1;
    do while j <= 3;
        print space i comma j;;
        j = j+1;
    endo;
    i = i+1;
    print;
endo;
```

The code above prints the following output:

```
1,1 1,2 1,3
2,1 2,2 2,3
3,1 3,2 3,3
4,1 4,2 4,3
```

In the example above, two nested loops are executed and the loop counter values are printed out. Note that the inner loop counter must be reset inside of the outer loop before entering the inner loop. An empty `print` statement is used to print a carriage return/line feed sequence after the inner loop finishes.

The following are examples of simple loops that execute a predetermined number of times. These loops will both have the result shown.

First loop:

```
format /rd 1,0;
i = 1;
do while i <= 10;
```

dos

```
        print i;;  
        i = i+1;  
    endo;
```

produces:

```
1 2 3 4 5 6 7 8 9 10
```

Second loop:

```
format /rd 1,0;  
i = 1;  
do until i > 10;  
    print i;;  
    i = i+1;  
endo;
```

produces:

```
1 2 3 4 5 6 7 8 9 10
```

See Also

[continue](#), [break](#)

dos

Purpose

Provides access to the operating system from within **GAUSS**.

Format

```
dos command;
```

Input

<i>cmd</i>	literal or ^string, the OS command to be executed.
------------	--

Portability

Linux

Control and output go to the controlling terminal, if there is one.

This function may be used in terminal mode.

Windows

The `dos` function opens a new terminal.

Running programs in the background is allowed on both of the aforementioned platforms.

Remarks

This allows all operating system commands to be used from within **GAUSS**. It allows other programs to be run even though **GAUSS** is still resident in memory.

If no operating system command (for instance, **dir** or **copy**) or program name is specified, then a shell of the operating system will be entered which can be used just like the base level OS. The **exit** command must be given from the shell to get back into **GAUSS**. If a command or program name is included, the return to **GAUSS** is automatic after the OS command has been executed.

All matrices are retained in memory when the OS is accessed in this way. This command allows the use of word processing, communications, and other programs from within **GAUSS**.

Do not execute programs that terminate and remain resident because they will be left resident inside of **GAUSS**'s workspace. Some examples are programs that create RAM disks or print spoolers.

dos

If the command is to be taken from a string variable, the ^ (caret) must precede the string.

The shorthand ">" can be used in place of "dos".

Example

```
cmdstr = "atog mycfile";  
dos ^cmdstr;
```

This will run the ATOG utility, using `mycfile.cmd` as the ATOG command file. For more information, see **ATOG**, CHAPTER 1.

```
> dir *.prg;
```

This will use the DOS **dir** command to print a directory listing of all files with a `.prg` extension on Windows. When the listing is finished, control will be returned to **GAUSS**.

```
> ls *.prg
```

This will perform the same operation on Linux.

```
dos;
```

This will cause a second level OS shell to be entered. The OS prompt will appear and OS commands or other programs can be executed. To return to **GAUSS**, type **exit**.

See Also

[exec](#)

doswin

Purpose

Opens the DOS compatibility window with default settings. NOTE: This function is no longer supported. This documentation is provided as a reference for understanding legacy code. In many cases, you may simply comment out calls to **doswin** and the program will run successfully in the program input/output window.

Format

```
doswin;
```

Portability

Windows only

Remarks

Calling **doswin** is equivalent to:

```
call  
  DOSWinOpen ( "", error (0) );
```

Source

gauss.src

DOSWinCloseall

Purpose

Closes the DOS compatibility window. NOTE: The DOS compatibility window is no longer supported. This documentation is provided as a reference for

DOSWinOpen

understanding legacy code.

Format

```
DOSWinCloseall;
```

Portability

Windows only

Remarks

Calling **DOSWinCloseall** closes the DOS window immediately, without asking for confirmation. If a program is running, its I/O reverts to the Command window.

Example

```
let attr = 50 50 7 0 7;
if not DOSWinOpen("Legacy Window", attr);
    errorlog "Failed to open DOS window, aborting";
    stop;
endif;
.
.
.
DOSWinCloseall;
```

DOSWinOpen

Purpose

Opens the DOS compatibility window and gives it the specified title and attributes. NOTE: This function is no longer supported. This documentation is provided as a reference for understanding legacy code. In many cases, you may simply

comment out calls to **DOSWinOpen** and the program will run successfully in the program input/output window.

Format

```
ret = DOSWinOpen(title, attr);
```

Input

<i>title</i>	string, window title.
<i>attr</i>	5x1 vector or scalar missing, window attributes.
	[1] window x position
	[2] window y position
	[3] text foreground color
	[4] text background color
	[5] close action bit flags
	bit 0 (1's issue dialog
	bit)
	bit 1 (2's close window
	bit)
	bit 2 (4's stop program
	bit)

Output

<i>ret</i>	scalar, success flag, 1 if successful, 0 if not.
------------	--

Portability

Windows only

DOSWinOpen

Remarks

If `title` is a null string (""), the window will be titled "GAUSS-DOS".

Defaults are defined for the elements of `attr`. To use the default, set an element to a missing value. Set `attr` to a scalar missing to use all defaults. The defaults are defined as follows:

[1]	varies	use x position of previous DOS window
[2]	varies	use y position of previous DOS window
[3]	7	white foreground
[4]	0	black background
[5]	6	4+2: stop program and close window without confirming

If the DOS window is already open, the new `title` and `attr` will be applied to it. Elements of `attr` that are missing are not reset to the default values, but are left as is.

To set the close action flags value (`attr[5]`), just sum the desired bit values. For example:

stop program (4) + close window (2) + confirm close (1) = 7

The close action flags are only relevant when a user attempts to interactively close the DOS window while a program is running. If **GAUSS** is idle, the window will be closed immediately. Likewise, if a program calls **DOSWinCloseall**, the window is closed, but the program does not get terminated.

Example

```
let attr = 50 50 7 0 7;

if not DOSWinOpen("Legacy Window", attr);
    errorlog "Failed to open DOS window, aborting";
```

```
stop;  
endif;
```

This example opens the DOS window at screen location (50,50), with white text on a black background. The close action flags are $4 + 2 + 1$ (stop program + close window + issue confirm dialog) = 7. Thus, if the user attempts to close the window while a program is running, he/she will be asked for confirmation. Upon confirmation, the window will be closed and the program terminated.

dot

Purpose

Returns a scalar dot product of the columns of two matrices.

Format

```
z = dot(x, y);
```

Input

x	$N \times 1$ vector or $N \times K$ matrix.
y	$N \times 1$ vector or $N \times K$ matrix.

Output

z	scalar or $K \times 1$ dot product
-----	------------------------------------

Examples

Example 1: Basic usage

dot

```
//Create two 4x1 column vectors
x = { 5,
      9,
      3,
      4 };

y = { 9,
      -6,
      8,
      1  };

//Compute dot product
z = dot(x,y);

print "z = " z;
```

After the code above:

```
z = 19
```

Example 2: Dot product of the corresponding columns of two matrices

```
//Create two 4x2 matrices
x = { 5 1,
      9 3,
      3 8,
      4 2 };

y = { 9 8,
      -6 4,
      8 3,
      1 -2 };

//Compute dot product
z = dot(x, y);
```

```
print "z = " z;
```

After the code above:

```
z = 19
    40
```

Remarks

Inputs x and y should have the same columns.

See Also

[crossprd](#)

dotfeq, dotfge, dotfgt, dotfle, dotflt, dotfne

Purpose

Fuzzy comparison functions. These functions use `_fcmptol` to fuzz the comparison operations to allow for roundoff error.

Format

```
y = dotfeq(a, b);
y = dotfge(a, b);
y = dotfgt(a, b);
y = dotfle(a, b);
y = dotflt(a, b);
y = dotfne(a, b);
```

dotfeq,dotfge, dotfgt,dotfle,dotflt,dotfne

Input

<i>a</i>	NxK matrix, first matrix.
<i>b</i>	LxM matrix, second matrix, ExE compatible with <i>a</i> .

Global Input

<i>_fcmtol</i>	scalar, comparison tolerance. The default value is 1.0e-15.
----------------	---

Output

<i>y</i>	max(N,L) by max(K,M) matrix of 1's and 0's.
----------	---

Remarks

The return value is 1 if TRUE and 0 if FALSE.

The statement:

```
y = dotfeq(a,b);
```

is equivalent to:

```
y = a .eq b;
```

The calling program can reset *_fcmtol* before calling these procedures:

```
_fcmtol = 1e-12;
```

Example

```
x = pi*ones(2,2);
```

dotfeqmt, dotfgemt, dotfgtmt, dotflemt, dotfltmt, dotfnemt

```
y = x;
y[1,1] = 2*pi;

//Test for elements where 'x' is > 'y'
t = dotfge(x,y);

x = 3.14 3.14  y = 6.28 3.14  t = 0.00 1.00
    3.14 3.14      3.14 3.14      1.00 1.00
```

Continuing with the data above:

```
//Test for elements where 'x' is < 'y '
t = dotflt(x,y);

t = 1.00 0.00
    0.00 0.00
```

Source

fcompare.src

Globals

`_fcmptol`

See Also

[feq-fne](#)

dotfeqmt, dotfgemt, dotfgtmt, dotflemt, dotfltmt, dotfnemt

Purpose

Fuzzy comparison functions. These functions use the *fcmptol* argument to fuzz the comparison operations to allow for roundoff error.

`dotfeqmt, dotfgemt, dotfgtmt, dotflemt, dotflmt, dotfnemt`

Format

```
y = dotfeqmt(a, b, fcmptol);  
y = dotfgemt(a, b, fcmptol);  
y = dotfgtmt(a, b, fcmptol);  
y = dotflemt(a, b, fcmptol);  
y = dotflmt(a, b, fcmptol);  
y = dotfnemt(a, b, fcmptol);
```

Input

<i>a</i>	NxK matrix, first matrix.
<i>b</i>	LxM matrix, second matrix, ExE compatible with <i>a</i> .
<i>fcmptol</i>	scalar, comparison tolerance.

Output

<i>y</i>	max(N,L) by max(K,M) matrix of 1's and 0's.
----------	---

Remarks

The return value is 1 if TRUE and 0 if FALSE.

The statement:

```
y = dotfeqmt(a,b,1e-13);
```

is equivalent to:

```
y = a .eq b;
```

Example

```
x = rndu(2,2);
```

```
y = x;
y[1,1] = y[1,1] + 0.000000002;
t = dotfgemt(x,y,1e-15);
```

```
t = 0 1    x-y = -2e-8    0
    1 1          0      0
```

Source

fcomparemt.src

See Also

[feqmt-fnemt](#)

draw

Purpose

Graphs lines, symbols, and text using the PQG global variables. This procedure does not require actual X, Y, or Z data since its main purpose is to manually build graphs using *_pline*, *_pmsgctl*, *_psym*, *_paxes*, *_parrow* and other globals.

NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
draw;
```

draw

Remarks

draw is especially useful when used in conjunction with transparent windows.

Example

```
library pgraph;
graphset;

begwind;
makewind(9,6.855,0,0,0); /* make full size window for
                           plot */
makewind(3,1,3,3,0);     /* make small overlapping window
                           for text */

setwind(1);
  x = seqa(.1,.1,100);
  y = sin(x);
  xy(x,y);                /* plot data in first window */
nextwind;
  _pbox = 15;
  _paxes = 0;
  _pnum = 0;
  _ptitlht = 1;
  margin(0,0,2,0);
  title("This is a text window.");
  draw;                    /* add a smaller text window */
endwind;                  /* create graph */
```

Source

pdraw.src

See Also

[window](#), [makewind](#)

drop (dataloop)

Purpose

Specifies columns to be dropped from the output data set in a data loop.

Format

```
drop variable_list;
```

Remarks

Commas are optional in *variable_list*.

Deletes the specified variables from the output data set. Any variables referenced must already exist, either as elements of the source data set, or as the result of a previous [make](#), [vector](#), or **code** statement.

If neither [keep](#) nor [drop](#) is used, the output data set will contain all variables from the source data set, as well as any defined variables. The effects of multiple [keep](#) and [drop](#) statements are cumulative.

Example

```
drop age, pay, sex;
```

See Also

[keep \(dataloop\)](#)

dsCreate

Purpose

Creates an instance of a structure of type **DS** set to default values.

dstat

Include

`ds.sdf`

Format

```
s = dsCreate();
```

Output

s instance of structure of type **DS**.

Example

```
//Declare 'myData' as instance of 'DS' structure
struct DS myData;

//Apply default settings
myData = dsCreate();
```

Source

`ds.src`

dstat

Purpose

Computes descriptive statistics.

Format

```
{ vnam, mean, var, std, min, max, valid, mis } = dstat(dataset,  
vars);
```

Input

<i>dataset</i>	string, name of data set. If <i>dataset</i> is null or 0, <i>vars</i> will be assumed to be a matrix containing the data.
<i>vars</i>	the variables. If <i>dataset</i> contains the name of a data set, <i>vars</i> will be interpreted as: Kx1 character vector, names of variables. - or - Kx1 numeric vector, indices of variables. - or - formula string , e.g. "PAY + WT" or ". - sex" These can be any size subset of the variables in the data set and can be in any order. If a scalar 0 is passed, all columns of the data set will be used. If <i>dataset</i> is null or 0, <i>vars</i> will be interpreted as: NxK matrix, the data on which to compute the descriptive statistics.

Global Input

<code>__altnam</code>	matrix, default 0. This can be a Kx1 character vector of alternate variable names for the output.
-----------------------	--

dstat

<code>__maxbytes</code>	scalar, the maximum number of bytes to be read per iteration of the read loop. Default = 1e9.
<code>__maxvec</code>	scalar, the largest number of elements allowed in any one matrix. Default = 20000.
<code>__miss</code>	scalar, default 0. 0 there are no missing values (fastest). 1 listwise deletion, drop a row if any missings occur in it. 2 pairwise deletion.
<code>__row</code>	scalar, the number of rows to read per iteration of the read loop.
<code>__output</code>	if 0, (default) the number of rows will be calculated using <code>__maxbytes</code> and <code>__maxvec</code> . scalar, controls output, default 1. 1 print output table. 0 do not print output.

Output

<code>vnam</code>	Kx1 character vector, the names of the variables used in the statistics.
<code>mean</code>	Kx1 vector, means.
<code>var</code>	Kx1 vector, variance.
<code>std</code>	Kx1 vector, standard deviation.
<code>min</code>	Kx1 vector, minima.
<code>max</code>	Kx1 vector, maxima.
<code>valid</code>	Kx1 vector, the number of valid cases.
<code>mis</code>	Kx1 vector, the number of missing cases.

Example

Example 1

Calculate statistics on all variables in dataset

```
file = getGAUSShome() $+ "examples/freqdata.dat";
//Calculate statistics on all variables in dataset: AGE,
PAY, sex and WT
vars = 0;
{ vnam, mean, var, std, min, max, valid, mis } = dstat
(file, vars);
```

After the above code,

```
-----
-----
Variable      Mean    Std Dev    Variance    Minimum    Max-
imum      Valid Missing
-----
-----
AGE           10.0000    400      0          1.0000
PAY           1.9675    0.8019    0.6431     1.0000
3.0000       400      0
sex           1.4699    0.3007    0.0904     1.0000
1.9900       400      0
WT           1.9675    0.8019    0.6431     1.0000
3.0000       400      0
sex           1.4699    0.3007    0.0904     1.0000
1.9900       400      0
WT           1.9675    0.8019    0.6431     1.0000
3.0000       400      0
```

Example 2

```
//Calculate statistics on just AGE and PAY
vars = { AGE, PAY };
```

dstat

```
{ vnam, mean, var, std, min, max, valid, mis } = dstat
(file, vars);
```

After the above code,

Variable	Mean	Std Dev	Variance	Minimum	Max-
imum	Valid	Missing			

AGE	-----	-----	-----	1.0000	
10.0000	400	0			
PAY	1.9675	0.8019	0.6431	1.0000	
3.0000	400	0			

Example 3

```
//Calculate statistics on just AGE and PAY using numerical
indices
vars = { 1, 2 };
{ vnam, mean, var, std, min, max, valid, mis } = dstat
(file, vars);
```

After the above code,

Variable	Mean	Std Dev	Variance	Minimum	Max-
imum	Valid	Missing			

AGE	-----	-----	-----	1.0000
10.0000	400	0		
PAY	1.9675	0.8019	0.6431	1.0000
3.0000	400	0		

Example 4

```
//Calculate statistics on just AGE and PAY using __miss
vars = { 1, 2 };
//Drop rows with missing values
__miss = 1;
{ vnam, mean, var, std, min, max, valid, mis } = dstat
(file, vars);
```

After the above code,

Variable	Mean	Std Dev	Variance	Minimum	Maximum
Valid	Missing				
AGE	5.6784	2.9932	8.9593	1.0000	
10.0000	398	2			
PAY	1.9623	0.8006	0.6409	1.0000	
3.0000	398	2			

Example 5

```
//Calculate statistics using formula string and __miss
//Set up a formula string with all variables exclude "sex"
vars = ". - sex";
//Drop rows with missing values
```

dstat

```
__miss = 1;
{ vnam, mean, var, std, min, max, valid, mis } = dstat
(file, vars);
```

After the above code,

Variable	Mean	Std Dev	Variance	Minimum	Max-
imum	Valid	Missing			

AGE	5.6784	2.9932	8.9593	1.0000	
10.0000	398	2			
PAY	1.9623	0.8006	0.6409	1.0000	
3.0000	398	2			
WT	1.4713	0.3009	0.0906	1.0000	
1.9900	398	2			

Remarks

- 1. If pairwise deletion is used, the minima and maxima will be the true values for the valid data. The means and standard deviations will be computed using the correct number of valid observations for each variable.
 - 2. The supported data set types are [CSV](#), [XLS](#), [XLSX](#), [HDF5](#), [FMT](#), [DAT](#).
- For HDF5 file, the *dataset* must include [file schema](#) and both file name and data set name must be provided, e.g. **dstat** ("h5://C:/gauss17/examples/testdata.h5/mydata", *formula*).

See also

[Formula String](#)

Source

dstat.src

dstatmt

Purpose

Compute descriptive statistics.

Format

```
dout = dstatmt(dataset);  
dout = dstatmt(dataset, vars);  
dout = dstatmt(dataset, vars, ctl);
```

Input

<i>dataset</i>	string, name of data set. If <i>dataset</i> is null or 0, <i>vars</i> will be assumed to be a matrix containing the data.
<i>vars</i>	Optional input, the variables. If <i>dataset</i> contains the name of a data set, <i>vars</i> will be interpreted as: If <i>dataset</i> is null or 0, <i>vars</i> will be interpreted as: Kx1 string array, names of variables. - or - Kx1 numeric vector, indices of variables. - or - formula string , e.g. "PAY + WT" or ". - sex"
<i>ctl</i>	Optional input, instance of a dstatmtControl structure containing the following members:

<i>ctl.altnames</i>	Kx1 string array of alternate variable names to be used if a matrix in memory is analyzed (i.e., <i>dataset</i> is a null string or 0). Default = "".	
<i>ctl.maxbytes</i>	scalar, the maximum number of bytes to be read per iteration of the read loop. Default = 1e9.	
<i>ctl.vartype</i>	Scalar, unused in dstatmt .	
<i>ctl.miss</i>	scalar, default 0.	
	0	there are no missing values (fastest).
	1	listwise deletion, drop a row if any missings occur in it.
	2	pairwise deletion.
<i>ctl.row</i>	scalar, the number of rows to read per iteration of the read loop.	
	If 0, (default) the number of rows will be calculated using <i>ctl.maxbytes</i> and maxvec .	
<i>ctl.output</i>	scalar, controls output, default 1.	
	1	print output table.
	0	do not print output.

These can be any size subset of the variables in the data set and can be in any order. If a scalar 0 is passed, all columns of the data set will be used.

NxK matrix, the data on which to compute the descriptive statistics.

Output

<i>dout</i>	instance of a dstatmtOut structure containing the following members:	
<i>dout.vnames</i>	Kx1	string array, the names of the variables used in the statistics.
<i>dout.mean</i>	Kx1	vector, means.
<i>dout.var</i>	Kx1	vector, variance.
<i>dout.std</i>	Kx1	vector, standard deviation.
<i>dout.min</i>	Kx1	vector, minima.
<i>dout.max</i>	Kx1	vector, maxima.
<i>dout.valid</i>	Kx1	vector, the number of valid cases.
<i>dout.missing</i>	Kx1	vector, the number of missing cases.
<i>dout.errcode</i>	scalar,	error code, 0 if successful; otherwise, one of the following:
	2	Can't open file.
	7	Too many missings - no data left after packing.
	9	<i>altnames</i> member of dstatmtControl structure wrong size.
	10	<i>vartype</i> member of dstatmtControl structure wrong size.

Examples

Example 1: Computing statistics on a GAUSS dataset

dstatmt

```
//Create file name with full path
file_name = getGAUSSHome() $+ "examples/fueleconomy.dat";

//Compute statistics for all variables in the dataset
//The 'call' keyword disregards return values from the func-
tion
call dstatmt(file_name);
```

The above example will print the following report to the program input/output window:

```
-----
-----
Variable              Mean      Std Dev      Variance
Minimum      Maximum      Valid      Missing
-----
-----
annual_fuel_cost      2.5371      0.6533      0.4267
1.0500      5.7000      978      0
engine_displacement    3.2333      1.3757      1.8925
1.0000      8.4000      978      0
```

The code below uses the second input, *vars*, to compute only the descriptive statistics for the second variable.

```
//Create file name with full path
file_name = getGAUSSHome() $+ "examples/fueleconomy.dat";

//Only calculate statistics on the second variable
vars = 2;

//Compute statistics for only the second variable in the
dataset
call dstatmt(file_name, vars);
```


The following report is printed to the program input/output window.

```

-----
-----
Variable              Mean      Std Dev   Variance
Minimum      Maximum  Valid   Missing
-----
engine_displacement  3.2333    1.3757    1.8925
1.0000      8.4000    978      0

```

Example 2: Computing statistics on a csv dataset with formula string

```

//Create file name with full path
file_name = getGAUSSHome() $+ "examples/binary.csv";

//Set up a formula string with variables "gre" and "gpa"
vars = "gre + gpa";

//Compute statistics for all variables in the dataset
//The 'call' keyword disregards return values from the func-
tion
call dstatmt(file_name, vars);

```

The above example will print the following report to the program input/output window:

```

-----
-----
Variable              Mean      Std Dev   Variance   Minimum
      Maximum  Valid   Missing
-----
-----

```

dstatmt

gre	587.7000	115.5165	13344.0702	220.0000
800.0000	400	0		
gpa	3.3899	0.3806	0.1448	2.2600
4.0000	400	0		

Example 3: Using control and out structures

```
//Create file name with full path
file_name = getGAUSSHome() $+ "examples/credit.dat";

//Declare control structure and fill in with defaults
struct dstatmtControl dct1;
dct1 = dstatmtControlCreate();

//Do not print output to the screen
dct1.output = 0;

//Declare output structure
struct dstatmtOut dout;

//Calculate statistics on the 1st, 3rd and 6th variables
vars = { 1, 3, 6 };

//Calculate statistics, and place output in 'dout'
dout = dstatmt(file_name, vars, dct1);

//Print calculated means and variable names
print dout.mean;
print dout.vnames;
```

The code above should print the following output:

```

45.218885
354.94000
13.450000

```

```

Income
Rating
Education

```

Example 4: Computing statistics on a matrix

```

//Set random number seed for repeatable random numbers
rndseed 32452;

//Create a random matrix on which to compute statistics
X = rndn(10,3);

//The empty string as the second input tells GAUSS to
//compute statistics on a matrix rather than a dataset
call dstatmt("", X);

```

The code above will print out the following report:

```

-----
-----
Variable      Mean      Std Dev      Variance      Minimum
Maximum Valid Missing
-----
-----
X1             0.2348      0.8164      0.6664      -1.0736
1.4604        10         0
X2            -0.5062      1.1256      1.2669      -2.2231
1.2695        10         0

```

dstatmt

X3	0.5011	0.7758	0.6018	-0.6119
1.8235	10	0		

Example 5: Computing statistics on a matrix, using structures

```
//Set random number seed for repeatable random numbers
rndseed 32452;

//Declare control structure and fill with default values
struct dstatmtControl dct1;
dct1 = dstatmtControlCreate();

//Variable names for printed output
dct1.altnames = "Alpha"$|"Beta"$|"Gamma";

//Declare structure to hold output values
struct dstatmtOut dout;

//Create a random matrix on which to compute statistics
X = rndn(10,3);

//The empty string as the second input tells GAUSS to
//compute statistics on a matrix rather than a dataset
dout = dstatmt("", X, dct1);
```

This time, the following output will be printed to the screen:

Variable	Mean	Std Dev	Variance	Minimum
Maximum	Valid	Missing		

Alpha	0.2348	0.8164	0.6664	-1.0736
1.4604	10	0		
Beta	-0.5062	1.1256	1.2669	-2.2231
1.2695	10	0		
Gamma	0.5011	0.7758	0.6018	-0.6119
1.8235	10	0		

Remarks

- If pairwise deletion is used, the minima and maxima will be the true values for the valid data. The means and standard deviations will be computed using the correct number of valid observations for each variable.
- For backwards compatibility, the following format is still supported:

```
dout = dstatmt(dctl, dataset, vars);
```

However, all new code should use one of the formats listed at the top of this document.

- The supported data set types are [CSV](#), [XLS](#), [XLSX](#), [HDF5](#), [FMT](#), [DAT](#).
- For HDF5 files, the *dataset* must include a [file schema](#) and both file name and data set name must be provided, e.g. `dstatmt("h5://testdata.h5/mydata")`.

Source

dstatmt.src

See Also

[dstatmtControlCreate](#), [Formula String](#)

dstatmtControlCreate

Purpose

Creates default **dstatmtControl** structure.

dtdate

Include

dstatmt.sdf

Format

```
c = dstatmtControlCreate();
```

Output

c	instance of dstatmtControl structure with members set to default values.
---	---

Example

```
//Declare 'dsm' as an instance of a
//'dstatmtControl' structure
struct dstatmtControl dsm;

//Apply default values to 'dsm'
dsm = dstatmtControlCreate();
```

Source

dstatmt.src

See Also

[dstatmt](#)

dtdate

Purpose

Creates a matrix in DT scalar format.

Format

```
dt = dtdate(year, month, day, hour, minute, second);
```

Input

<i>year</i>	NxK matrix of years.
<i>month</i>	NxK matrix of months, 1-12.
<i>day</i>	NxK matrix of days, 1-31.
<i>hour</i>	NxK matrix of hours, 0-23.
<i>minute</i>	NxK matrix of minutes, 0-59.
<i>second</i>	NxK matrix of seconds, 0-59.

Output

<i>dt</i>	NxK matrix of DT scalar format dates.
-----------	---------------------------------------

Remarks

The arguments must be ExE conformable.

Source

time.src

See Also

[dtday](#), [dttime](#), [utctodt](#), [dttostr](#), [dayofweek](#)

dtday

dtday

Purpose

Creates a matrix in DT scalar format containing only the year, month and day. Time of day information is zeroed out.

Format

```
dt = dtday(year, month, day);
```

Input

<i>year</i>	NxK matrix of years.
<i>month</i>	NxK matrix of months, 1-12.
<i>day</i>	NxK matrix of days, 1-31.

Output

<i>dt</i>	NxK matrix of DT scalar format dates.
-----------	---------------------------------------

Remarks

This amounts to 00:00:00 or midnight on the given day. The arguments must be ExE conformable.

Source

time.src

See Also

[dttime](#), [dtdate](#), [utctodt](#), [dttostr](#)

dttime

Purpose

Creates a matrix in DT scalar format containing only the hour, minute and second. The date information is zeroed out.

Format

```
dt = dttime(hour, minute, second);
```

Input

<i>hour</i>	NxK matrix of hours, 0-23.
<i>minute</i>	NxK matrix of minutes, 0-59.
<i>second</i>	NxK matrix of seconds, 0-59.

Output

<i>dt</i>	NxK matrix of DT scalar format times.
-----------	---------------------------------------

Remarks

The arguments must be ExE conformable.

Source

time.src

See Also

[dtday](#), [dtdate](#), [utctodt](#), [dttostr](#)

dttodtv

dttodtv

Purpose

Converts DT scalar format to DTV vector format.

Format

$dtv = \text{dttodtv}(dt);$

Input

dt	Nx1 vector, DT scalar format.
------	-------------------------------

Output

dtv	Nx8 matrix, DTV vector format.
-------	--------------------------------

Remarks

In DT scalar format, 15:10:55 on July 3, 2005 is 20050703151055.

Each row of dtv , in DTV vector format, contains:

[N,1]	Year
[N,2]	Month in Year, 1-12
[N,3]	Day of month, 1-31
[N,4]	Hours since midnight, 0-23
[N,5]	Minutes, 0-59
[N,6]	Seconds, 0-59
[N,7]	Day of week, 0-6, 0 = Sunday
[N,8]	Days since Jan 1 of current year, 0-365

Example

```
dt = 20100326110722;
print "dt = " dt;
```

```
20100326110722
```

```
dtv = dttodtv(dt);
print "dtv = " dtv;
```

```
2010 3 26 11 7 22 1 84
```

Source

time.src

See Also

[dtvnormal](#), [timeutc](#), [utctodtv](#), [dtvtodt](#), [dttoutc](#), [dtvtodt](#), [strtodt](#), [dttostr](#)

dttostr

Purpose

Converts a matrix containing dates in DT scalar format to a string array.

Format

```
sa = dttostr(x, fmt);
```

Input

<i>x</i>	NxK matrix containing dates in DT scalar format.
<i>fmt</i>	string, or ExE conformable string array containing date/time format characters.

dttostr

Output

<i>sa</i>	NxK string array.
-----------	-------------------

Remarks

The DT scalar format is a double precision representation of the date and time. In the DT scalar format, the number

20120703105031

represents 10:50:31 or 10:50:31 AM on July 3, 2012. **dttostr** converts a date in DT scalar format to a character string using the format string in *fmt*.

The following formats are supported:

YYYY	4 digit year
YR	Last two digits of year
QQ	Quarter of the year. This is calculated from the month number.
MO	Number of month, 01-12
DD	Day of month, 01-31
HH	Hour of day, 00-23
MI	Minute of hour, 00-59
SS	Second of minute, 00-59

A complete DT scalar format number will have 14 digits all to the left of the decimal point. However, **dttostr** will accept numbers with fewer digits. It will assume that the first four digits are the year, the next two the month and so on.

Example

Example 1

```
dt = 201202;  
print dttostr(dt, "QQ-YYYY");
```

produces the output:

```
Q1-2012
```

Example 2

```
s0 = dttostr(utctodt(timeutc), "YYYY-MO-DD HH:MI:SS");  
print ("Date and Time are: " $+ s0);
```

produces the output:

```
Date and time are: 2012-09-14 11:49:10
```

Example 3

```
print  
dttostr(utctodt(timeutc), "Today is DD-MO-YR");
```

produces the output:

```
Today is 14-09-12
```

Example 4

```
x = { 19120317060424, 19370904010928, 19510221031129 };  
s = dttostr(x, "YYYY-MO-DD");
```

produces *s* equal to:

```
1912-03-17  
1937-09-04  
1951-02-21
```

dtoutc

Using the same x from above:

```
s = dttostr(x, "DD/MO/YYYY");
```

produces s equal to:

```
03/17/1912
09/04/1937
02/21/1951
```

Continuing with the same x from above:

```
string fmt = { "YYYY-QQ", "YYYY-QQ-DD", "DD/MO/YYYY" };
s = dttostr(x, fmt);
```

produces s equal to:

```
1912-Q1
1937-Q3-04
21/02/1951
```

See Also

[strtodt](#), [dtoutc](#), [utctodt](#)

dtoutc

Purpose

Converts DT scalar format to UTC scalar format.

Format

```
utc = dtoutc(dt);
```

Input

<i>dt</i>	Nx1 vector, DT scalar format.
-----------	-------------------------------

Output

<i>utc</i>	Nx1 vector, UTC scalar format.
------------	--------------------------------

Remarks

In DT scalar format, 10:50:31 on July 15, 2010 is 20100703105031. A UTC scalar gives the number of seconds since or before January 1, 1970 Greenwich Mean Time.

Example

```
dt = 20010326085118;  
tc = dttoutc(dt);  
  
print "tc = " tc;
```

The above code produces the following output:

```
tc = 985633642;
```

Source

time.src

See Also

[dtvnormal](#), [timeutc](#), [utctodtv](#), [dttodtv](#), [dvtvtdt](#), [dvtvtoutc](#), [dvtvtdt](#), [strtdt](#), [dttostr](#)

dtvnormal

dtvnormal

Purpose

Normalizes a date and time (DTV) vector.

Format

$d = \text{dtvnormal}(t);$

Input

t	1x8 date and time vector that has one or more elements outside the normal range.
-----	--

Output

d	Normalized 1x8 date and time vector.
-----	--------------------------------------

Remarks

The date and time vector is a 1x8 vector whose elements consist of:

Year	Year, four digit integer.
Month	1-12, Month in year.
Day	1-31, Day of month.
Hour	0-23, Hours since midnight.
Min	0-59, Minutes.
Sec	0-59, Seconds.
DoW	0-6, Day of week, 0 = Sunday.
DiY	0-365, Days since Jan 1 of year.

On input missing values are treated as zeros and the last two elements are ignored.

Example

```
format /rd 4,0;

dStart = { 2011 08 21 6 21 37 0 0 };
mnth = { 0 1 0 0 0 0 0 0 };

//Add 6 months to 'dStart' which will give a 14 for the
//month
dEnd = dStart + 6*mnth;

//Normalize the date vector
dEnd2 = dtvnormal(dEnd);
```

After the code above:

dEnd	=	2011	14	21	6	21	37	0	0
dEnd2	=	2012	2	21	6	21	37	2	51

See Also

[date](#), [ethsec](#), [etstr](#), [time](#), [timestr](#), [timeutc](#), [utctodtv](#)

dtvtodt

Purpose

Converts DT vector format to DT scalar format.

Format

```
dt = dtvtodt(dtv);
```

dtvtodt

Input

dtv	Nx8 matrix, DTV vector format.
-------	--------------------------------

Output

dt	Nx1 vector, DT scalar format.
------	-------------------------------

Remarks

In DT scalar format, 11:06:47 on March 15, 2012 is 20120315110647.

Each row of dtv , in DTV vector format, contains:

$[N, 1]$	Year
$[N, 2]$	Month in Year, 1-12
$[N, 3]$	Day of month, 1-31
$[N, 4]$	Hours since midnight, 0-23
$[N, 5]$	Minutes, 0-59
$[N, 6]$	Seconds, 0-59
$[N, 7]$	Day of week, 0-6, 0 = Sunday
$[N, 8]$	Days since Jan 1 of current year, 0-365

Example

```
let dtv = { 2012 9 16 11 7 22 1 84 };
dt = dtvtodt(dtv);
```

The code above assigns dt as follows:

```
20120916110722
```

Source

time.src

See Also

[dtvnormal](#), [timeutc](#), [utctodtv](#), [dttodtv](#), [dttoutc](#), [strtodt](#), [dttostr](#)

dtvtoutc

Purpose

Converts DTV vector format to UTC scalar format.

Format

`utc = dtvtoutc(dtv);`

Input

<i>dtv</i>	Nx8 matrix, DTV vector format.
------------	--------------------------------

Output

<i>utc</i>	Nx1 vector, UTC scalar format.
------------	--------------------------------

Remarks

A UTC scalar gives the number of seconds since or before January 1, 1970 Greenwich Mean Time.

Each row of *dtv*, in DTV vector format, contains:

$[N, 1]$	Year
$[N, 2]$	Month in Year, 1-12

dummy

$[N, 3]$	Day of month, 1-31
$[N, 4]$	Hours since midnight, 0-23
$[N, 5]$	Minutes, 0-59
$[N, 6]$	Seconds, 0-59
$[N, 7]$	Day of week, 0-6, 0 = Sunday
$[N, 8]$	Days since Jan 1 of current year, 0-365

Example

```
dtv = utctodtv(timeutc);  
utc = dtvtoutc(dtv);
```

```
dtv = 2012      7      17      10      13      48      2      198  
utc = 1342545228
```

See Also

[dtvnormal](#), [timeutc](#), [utctodt](#), [dttodtv](#), [dttoutc](#), [dtvtodt](#), [dtvtoutc](#), [strtodt](#), [dttostr](#)

dummy

Purpose

Creates a set of dummy (0/1) variables by breaking up a variable into specified categories. The highest (rightmost) category is unbounded on the right.

Format

```
y = dummy(x, v);
```

Input

x	$N \times 1$ vector of data that is to be broken up into dummy variables.
v	$(K-1) \times 1$ vector specifying the $K-1$ breakpoints (these must be in ascending order) that determine the K categories to be used. These categories should not overlap.

Output

y	$N \times K$ matrix containing the K dummy variables.
-----	---

Remarks

Missings are deleted before the dummy variables are created.

All categories are open on the left (i.e., do not contain their left boundaries) and all but the highest are closed on the right (i.e., do contain their right boundaries). The highest (rightmost) category is unbounded on the right. Thus, only $K-1$ breakpoints are required to specify K dummy variables.

The function **dummybr** is similar to **dummy**, but in that function the highest category is bounded on the right. The function **dummydn** is also similar to **dummy**, but in that function a specified column of dummies is dropped.

Example

```
//Set seed for repeatable random numbers
rndseed 135345;

//Create uniform random integers between 1 and 9
x = ceil(9*rndu(5,1));
```

dummybr

```
//Set the breakpoints  
v = { 1, 5, 7 };  
  
dm = dummy(x,v);
```

The code above produces four dummies based upon the breakpoints in the vector `v`:

```
      x < 1  
1 < x < 5  
5 < x < 7  
7 < x
```

which look like:

```
      0 1 0 0      2  
      0 0 0 1      9  
dm = 0 1 0 0      x = 4  
      0 0 1 0      7  
      1 0 0 0      1
```

Source

`datatran.src`

See Also

[dummybr](#), [dummydn](#), [code](#), [recode](#), [reclassifyCuts](#), [substute](#), [rescale](#), [reclassify](#)

dummybr

Purpose

Creates a set of dummy (0/1) variables. The highest (rightmost) category is bounded on the right.

Format

```
y = dummybr(x, v);
```

Input

x	Nx1 vector of data that is to be broken up into dummy variables.
v	Kx1 vector specifying the K breakpoints (these must be in ascending order) that determine the K categories to be used. These categories should not overlap.

Output

y	NxK matrix containing the K dummy variables. Each row will have a maximum of one 1.
---	---

Remarks

Missings are deleted before the dummy variables are created.

All categories are open on the left (i.e., do not contain their left boundaries) and are closed on the right (i.e., do contain their right boundaries). Thus, K breakpoints are required to specify K dummy variables.

The function **dummy** is similar to **dummybr**, but in that function the highest category is unbounded on the right.

Example

```
//Set seed for repeatable random numbers  
rndseed 135345;  
  
//Create uniform random integers between 1 and 9
```

dummydn

```
x = ceil (9*randu (5,1)) ;

//Set the breakpoints
v = { 1, 5, 7 } ;

dm = dummybr (x,v) ;
```

The code above produces three dummies based upon the breakpoints in the vector *v*:

```
      x < 1
1 < x < 5
5 < x < 7
```

which look like:

```
      0 1 0      2
      0 0 0      9
dm = 0 1 0    x = 4
      0 0 1      7
      1 0 0      1
```

Source

datatran.src

See Also

[dummydn](#), [dummy](#), [code](#), [recode](#), [reclassifyCuts](#), [substute](#), [rescale](#), [reclassify](#)

dummydn

Purpose

Creates a set of dummy (0/1) variables by breaking up a variable into specified categories. The highest (rightmost) category is unbounded on the right, and a specified column of dummies is dropped.

Format

```
 $y = \text{dummydn}(x, \ v, \ p);$ 
```

Input

x	$N \times 1$ vector of data to be broken up into dummy variables.
v	$(K-1) \times 1$ vector specifying the $K-1$ breakpoints (these must be in ascending order) that determine the K categories to be used. These categories should not overlap.
p	positive integer in the range $[1, K]$, specifying which column should be dropped in the matrix of dummy variables.

Output

y	$N \times (K-1)$ matrix containing the $K-1$ dummy variables.
-----	---

Remarks

This is just like the function **dummy**, except that the p th column of the matrix of dummies is dropped. This ensures that the columns of the matrix of dummies do not sum to 1, and so these variables will not be collinear with a vector of ones.

Missings are deleted before the dummy variables are created.

All categories are open on the left (i.e., do not contain their left boundaries) and all but the highest are closed on the right (i.e., do contain their right boundaries). The highest (rightmost) category is unbounded on the right. Thus, only $K-1$ breakpoints are required to specify K dummy variables.

dummydn

Example

```
//Set seed for repeatable random numbers
rndseed 135345;

//Create uniform random integers between 1 and 9
x = ceil(9*rndu(5,1));

//Set the breakpoints
v = { 1, 5, 7 };

//Column to drop
p = 2;

dm = dummydn(x,v,p);
```

The code above produces four dummies based upon the breakpoints in the vector *v*:

```
      x < 1
1 < x < 5
5 < x < 7
7 < x
```

and then remove the *p*th column which will result in:

```
      0 0 0      2
      0 0 1      9
dm = 0 0 0      x = 4
      0 1 0      7
      1 0 0      1
```

Source

datatran.src

See Also

[dummy](#), [dummybr](#), [code](#), [recode](#), [reclassifyCuts](#), [substute](#), [rescale](#), [reclassify](#)

e

ed

Purpose

Accesses an alternate editor.

Format

```
ed filename;
```

Input

<i>filename</i>	literal, the name of the file to be edited.
-----------------	---

Remarks

The default name of the editor is set in `gauss.cfg`. To change the name of the editor used from within a **GAUSS** session, enter:

```
ed = editor_name flags;
```

or

```
ed = "editor_nameflags";
```

The flags are any command line flags you may want between the name of the editor and the filename when your editor is invoked. The quoted version will prevent the flags, if any, from being forced to uppercase.

edit

This command can be placed in the startup file, so it will be set for you automatically when you start **GAUSS**.

See the `edit` command to open a file in the **GAUSS** editor from the command line.

edit

Purpose

Edits a disk file.

Format

```
edit filename;
```

Input

filename literal, the name of the file to be edited.

This command loads a disk file in a **GAUSS** edit window. It is available only in the **GAUSS** graphical user interface.

Remarks

The edit command does not follow the *src_path* to locate files. You must specify the location in the *filename*. The default location is the current directory.

To edit the last run file, use F7 or the Action List toolbar.

Example

```
edit test1.e;
```

See Also

[run](#)

erfInv,erfcInv

Purpose

Computes the inverse of the Gaussian error function (**erfInv**) and its complement (**erfcInv**).

Format

```
x = erfInv(y);  
x = erfcInv(y);
```

Input

y scalar or NxK matrix. $-1 < y < 1$.

Output

x scalar or NxK matrix.

Example

```
x = seqa(.1,.1,10);  
y = erf(x);
```

0.1000	0.1125
0.2000	0.2227
0.3000	0.3286
0.4000	0.4284

eig

```
x = 0.5000    y = 0.5205
    0.6000    0.6039
    0.7000    0.6778
    0.8000    0.7421
    0.9000    0.7969
    1.0000    0.8427
```

```
print erfInv(y);
```

```
0.1000
0.2000
0.3000
0.4000
0.5000
0.6000
0.7000
0.8000
0.9000
1.0000
```

See Also

[erf](#), [erfc](#), [cdfN](#), [cdfNC](#), [cdfNi](#)

eig

Purpose

Computes the eigenvalues of a general matrix.

Format

```
lambda = eig(A);
```

Input

A	NxN matrix or K-dimensional array where the last two dimensions are NxN.
-----	--

Output

λ	Nx1 vector or K-dimensional array where the last two dimensions are Nx1, the eigenvalues of A .
-----------	---

Example

```
A = { 0.5  1.2  0.3,
      0.6  0.9  0.2,
      0.8  1.5  0.0 };

lambda = eig(A);
```

After the above code, λ will equal:

```
1.8626
-0.1871
-0.2754
```

To calculate eigenvalues and eigenvectors see **eigv**. To calculate generalized eigenvalues and eigenvectors, see **lapgeig**, or **lapgeigv**.

Remarks

If A is an array, the result will be an array containing the eigenvalues of each 2-dimensional array described by the two trailing dimensions of A . In other words, for a 10x4x4 array, the result will be a 10x4x1 array containing the eigenvalues of each of the 10 4x4 arrays contained in A .

Errors

eigh

If the eigenvalues cannot all be determined, `lambda[1]` is set to an error code. Passing `lambda[1]` to the **scalerr** function will return the index of the eigenvalue that failed. The eigenvalues for indices **scalerr**(`lambda[1]`)+1 to N should be correct.

Error handling is controlled with the low bit of the trap flag.

trap 0	set <code>lambda[1]</code> and terminate with message
trap 1	set <code>lambda[1]</code> and continue execution

Invalid inputs, such as an infinity, missing value or Nan will cause an error. If the **trap** is set to 1, `lambda` will be set to a scalar error code and program execution will continue. Passing this scalar error code to the **scalerr** function will return -1.

Eigenvalue ordering

The eigenvalues are unordered except that complex conjugate pairs of eigenvalues will appear consecutively with the eigenvalue having the positive imaginary part first.

See Also

[eigh](#), [eighv](#), [eigv](#)

eigh

Purpose

Computes the eigenvalues of a complex hermitian or real symmetric matrix.

Format

```
va = eigh(x);
```

Input

<code>x</code>	NxN matrix or K-dimensional array where the last two
----------------	--

dimensions are $N \times N$.

Output

va

$N \times 1$ vector or K -dimensional array where the last two dimensions are $N \times 1$, the eigenvalues of x .

Remarks

If x is an array, the result will be an array containing the eigenvalues of each 2-dimensional array described by the two trailing dimensions of x . In other words, for a $10 \times 4 \times 4$ array, the result will be a $10 \times 4 \times 1$ array containing the eigenvalues of each of the 10 4×4 arrays contained in x .

Errors

If the eigenvalues cannot all be determined, $va[1]$ is set to an error code. Passing $va[1]$ to the **scalerr** function will return the index of the eigenvalue that failed. The eigenvalues for indices 1 to **scalerr**($va[1]$)-1 should be correct.

Error handling is controlled with the low bit of the trap flag.

trap 0	set $va[1]$ and terminate with message
trap 1	set $va[1]$ and continue execution

Invalid inputs, such as an infinity, missing value or Nan will cause an error. If the **trap** is set to 1, va will be set to a scalar error code and program execution will continue. Passing this scalar error code to the **scalerr** function will return -1.

Eigenvalue ordering

The eigenvalues are in ascending order.

The eigenvalues of a complex hermitian or real symmetric matrix are always real.

See Also

[eig](#), [eighv](#), [eigv](#)

eighv

eighv

Purpose

Computes eigenvalues and eigenvectors of a complex hermitian or real symmetric matrix.

Format

$\{ va, ve \} = \text{eighv}(x);$

Input

x	NxN matrix or K-dimensional array where the last two dimensions are NxN.
-----	--

Output

va	Nx1 vector or K-dimensional array where the last two dimensions are Nx1, the eigenvalues of x .
ve	NxN matrix or K-dimensional array where the last two dimensions are NxN, the eigenvectors of x .

Remarks

If x is an array, va will be an array containing the eigenvalues of each 2-dimensional array described by the two trailing dimensions of x , and ve will be an array containing the corresponding eigenvectors. In other words, for a 10x4x4 array, va will be a 10x4x1 array containing the eigenvalues and ve a 10x4x4 array containing the eigenvectors of each of the 10 4x4 arrays contained in x .

Errors

If the eigenvalues cannot all be determined, `va[1]` is set to an error code. Passing `va[1]` to the **scalerr** function will return the index of the eigenvalue that failed. The eigenvalues for indices 1 to **scalerr**(`va[1]`)-1 should be correct. The eigenvectors are not computed.

Error handling is controlled with the low bit of the trap flag.

trap 0	set <code>va[1]</code> and terminate with message
trap 1	set <code>va[1]</code> and continue execution

Invalid inputs, such as an infinity, missing value or Nan will cause an error. If the **trap** is set to 1, `va` will be set to a scalar error code and program execution will continue. Passing this scalar error code to the **scalerr** function will return -1.

Eigenvalue ordering

The eigenvalues are in ascending order. The columns of `ve` contain the eigenvectors of `x` in the same order as the eigenvalues. The eigenvectors are orthonormal.

The eigenvalues of a complex hermitian or real symmetric matrix are always real.

See Also

[eig](#), [eigh](#), [eigv](#)

eigv

Purpose

Computes eigenvalues and eigenvectors of a general matrix.

Format

```
{ lambda, v } = eigv(x);
```

eigv

Input

x	NxN matrix or K-dimensional array where the last two dimensions are NxN.
-----	--

Output

λ	Nx1 vector or K-dimensional array where the last two dimensions are Nx1, the eigenvalues of x .
v	NxN matrix or K-dimensional array where the last two dimensions are NxN, the eigenvectors of x .

Example

```

x = { 0.5  1.2  0.3,
      0.6  0.9  0.2,
      0.8  1.5  0.0 };

{ lambda, v } = eigv(x);

```

After the above code:

	1.8626		0.5044	-0.7122	-0.6152
lambda =	-0.1871	v =	0.4317	0.3520	0.1361
	-0.2754		0.5643	0.2234	1.0458

Remarks

If x is an array, λ will be an array containing the eigenvalues of each 2-dimensional array described by the two trailing dimensions of x , and v will be an array containing the corresponding eigenvectors. In other words, for a 10x4x4 array, λ

will be a 10x4x1 array containing the eigenvalues and v a 10x4x4 array containing the eigenvectors of each of the 10 4x4 arrays contained in x .

Errors

If the eigenvalues cannot all be determined, $lambda[1]$ is set to an error code. Passing $lambda[1]$ to the **scalerr** function will return the index of the eigenvalue that failed. The eigenvalues for indices **scalerr**($lambda[1]$)+1 to N should be correct. The eigenvectors are not computed.

Error handling is controlled with the low bit of the trap flag.

trap 0	set $lambda[1]$ and terminate with message
trap 1	set $lambda[1]$ and continue execution

Invalid inputs, such as an infinity, missing value or Nan will cause an error. If the **trap** is set to 1, $lambda$ will be set to a scalar error code and program execution will continue. Passing this scalar error code to the **scalerr** function will return -1.

Eigenvalue ordering

The eigenvalues are unordered except that complex conjugate pairs of eigenvalues will appear consecutively with the eigenvalue having the positive imaginary part first. The columns of v contain the eigenvectors of x in the same order as the eigenvalues. The eigenvectors are not normalized.

See Also

[eig](#), [eigh](#), [eighv](#)

elapsedTradingDays

Purpose

Computes number of trading days between two dates inclusively.

elapsedTradingDays

Format

```
n = elapsedTradingDays(a, b);
```

Input

<i>a</i>	scalar, date in DT scalar format.
<i>b</i>	scalar, date in DT scalar format.

Output

<i>n</i>	number of trading days between dates inclusively, that is, elapsed time includes the dates <i>a</i> and <i>b</i> .
----------	--

Remarks

A trading day is a weekday that is not a holiday as defined by the New York Stock Exchange from 1888 through 2013. Holidays are defined in `holidays.asc`. You may edit that file to modify or add holidays.

Example

```
//September 10, 2015
tStart = 20150910110231;

//September 28, 2015
tEnd = 20150928080722;

nDays = elapsedTradingDays(tStart, tEnd);

nDays = 12
```

Source

`finutils.src`

Globals

_fin_holidays

See Also

[getNextTradingDay](#), [getPreviousTradingDay](#), [getNextWeekDay](#), [getPreviousWeekDay](#)

end

Purpose

Terminates a program.

Format

```
end;
```

Remarks

`end` causes **GAUSS** to revert to interactive mode, and closes all open files. `end` also closes the auxiliary output file and turns the window on. It is not necessary to put an `end` statement at the end of a program.

An `end` command can be placed above a label which begins a subroutine to make sure that a program does not enter a subroutine without a `gosub`.

`stop` also terminates a program but closes no files and leaves the window setting as it is.

Example

```
output on;  
screen off;  
print x;  
end;
```

endp

In this example, a matrix x is printed to the auxiliary output. The output to the window is turned off to speed up the printing. The `end` statement is used to terminate the program, so the output file will be closed and the window turned back on.

See Also

[new](#), [stop](#), [system](#)

endp

Purpose

Closes a procedure or keyword definition.

Format

```
endp;
```

Remarks

`endp` marks the end of a procedure definition that began with a `proc` or `keyword` statement. (For details on writing and using procedures, see **PROCEDURES AND KEYWORDS**, CHAPTER 1.)

Example

```
proc regress (y,x) ;
    retp (inv (x'x) *x'y) ;
endp;

x = { 1 3 2, 7 4 9, 1 1 6, 3 3 2 };
```



```
y = { 3, 5, 2, 7 };  
  
b = regress(y,x);
```

After executing the above code:

```
0.1546  
b = 1.5028  
-0.1284
```

See Also

[proc](#), [keyword](#), [retp](#)

endwind

Purpose

Ends graphic panel manipulation; displays graphs with **rerun**. Note: This function is for use with the deprecated PQG graphics.

Library

pgraph

Format

```
endwind;
```

Remarks

This function uses **rerun** to display the most recently created `.tkf` file.

Source

pwindow.src

envget

See Also

[begwind](#), [window](#), [makewind](#), [setwind](#), [nextwind](#), [getwind](#)

envget

Purpose

Searches the environment table for a defined name.

Format

```
y = envget(s);
```

Input

<i>s</i>	string, the name to be searched for.
----------	--------------------------------------

Output

<i>y</i>	string, the string that corresponds to that name in the environment table or a null string if it is not found.
----------	--

Example

Example 1

```
//%USERPROFILE% is the user's home
//directory on most Windows systems
hm_dir = envget("USERPROFILE");
```

Example 2

Below is an example of a procedure that will open a data file using a path stored in an environment string called DPATH.

```
proc dopen(file);
  local fname,fp;
  fname = envget("DPATH");
  //Check to see if DPATH is set or empty
  if fname == "";
    fname = file;
  else;
    //Check to see if 'fname' ends with
    //a path separator
    if strsect(fname,strlen(fname),1) == "\\ ";
      fname = fname $+ file;
    else;
      fname = fname $+ "\\ " $+ file;
    endif;
  endif;
  open fp = ^fname;
  retp(fp);
endp;
```

The procedure returns the file handle and is called as follows:

```
fp = dopen("myfile");
```

See Also

[cdir](#)

eof

Purpose

Tests if the end of a file has been reached.

eof

Format

```
y = eof(fh);
```

Input

<i>fh</i>	scalar, file handle.
-----------	----------------------

Output

<i>y</i>	scalar, 1 if end of file has been reached, else 0.
----------	--

Remarks

This function is used with **readr** and the **fgets** `xxx` commands to test for the end of a file.

The **seekr** function can be used to set the pointer to a specific row position in a data set; the **fseek** function can be used to set the pointer to a specific byte offset in a file opened with **fopen**.

Example

```
open f1 = dat1;
xx = 0;
do until eof(f1);
    xx = xx + moment(readr(f1,100),0);
endo;
```

In this example, the data file `dat1.dat` is opened and given the handle `f1`. Then the data are read from this data set and are used to create the moment matrix ($x'x$) of the data. On each iteration of the loop, 100 additional rows of data are read in, and the moment matrix for this set of rows is computed and added to the matrix `xx`. When all the data have been read, `xx` will contain the entire moment matrix for the data set.

GAUSS will keep reading until **eof**(*f1*) returns the value 1, which it will when the end of the data set has been reached. On the last iteration of the loop, all remaining observations are read in if there are 100 or fewer left.

See Also

[open](#), [readr](#), [seekr](#)

eqSolve

Purpose

Solves a system of nonlinear equations.

Format

```
{ x, retcode } = eqSolve(&F, start);
```

Input

<i>&F</i>	scalar, a pointer to a procedure which computes the value at <i>x</i> of the equations to be solved.
<i>start</i>	Kx1 vector, starting values.

Global Input

The following are set by **eqSolveSet**:

<code>_eqs_JacobianProc</code>	pointer to a procedure which computes the analytical Jacobian. By default, eqSolve will compute the Jacobian numerically.
<code>_eqs_MaxIters</code>	scalar, the maximum number of iterations. Default = 100.

eqSolve

<code>__eqs_StepTol</code>	scalar, the step tolerance. Default = <code>__macheps</code> ^{2/3} .
<code>__eqs_TypicalF</code>	Kx1 vector of the typical $\mathbf{F}(x)$ values at a point not near a root, used for scaling. This becomes important when the magnitudes of the components of $\mathbf{F}(x)$ are expected to be very different. By default, function values are not scaled.
<code>__eqs_TypicalX</code>	Kx1 vector of the typical magnitude of x , used for scaling. This becomes important when the magnitudes of the components of x are expected to be very different. By default, variable values are not scaled.
<code>__eqs_IterInfo</code>	scalar, if nonzero, iteration information is printed. Default = 0.

The following are set by **gausset**:

<code>__Tol</code>	scalar, the tolerance of the scalar function $f = 0.5 * F(x) ^2$ required to terminate the algorithm. Default = 1e-5.
<code>__altnam</code>	Kx1 character vector of alternate names to be used by the printed output. By default, the names "X1, X2,X3..." or "X01,X02,X03..." (depending on how <code>__vpad</code> is set) will be used.
<code>__output</code>	scalar. If non-zero, final results are printed.
<code>__title</code>	string, a custom title to be printed at the top of the iterations report. By default, only a generic title will be printed.
<code>__vpad</code>	scalar. If <code>__altnam</code> is not set, variable names are automatically created. Two types of names can be created: <div style="margin-left: 40px;">0 Variable names are not padded to give them equal length. For example, $x_1, x_2, \dots, x_{10}, \dots$</div>

- 1 Variable names are padded with zeros to give them an equal number of characters. For example, $x01, x02, \dots, x10, \dots$. This is useful if you want the variable names to sort properly.

Output

x	Kx1 vector, solution.	
<i>retcode</i>	scalar, the return code:	
	1	Norm of the scaled function value is less than <code>__Tol</code> . x given is an approximate root of $\mathbf{F}(x)$ (unless <code>__Tol</code> is too large).
	2	The scaled distance between the last two steps is less than the step-tolerance (<code>_eqs_StepTol</code>). x may be an approximate root of $\mathbf{F}(x)$, but it is also possible that the algorithm is making very slow progress and is not near a root, or the step-tolerance is too large.
	3	The last global step failed to decrease <code>norm2(F(x))</code> sufficiently; either x is close to a root of $\mathbf{F}(x)$ and no more

eqSolve

4

5

6

accuracy is possible, or an incorrectly coded analytic Jacobian is being used, or the secant approximation to the Jacobian is inaccurate, or the step-tolerance is too large.

Iteration limit exceeded.

Five consecutive steps of maximum step length have been taken; either **norm2(F(x))** asymptotes from above to a finite value in some direction or the maximum step length is too small.

x seems to be an approximate local minimizer of **norm2(F(x))** that is not a root of **F(x)**. To find a root of **F(x)**, restart **eqSolve** from a different region.

Remarks

The equation procedure should return a column vector containing the result for each equation. For example:

Equation 1: $x_1^2 + x_2^2 - 2 = 0$
Equation 2: $\exp(x_1 - 1) + x_2^3 - 2 = 0$

```
proc (1) = f(var);  
  local x1,x2,eqns;
```



```

x1 = var[1];
x2 = var[2];
eqns[1] = x1^2 + x2^2 - 2;          /* Equation 1 */
eqns[2] = exp(x1-1) + x2^3 - 2;    /* Equation 2 */
retp(eqns);
endp;

```

Example

```

eqSolveSet();

proc (1) = f(x);
  local f1,f2,f3;
  f1 = 3*x[1]^3 + 2*x[2]^2 + 5*x[3] - 10;
  f2 = -x[1]^3 - 3*x[2]^2 + x[3] + 5;
  f3 = 3*x[1]^3 + 2*x[2]^2 - 4*x[3];
  retp(f1|f2|f3);
endp;

proc (1) = fjc(x);
  local fjc1,fjc2, fjc3;
  fjc1 = 9*x[1]^2 ~ 4*x[2] ~ 5;
  fjc2 = -3*x[1]^2 ~ -6*x[2] ~ 1;
  fjc3 = 9*x[1]^2 ~ 4*x[2] ~ -4;
  retp(fjc1|fjc2|fjc3);
endp;

start = { -1, 12, -1 };

_eqs_JacobianProc = &fjc;

{ x,tcode } = eqSolve(&f,start);

```

produces:

eqSolve

```
=====
EqSolve Version 11.0.5          7/17/2015   5:47 pm
=====

||F(X)|| at final solution:          0.93699762
-----
Termination Code = 1:

Norm of the scaled function value is less than __Tol;
-----

-----
VARIABLE      START      ROOTS      F (ROOTS)
-----
X1            -1.00000    0.54144351  4.4175402e-006
X2            12.00000    1.4085912  -6.6263102e-006
X3            -1.00000    1.1111111  4.4175402e-006
-----
```

Source

eqsolve.src

eqSolve

Purpose

Solves a system of nonlinear equations.

Format

```
out = eqSolve(&fct, par);
out = eqSolve(&fct, par, ...);
out = eqSolve(&fct, par, c);
out = eqSolve(&fct, par, ..., c);
```

Input

<i>&fct</i>	pointer to a procedure that computes the function to be minimized. This procedure must have two input arguments, an instance of a PV structure containing the parameters, and an instance of a DS structure containing data, if any. And, one output argument, a column vector containing the result of each equation.
<i>par</i>	an instance of a PV structure. The <i>par</i> instance is passed to the user-provided procedure pointed to by <i>&fct</i> . <i>par</i> is constructed using the pvPack functions.
<i>...</i>	Optional extra arguments. These arguments are passed untouched to the user-provided objective function, by sqpSolveMT .
<i>c</i>	Optional, an instance of an eqSolvemtControl structure. Normally an instance is initialized by calling eqSolvemtControlCreate and members of this instance can be set to other values by the user. For an instance named <i>c</i> , the members are:
<i>c.jacobianProc</i>	pointer to a procedure which computes the analytical Jacobian. By default, eqSolvemt will compute the Jacobian numerically.
<i>c.maxIters</i>	scalar, the maximum number of iterations. Default = 100.
<i>c.stepTolerance</i>	scalar, the step tolerance. Default = $\text{macheps}^{2/3}$.
<i>c.typicalF</i>	Kx1 vector of the typical fct(x) values at a point not near a root,

c.typicalX

used for scaling. This becomes important when the magnitudes of the components of $\mathbf{fct}(\mathbf{x})$ are expected to be very different. By default, function values are not scaled.

Kx1 vector of the typical magnitude of \mathbf{x} , used for scaling. This becomes important when the magnitudes of the components of \mathbf{x} are expected to be very different. By default, variable values are not scaled.

c.printIters

scalar, if nonzero, iteration information is printed. Default = 0.

c.tolerance

scalar, the tolerance of the scalar function $f = 0.5 * ||\mathbf{fct}(\mathbf{X})||^2$ required to terminate the algorithm. That is, the condition that $||\mathbf{f}(\mathbf{x})|| \leq c.tolerance$ must be met before that algorithm can terminate successfully. Default = 1e-5.

c.altNames

Kx1 string array of alternate names to be used by the printed output. By default, the names "X1,X2,X3..." will be used.

c.title

string, printed as a title in output.

c.output

scalar. If non-zero, final results are

printed.

Output

out an instance of an **eqSolvemtOut** structure. For an instance named *out*, the members are:

<i>out.par</i>	an instance of a PV structure containing the parameter estimates.
<i>out.fct</i>	scalar, function evaluated at <i>x</i>
<i>out.retcode</i>	scalar, return code:
-1	Jacobian is singular.
1	Norm of the scaled function value is less than <i>c.tolerance</i> . <i>x</i> given is an approximate root of <i>fct(x)</i> (unless <i>c.tolerance</i> is too large).
2	The scaled distance between the last two steps is less than the step-tolerance (<i>c.stepTolerance</i>). <i>x</i> may be an approximate root of fct (<i>x</i>), but it is also possible that the algorithm is making very slow progress and is not near a root, or the step-tolerance is too large.
3	The last global step failed to decrease norm2 (<i>fct(x)</i>) sufficiently; either <i>x</i> is close to a

eqSolvemt

- root of **fct**(x) and no more accuracy is possible, or an incorrectly coded analytic Jacobian is being used, or the secant approximation to the Jacobian is inaccurate, or the step-tolerance is too large.
- 4 Iteration limit exceeded.
- 5 Five consecutive steps of maximum step length have been taken; either **norm2**($fct(x)$) asymptotes from above to a finite value in some direction or the maximum step length is too small.
- 6 x seems to be an approximate local minimizer of **norm2**($fct(x)$) that is not a root of **fct**(x). To find a root of **fct**(x), restart **eqSolvemt** from a different region.

Remarks

The equation procedure should return a column vector containing the result for each equation.

Examples

Example 1: Basic usage

Equation 1: $x_1^2 + x_2^2 - 5 = 0$
 Equation 2: $\exp(x_1 - 1) + x_2^3 - 5 = 0$

```
//Declare 'par' to be an instance of a PV vector
struct PV par;

//Create default PV struct and add a parameter
//named 'x1' with a starting value of 1
par = pvPack(pvCreate(), 1, "x1");

//Add a parameter named 'x2' to 'par'
//with a starting value of 1
par = pvPack(par, 1, "x2");

//Solve the system of equations
//and print the output to the screen
call eqSolvemt(&fct, par);

//The definition of the function to be minimized
proc fct(struct PV p);
    local x1, x2, z;
    x1 = pvUnpack(p, "x1");
    x2 = pvUnpack(p, "x2");
    z = (x1^2 + x2^2 - 5) | (exp(x1 - 1) + x2^3 - 5);
    retp(z);
endp;
```

After the code above, a short report will be printed to the program input/output window. Part of the output is displayed below:

eqSolvemt

```

-----
-----
VARIABLE          START          ROOTS          F
(ROOT)
-----
-----

X1                1.00000        1.7146639625
0.00000000001
X2                1.00000        1.4352447511
0.00000000002
-----
-----

```

Example 2: Using control and output structures

Equation 1: $x_1^2 + x_2^2 - 5 = 0$
Equation 2: **exp**($x_1 - 1$) + $x_2^3 - 5 = 0$

```

//Declare control structure and fill with defaults
struct eqSolvemtControl c;
c = eqSolvemtControlCreate();

//Turn on printing of iteration information
c.printIters = 1;

//Assign variable names printed output
c.altNames = "alpha" $| "beta";

//Declare 'par' to be an instance of a PV vector
struct PV par;

//Create default PV struct and add a parameter
//named 'x1' with a starting value of 1
par = pvPack(pvCreate(), 1, "x1");

```



```

//Add a parameter named 'x2' to 'par'
//with a starting value of 1
par = pvPack(par,1, "x2");

//Declare output structure to hold results
struct eqSolvemtOut out;

//Solve the system of equations
out = eqSolvemt(&fct,par,c);

//The definition of the function to be minimized
proc fct(struct PV p);
    local x1, x2, z;
    x1 = pvUnpack(p, "x1");
    x2 = pvUnpack(p, "x2");
    z = (x1^2 + x2^2 - 5) | (exp(x1 - 1) + x2^3 - 5);
    retp(z);
endp;

```

The code above will print out a report similar to the previous example. Notice that the variable names in the report are what we assigned to the *altNames* member of the control structure.

```

-----
-----
VARIABLE          START          ROOTS
      F (ROOTS)
-----
-----

alpha             1.00000          1.7146639625
      0.00000000001
beta              1.00000          1.4352447511
      0.00000000002

```

eqSolveMtControlCreate

The parameter values returned by **eqSolveMT** are located in the *par* member of the **eqsolveMTOut** struct. They can be accessed with **pvGetParVector** or **pvUnpack** like this:

```
//Return the values of 'x1' and 'x2' as a 2x1 vector
x_all = pvGetParVector(out.par);

//Return the value of 'x1'
x1 = pvUnpack(out.par, "x1");

//Return the value of 'x2'
x2 = pvUnpack(out.par, "x2");
```

Source

eqsolveMt.src

See Also

[eqSolveMtControlCreate](#), [eqSolveMtOutCreate](#)

eqSolveMtControlCreate

Purpose

Creates default **eqSolveMtControl** structure.

Format

```
c = eqSolveMtControlCreate();
```

Output

`c` instance of **eqSolvemtControl** structure with members set to default values.

Example

Since structures are strongly typed in **GAUSS**, each structure must be declared before it can be used.

```
//declare 'c' as an
//eqSolvemtControl structure
struct eqSolvemtControl c;

//initialize structure c
c = eqSolvemtControlCreate();
```

The members of an **eqSolvemtControl** structure and default values are described in the manual entry for **eqSolvemt**.

Source

eqsolvemt.src

See Also

[eqSolvemt](#)

eqSolvemtOutCreate

Purpose

Creates default **eqSolvemtOut** structure.

eqSolvemtOutCreate

Include

eqsolvemt.sdf

Format

```
c = eqSolvemtOutCreate();
```

Output

c	instance of eqSolvemtOut structure with members set to default values.
---	---

Example

Since structures are strongly typed in **GAUSS**, each structure must be declared before it can be used.

```
//declare structure
struct eqSolvemtOut c;

//Initialize structure
c = eqSolvemtOutCreate();
```

The members of an **eqSolvemtOut** structure and default values are described in the manual entry for **eqSolvemt**.

Source

eqsolvemt.src

See Also

[eqSolvemt](#)

eqSolveSet

Purpose

Sets global input used by **eqSolve** to default values.

Format

```
eqSolveset;
```

Global Output

<code>__eqs_</code> <i>TypicalX</i>	Set to 0.
<code>__eqs_</code> <i>TypicalF</i>	Set to 0.
<code>__eqs_</code> <i>IterInfo</i>	Set to 0.
<code>__eqs_</code> <i>JacobianProc</i>	Set to 0.
<code>__eqs_</code> <i>MaxIters</i>	Set to 100.
<code>__eqs_StepTol</code>	Set to <code>__macheps</code> ^{2/3}

erf,erfc

Purpose

Computes the Gaussian error function (**erf**) and its complement (**erfc**).

Format

```
y = erf(x);  
y = erfc(x);
```

erf,erfc

Input

x NxK matrix.

Output

y NxK matrix.

Remarks

The **erf** and **erfc** functions are closely related to the Normal distribution:

```
if x > 0
    cdfn(x) = 0.5 * (1 + erf(x / sqrt(2)));

if x ≤ 0
    cdfn(x) = 0.5 * erfc(-x / sqrt(2));
```

Example

```
//Print 3 digits after the decimal point
format /rd 5,3;

x = { .5 .4 .3,
      .6 .8 .3 };
y = erf(x);
yc = erfc(x);

//The '~' operator performs horizontal concatenation
//and causes this print statement to format 'x',
//'y' and 'yc' as if they were one 2x9 matrix rather
//than 3 2x3 matrices
//This does not change the variable values, only
//their appearance for this print statement
print x~y~yc;
```

produces the following output:

```
0.500 0.400 0.300 0.520 0.428 0.329 0.480 0.572 0.671
0.600 0.800 0.300 0.604 0.742 0.329 0.396 0.258 0.671
```

See Also

[cdfN](#), [cdfNc](#)

Technical Notes

erf and **erfc** are computed by summing the appropriate series and continued fractions. They are accurate to about 12 or more digits.

erfcplx,erfccplx

Purpose

Computes the Gaussian error function (**erfcplx**) and its complement (**erfccplx**) for complex inputs.

Format

```
f = erfcplx(z);
f = erfccplx(z);
```

Input

z NxK complex matrix; *z* must be > 0.

Output

f NxK complex matrix.

error

Technical Notes

Accuracy is better than 12 significant digits.

References

1. Abramowitz & Stegun, section 7.1, equations 7.1.9, 7.1.23, and 7.1.29
2. Main author Paul Godfrey
3. Small changes by Peter J. Acklam

error

Purpose

Allows the user to generate a user-defined error code which can be tested quickly with the **scalerr** function.

Format

```
y = error(x);
```

Input

x	scalar, in the range 0-65535.
-----	-------------------------------

Output

y	scalar error code which can be interpreted as an integer with the scalerr function.
-----	--

Remarks

The user may assign any number in the range 0-65535 to denote particular error

conditions. This number may be tested for as an error code by **scalerr**.

The **scalerr** function will return the value of the error code and so is the reverse of **error**. These user-generated error codes work in the same way as the intrinsic **GAUSS** error codes which are generated automatically when **trap 1** is on and certain **GAUSS** functions detect a numerical error such as a singular matrix.

```
error (0) ;
```

is equal to the missing value code.

Example

Example 1: Basic usage

```
//Set 'err_code' to contain a scalar error
//code, holding the value 28
err_code = error (28) ;

//Decode error code
err_num = scalerr (err_code) ;

print err_num;
```

The above code will print out the value:

```
28
```

Example 2

The procedure **syminv**, below, returns error code 99 if the matrix is not symmetric. If **invpd** fails, it returns error code 20. If **inv** fails, it returns error code 50. The original trap state is restored before the procedure returns.

```
proc syminv (x) ;
```

error

```

local oldtrap,y;

//Check to see if 'x' is symmetric
if not x == x';
    retp(error(99));
endif;

//Store current error trap state
oldtrap = trapchk(0xffff);

//Turn on trapping of errors
trap 1;

//Attempt matrix inversion with 'invpd'
y = invpd(x);

//Attempt inversion with 'inv' if
//'invpd' returned an error code
if scalerr(y);
    y = inv(x);
endif;

//Reset trap state
trap oldtrap,0xffff;

retp(y);
endp;

```

See Also

[scalerr](#), [trap](#), [trapchk](#)

errorlog

Purpose

Prints an error message to the window and error log file.

Format

```
errorlog str;
```

Input

str string, the error message to print.

Remarks

This command enables you to do your own error handling in your **GAUSS** programs. To print an error message to the window and error log file along with file name and line number information, use [errorlogat](#).

See Also

[errorlogat](#)

errorlogat

Purpose

Prints an error message to the window and error log file, along with the file name and line number at which the error occurred.

Format

```
errorlogat str;
```

etdays

Input

<i>str</i>	string, the error message to print.
------------	-------------------------------------

Remarks

This command enables you to do your own error handling in your **GAUSS** programs. To print an error message to the window and error log file without file name and line number information, use [errorlog](#).

See Also

[errorlog](#)

etdays

Purpose

Computes the difference between two times, as generated by the **date** command, in days.

Format

```
days = etdays(tstart, tend);
```

Input

<i>tstart</i>	3x1 or 4x1 vector, starting date, in the order: yr, mo, day. (Only the first 3 elements are used.)
<i>tend</i>	3x1 or 4x1 vector, ending date, in the order: yr, mo, day. (Only the first 3 elements are used.) MUST be later than <i>tstart</i> .

Output

<i>days</i>	scalar, elapsed time measured in days.
-------------	--

Remarks

This will work correctly across leap years and centuries. The assumptions are a Gregorian calendar with leap years on the years evenly divisible by 4 and not evenly divisible by 100, unless divisible by 400.

Example

```
let date1 = 2008 1 2;  
let date2 = 2009 9 14;  
d = etdays (date1, date2);
```

After the code above, *d* is equal to:

621

Source

time.src

See Also

[dayinyr](#)

ethsec

Purpose

Computes the difference between two times, as generated by the **date** command, in hundredths of a second.

ethsec

Format

```
hs = ethsec(tstart, tend);
```

Input

<i>tstart</i>	4x1 vector, starting date, in the order: yr, mo, day, hundredths of a second.
<i>tend</i>	4x1 vector, ending date, in the order: yr, mo, day, hundredths of a second. MUST be later date than <i>tstart</i> .

Output

<i>hs</i>	scalar, elapsed time measured in hundredths of a second.
-----------	--

Remarks

This will work correctly across leap years and centuries. The assumptions are a Gregorian calendar with leap years on the years evenly divisible by 4 and not evenly divisible by 100, unless divisible by 400.

Example

```
let date1 = 2008 1 2 0;
let date2 = 2009 9 14 0;
t = ethsec(date1, date2);
```

After the code above, *t* is equal to:

```
5365440000
```

Source

time.src

See Also

[dayinyr](#)

etstr

Purpose

Formats an elapsed time measured in hundredths of a second to a string.

Format

```
str = etstr(tothsecs);
```

Input

<i>tothsecs</i>	scalar, an elapsed time measured in hundredths of a second, as given, for instance, by the ethsec function.
-----------------	--

Output

<i>str</i>	string containing the elapsed time in the form:		
#	#	#	#,## seconds
days	hours	minutes	

Example

```
d1 = { 2012, 1, 2, 0 };
```

EuropeanBinomCall

```
d2 = { 2012, 1, 14, 815642 };
t = ethsec(d1,d2);
str = etstr(t);

print "t    = " t;
print "str = " str;
```

Output:

```
t    = 104495642.000
str = 12 days  2 hours  15 minutes  56.42 seconds
```

Source

time.src

See Also

[ethsec](#)

EuropeanBinomCall

Purpose

Prices European call options using binomial method.

Format

```
c = EuropeanBinomCall(S0, K, r, div, tau, sigma, N);
```

Input

$S0$	scalar, current price.
K	Mx1 vector, strike prices.

<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.
<i>N</i>	number of time segments.

Output

<i>c</i>	Mx1 vector, call premiums.
----------	----------------------------

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach", *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;
sigma = .2493;
t0 = dtday(2001, 1, 30);
t1 = dtday(2001, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2012);
c = EuropeanBinomCall(S0,K,r,0,tau,sigma,60);
print c;
```

produces:

EuropeanBinomCall_Greeks

```
17.1325
14.8599
12.6383
```

Source

finprocs.src

EuropeanBinomCall_Greeks

Purpose

Computes Delta, Gamma, Theta, Vega, and Rho for European call options using binomial method.

Format

```
{ d, g, t, v, rh } = EuropeanBinomCall_Greeks(S0, K, r,
div, tau, sigma, N);
```

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.
<i>N</i>	number of time segments.

Global Input

<code>_fin_thetaType</code>	scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.
<code>_fin_epsilon</code>	scalar, finite difference stepsize. Default = 1e-8.

Output

<code>d</code>	Mx1 vector, delta.
<code>g</code>	Mx1 vector, gamma.
<code>t</code>	Mx1 vector, theta.
<code>v</code>	Mx1 vector, vega.
<code>rh</code>	Mx1 vector, rho.

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach", *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
S0 = 305;
K = 300;
r = .08;
sigma = .25;
tau = .33;
div = 0;
print EuropeanBinomcall_Greeks (S0,K,r,0,tau,sigma,30);
```

produces:

EuropeanBinomCall_ImpVol

```

0.670
0.000
-38.426
65.170
56.677

```

Source

finprocs.src

See Also

[EuropeanBinomCall_ImpVol](#), [EuropeanBinomCall](#), [EuropeanBinomPut Greeks](#), [EuropeanBSCall Greeks](#)

EuropeanBinomCall_ImpVol

Purpose

Computes implied volatilities for European call options using binomial method.

Format

```

sigma = EuropeanBinomCall_ImpVol(c, S0, K, r, div, tau,
N);

```

Input

<i>c</i>	Mx1 vector, call premiums.
<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.

τ	scalar, elapsed time to exercise in annualized days of trading.
N	number of time segments.

Output

σ	Mx1 vector, volatility.
----------	-------------------------

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach", *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
c = { 13.70, 11.90, 9.10 };
S0 = 718.46;
K = { 720, 725, 730 };
r = .0368;
div = 0;
t0 = dtday(2012, 1, 30);
t1 = dtday(2012, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2012);
sigma = EuropeanBinomCall_ImpVol(c,S0,K,r,0,tau,30);
print sigma;
```

produces:

```
0.2027
0.2081
0.1989
```

EuropeanBinomPut

Source

finprocs.src

EuropeanBinomPut

Purpose

Prices European put options using binomial method.

Format

$c = \text{EuropeanBinomPut}(S0, K, r, div, tau, sigma, N);$

Input

$S0$	scalar, current price.
K	Mx1 vector, strike prices.
r	scalar, risk free rate.
div	continuous dividend yield.
tau	scalar, elapsed time to exercise in annualized days of trading.
$sigma$	scalar, volatility.
N	number of time segments.

Output

c	Mx1 vector, put premiums.
-----	---------------------------

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified

approach", *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
S0 = 718.46;
K = { 720, 725, 730 };
r = .0398;
sigma = .2493;
t0 = dtday(2012, 1, 30);
t1 = dtday(2012, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2012);
c = EuropeanBinomPut(S0,K,r,0,tau,sigma,60); print c;
```

produces:

```
16.872213
19.606098
22.390831
```

Source

finprocs.src

EuropeanBinomPut_Greeks

Purpose

Computes Delta, Gamma, Theta, Vega, and Rho for European put options using binomial method.

EuropeanBinomPut_Greeks

Format

```
{ d, g, t, v, rh } = EuropeanBinomPut_Greeks(S0, K, r,
div, tau, sigma, N);
```

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.
<i>N</i>	number of time segments.

Global Input

<i>_fin_thetaType</i>	scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.
<i>_fin_epsilon</i>	scalar, finite difference stepsize. Default = 1e-8.

Output

<i>d</i>	Mx1 vector, delta.
<i>g</i>	Mx1 vector, gamma.
<i>t</i>	Mx1 vector, theta.
<i>v</i>	Mx1 vector, vega.
<i>rh</i>	Mx1 vector, rho.

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach", *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
S0 = 305;  
K = 300;  
r = .08;  
div = 0;  
sigma = .25;  
tau = .33;  
print EuropeanBinomPut_Greeks(S0,K,r,0,tau,sigma,60);
```

produces:

```
-0.350  
0.001  
7.237  
65.432  
-39.652
```

Source

finprocs.src

See Also

[EuropeanBinomPut ImpVol](#), [EuropeanBinomPut](#), [EuropeanBinomCall Greeks](#),
[EuropeanBSPut Greeks](#)

EuropeanBinomPut_ImpVol

EuropeanBinomPut_ImpVol

Purpose

Computes implied volatilities for European put options using binomial method.

Format

$\sigma = \text{EuropeanBinomPut_ImpVol}(c, S_0, K, r, \text{div}, \tau, N);$

Input

c	Mx1 vector, put premiums.
S_0	scalar, current price.
K	Mx1 vector, strike prices.
r	scalar, risk free rate.
div	continuous dividend yield.
τ	scalar, elapsed time to exercise in annualized days of trading.
N	number of time segments.

Output

σ	Mx1 vector, volatility.
----------	-------------------------

Remarks

The binomial method of Cox, Ross, and Rubinstein ("Option pricing: a simplified approach", *Journal of Financial Economics*, 7:229:264) as described in *Options, Futures, and other Derivatives* by John C. Hull is the basis of this procedure.

Example

```
p = { 14.60, 17.10, 20.10 };
S0 = 718.46;
K = { 720, 725, 730 };
r = .0398;
div = 0;
t0 = dtday(2012, 1, 30);
t1 = dtday(2012, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2012);
sigma = EuropeanBinomPut_ImpVol(p,S0,K,r,0,tau,30);
print sigma;
```

produces:

```
0.21609253
0.21139494
0.21407512
```

Source

finprocs.src

EuropeanBSCall

Purpose

Prices European call options using Black, Scholes and Merton method.

Format

```
c = EuropeanBSCall(S0, K, r, div, tau, sigma);
```

EuropeanBSCall

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.

Output

<i>c</i>	Mx1 vector, call premiums.
----------	----------------------------

Example

```
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;
sigma = .2493;
t0 = dtday(2012, 1, 30);
t1 = dtday(2012, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2012);
c = EuropeanBSCall(S0,K,r,0,tau,sigma);
print c;
```

produces:

```
17.1351
14.7955
12.6860
```

Source

finprocs.src

EuropeanBSCall_Greeks

Purpose

Computes Delta, Gamma, Theta, Vega, and Rho for European call options using Black, Scholes, and Merton method.

Format

```
{ d, g, t, v, rh } = EuropeanBSCall_Greeks(S0, K, r, div,
tau, sigma);
```

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.

Global Input

<i>_fin_thetaType</i>	scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.
<i>_fin_epsilon</i>	scalar, finite difference stepsize. Default = 1e-8.

EuropeanBSCall_Greeks

Output

d	Mx1 vector, delta.
g	Mx1 vector, gamma.
t	Mx1 vector, theta.
v	Mx1 vector, vega.
rh	Mx1 vector, rho.

Example

```
S0 = 305;
K = 300;
r = .08;
sigma = .25;
tau = .33;
print EuropeanBSCall_Greeks(S0,K,r,0,tau,sigma);
```

produce:

```
0.6446
0.0085
-38.5054
65.2563
56.8720
```

Source

finprocs.src

See Also

[EuropeanBSCall ImpVol](#), [EuropeanBSCall](#), [EuropeanBSPut Greeks](#),
[EuropeanBinomCall Greeks](#)

EuropeanBSCall_ImpVol

Purpose

Computes implied volatilities for European call options using Black, Scholes, and Merton method.

Format

```
sigma = EuropeanBSCall_ImpVol(c, S0, K, r, div, tau);
```

Input

<i>c</i>	Mx1 vector, call premiums.
<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.

Output

<i>sigma</i>	Mx1 vector, volatility.
--------------	-------------------------

Example

```
c = { 13.70, 11.90, 9.10 };  
S0 = 718.46;  
K = { 720, 725, 730 };  
r = .0498;  
t0 = dtday(2012, 1, 30);
```

EuropeanBSPut

```
t1 = dtday(2012, 2, 16);
tau = elapsedTradingDays(t0,t1) /
      annualTradingDays(2012);
sigma = EuropeanBSCall_ImpVol(c,S0,K,r,0,tau);
print sigma;
```

produces:

```
0.1986
0.2064
0.1951
```

Source

finprocs.src

EuropeanBSPut

Purpose

Prices European put options using Black, Scholes, and Merton method.

Format

$c = \text{EuropeanBSPut}(S0, K, r, div, tau, sigma);$

Input

$S0$	scalar, current price.
K	Mx1 vector, strike prices.
r	scalar, risk free rate.
div	continuous dividend yield.
tau	scalar, elapsed time to exercise in annualized days of

<i>sigma</i>	trading. scalar, volatility.
--------------	---------------------------------

Output

<i>c</i>	Mx1 vector, put premiums.
----------	---------------------------

Example

```
S0 = 718.46;  
K = { 720, 725, 730 };  
r = .0498;  
sigma = .2493;  
t0 = dtday(2012, 1, 30);  
t1 = dtday(2012, 2, 16);  
tau = elapsedTradingDays(t0,t1) /  
      annualTradingDays(2012);  
c = EuropeanBSPut(S0,K,r,0,tau,sigma);  
print c;
```

produces:

```
16.6700  
19.3164  
22.1930
```

Source

finprocs.src

EuropeanBSPut_Greeks

EuropeanBSPut_Greeks

Purpose

Computes Delta, Gamma, Theta, Vega, and Rho for European put options using Black, Scholes, and Merton method.

Format

```
{ d, g, t, v, rh } = EuropeanBSPut_Greeks(S0, K, r, div, tau, sigma);
```

Input

<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.
<i>sigma</i>	scalar, volatility.

Global Input

<i>_fin_thetaType</i>	scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.
<i>_fin_epsilon</i>	scalar, finite difference stepsize. Default = 1e-8.

Output

<i>d</i>	Mx1 vector, delta.
----------	--------------------

g	Mx1 vector, gamma.
t	Mx1 vector, theta.
v	Mx1 vector, vega.
rh	Mx1 vector, rho.

Example

```
S0 = 305;  
K = 300;  
r = .08;  
sigma = .25;  
tau = .33;  
print EuropeanBSPut_Greeks (S0,K,r,0,tau,sigma);
```

produces:

```
-0.3554  
 0.0085  
-15.1307  
 65.2563  
-39.54861
```

Source

finprocs.src

See Also

[EuropeanBSPut ImpVol](#), [EuropeanBSPut](#), [EuropeanBSCall Greeks](#),
[EuropeanBinomPut Greeks](#)

EuropeanBSPut_ImpVol

EuropeanBSPut_ImpVol

Purpose

Computes implied volatilities for European put options using Black, Scholes, and Merton method.

Format

```
sigma = EuropeanBSPut_ImpVol(c, S0, K, r, div, tau);
```

Input

<i>c</i>	Mx1 vector, put premiums
<i>S0</i>	scalar, current price.
<i>K</i>	Mx1 vector, strike prices.
<i>r</i>	scalar, risk free rate.
<i>div</i>	continuous dividend yield.
<i>tau</i>	scalar, elapsed time to exercise in annualized days of trading.

Output

<i>sigma</i>	Mx1 vector, volatility.
--------------	-------------------------

Example

```
p = { 14.60, 17.10, 20.10 };
S0 = 718.46;
K = { 720, 725, 730 };
r = .0498;
t0 = dtday(2012, 1, 30);
```

```
t1 = dtday(2012, 2, 16);  
tau = elapsedTradingDays(t0,t1) / annualTradingDays(2012);  
sigma = EuropeanBSPut_ImpVol(p,S0,K,r,0,tau);  
print sigma;
```

produce:

```
0.2188  
0.2165  
0.2177
```

Source

finprocs.src

exctsmpl

Purpose

Computes a random subsample of a data set.

Format

```
n = exctsmpl(infile, outfile, percent);
```

Input

<i>infile</i>	string, the name of the original data set.
<i>outfile</i>	string, the name of the data set to be created.
<i>percent</i>	scalar, the percentage random sample to take. This must be in the range 0-100.

exctsmpl

Output

n	scalar, number of rows in output data set.
Error returns are controlled by the low bit of the trap flag:	
trap 0	terminate with error message
trap 1	return scalar negative integer
-1	can't open input file
-2	can't open output file
-3	disk full

Remarks

Random sampling is done with replacement. Thus, an observation may be in the resulting sample more than once. If *percent* is 100, the resulting sample will not be identical to the original sample, though it will be the same size.

Example

```
n = exctsmpl (getGAUSSHome () $+
  "examples/freqdata.dat", "rout", 30);
```

freqdata.dat is an example data set provided with **GAUSS**. Switching to the examples subdirectory of your **GAUSS** installation directory will make it possible to do the above example as shown. Otherwise you will need to substitute another data set name for "freqdata.dat".

Source

exctsmpl.src

exec

Purpose

Executes an executable program and returns the exit code to **GAUSS**.

Format

```
y = exec(program, comline);
```

Input

<i>program</i>	string, the name of the program, including the extension, to be executed.
<i>comline</i>	string, the arguments to be placed on the command line of the program being executed.

Output

<i>y</i>	scalar, the exit code returned by <i>program</i> . If exec can't execute <i>program</i> , the error returns will be negative: -1 file not found -2 the file is not an executable file -3 not enough memory -4 command line too long
----------	---

Example

```
y = exec("atog", "comd1.cmd");  
  
//If 'y' is nonzero
```

execbg

```
if y;
    errorlog"atog failed";
end;
endif;
```

In this example the ATOG ASCII conversion utility is executed under the **exec** function. The name of the command file to be used, `comd1.cmd`, is passed to ATOG on its command line. The exit code `y` returned by **exec** is tested to see if ATOG was successful; if not, the program will be terminated after printing an error message. See ATOG, Section 1.

execbg

Purpose

Executes an executable program in the background and returns the process id to GAUSS.

Format

```
pid = execbg(program, comline);
```

Input

<i>program</i>	string, the name of the program, including the extension, to be executed.
<i>comline</i>	string, the arguments to be placed on the command line of the program being executed.

Output

<i>pid</i>	scalar, the process id of the executable returned by program.
------------	---

If **execbg** cannot execute program, the error returns will be negative:

- 1 file not found
- 2 the file is not an executable file
- 3 not enough memory
- 4 command line too long

Example

```
y = execbg ("atog.exe", "comd1.cmd");  
if (y < 0);  
    errorlog"atog failed";  
end;  
endif;
```

In this example, the ATOG ASCII conversion utility is executed under the **execbg** function. The name of the command file to be used, `comd1.cmd`, is passed to ATOG on its command line. The returned value, `y`, is tested to see whether ATOG was successful. If not successful the program terminates after printing an error message. See **ATOG**, CHAPTER 1.

exp

Purpose

Calculates the exponential function.

Format

```
y = exp(x);
```

exp

Input

x	NxK matrix or N-dimensional array.
-----	------------------------------------

Output

y	NxK matrix or N-dimensional array containing e , the base of natural logs, raised to the powers given by the elements of x .
-----	--

Example

```
x = eye(3);  
y = exp(x);
```

```
      1.000000  0.000000  0.000000  
x =  0.000000  1.000000  0.000000  
      0.000000  0.000000  1.000000
```

```
      2.718282  1.000000  1.000000  
y =  1.000000  2.718282  1.000000  
      1.000000  1.000000  2.718282
```

This example creates a 3x3 identity matrix and computes the exponential function for each one of its elements. Note that **exp**(1) returns e , the base of natural logs.

See Also

[ln](#)

extern (dataloop)

Purpose

Allows access to matrices or strings in memory from inside a data loop.

Format

```
extern variable_list;
```

Remarks

Commas in *variable_list* are optional.

extern tells the translator not to generate local code for the listed variables, and not to assume that they are elements of the input data set.

extern statements should be placed before any reference to the symbols listed. The specified names should not exist in the input data set, or be used in a **make** statement.

Example

This example shows how to assign the contents of an external vector to a new variable in the data set, by iteratively assigning a range of elements to the variable. The reserved variable `x_x` contains the data read from the input data set on each iteration. The external vector must have at least as many rows as the data set.

```
base = 1;      /* used to index a range of */
               /* elements from exvec */
dataloop oldata newdata;
extern base, exvec;
make ndvar = exvec[seqa(base,1, rows(x_x))];
# base = base + rows(x_x); /* execute command */
                        /* literally */
endata;
```

external

external

Purpose

Lets the compiler know about symbols that are referenced above or in a separate file from their definitions.

Format

```
external proc  dog,  cat;
external keyword  dog;
external fn  dog;
external matrix x, y,  z;
external string  mstr,  cstr;
external array  a,  b;
external sparse matrix  sma,  smb;
external struct  structure_type  sta,  stb;
```

Remarks

See **PROCEDURES AND KEYWORDS**, CHAPTER 1.

You may have several procedures in different files that reference the same global variable. By placing an `external` statement at the top of each file, you can let the compiler know what the type of the symbol is. If the symbol is listed and strongly typed in an active library, no `external` statement is needed.

If a matrix, string, N-dimensional array, sparse matrix, or structure appears in an `external` statement, it needs to appear once in a `declare` statement. If no declaration is found, an **Undefined symbol** error message will result.

Example

Let us suppose that you created a set of procedures defined in different files, which all set a global matrix `_errcode` to some scalar error code if errors were encountered.

You could use the following code to call one of the procedures in the set and check whether it succeeded:

```
external matrix _errcode;
x = rndn(10,5);
y = myproc1(x);
if _errcode;
    print "myproc1 failed";
end;
endif;
```

Without the `external` statement, the compiler would assume that `_errcode` was a procedure and incorrectly compile this program. The file containing the `myproc1` procedure must also contain an `external` statement that defines `_errcode` as a matrix, but this would not be encountered by the compiler until the `if` statement containing the reference to `_errcode` in the main program file had already been incorrectly compiled.

See Also

[declare](#)

eye

Purpose

Creates an identity matrix.

Format

```
y = eye(n);
```

Input

n	scalar, size of identity matrix to be created.
-----	--

eye

Output

y $n \times n$ identity matrix.

Remarks

If n is not an integer, it will be truncated to an integer.

The matrix created will contain 1's down the diagonal and 0's everywhere else.

Example

```
x = eye(3);
```

The code above assigns x to be equal to:

```
1.0000 0.0000 0.0000
0.0000 1.0000 0.0000
0.0000 0.0000 1.0000
```

See Also

[zeros](#), [ones](#)

f

fcheckerr

Purpose

Gets the error status of a file.

Format

```
err = fcheckerr(f);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
----------	--

Output

<i>err</i>	scalar, error status.
------------	-----------------------

Remarks

If there has been a read or write error on a file, **fcheckerr** returns 1, otherwise 0.

If you pass **fcheckerr** the handle of a file opened with [open](#) (i.e., a data set or matrix file), your program will terminate with a fatal error.

fclearerr

fclearerr

Purpose

Gets the error status of a file, then clears it.

Format

```
err = fclearerr(f);
```

Input

f scalar, file handle of a file opened with **fopen**.

Output

err scalar, error status.

Remarks

Each file has an error flag that gets set when there is an I/O error on the file. Typically, once this flag is set, you can no longer do I/O on the file, even if the error is a recoverable one. **fclearerr** clears the file's error flag, so you can attempt to continue using it.

If there has been a read or write error on a file, **fclearerr** returns 1, otherwise 0.

If you pass **fclearerr** the handle of a file opened with [open](#) (i.e., a data set or matrix file), your program will terminate with a fatal error.

The flag accessed by **fclearerr** is not the same as that accessed by **fstrerror**.

feq,fge,fgt,fle,flt,fne

Purpose

Fuzzy comparison functions. These functions use `__fcmptol` to fuzz the comparison operations to allow for roundoff error.

Format

```
y = feq(a, b);  
y = fge(a, b);  
y = fgt(a, b);  
y = fle(a, b);  
y = flt(a, b);  
y = fne(a, b);
```

Input

<i>a</i>	NxK matrix, first matrix.
<i>b</i>	LxM matrix, second matrix, ExE compatible with <i>a</i> .

Global Input

<code>__fcmptol</code>	scalar, comparison tolerance. The default value is 1.0e-15.
------------------------	---

Output

<i>y</i>	scalar, 1 (TRUE) or 0 (FALSE).
----------	--------------------------------

Remarks

The return value is TRUE if every comparison is TRUE.

feq,fge,fgt,fle,flt,fne

The statement:

```
y = feq(a,b);
```

is equivalent to:

```
y = a eq b;
```

For the sake of efficiency, these functions are not written to handle missing values. If *a* and *b* contain missing values, use **missrv** to convert the missing values to something appropriate before calling a fuzzy comparison function.

The calling program can reset *_fcmp_tol* before calling these procedures:

```
_fcmp_tol = 1e-12;
```

Example

```
_fcmp_tol = 1e-12;

x = rndu(2,2);

y = x + 0.5*(_fcmp_tol);

if fge(x,y);
    print "each element of x is greater than";
    print "or equal to each element of y";
else;
    print "at least one element of x is less";
    print "its corresponding element in y";
endif;
```

Source

fcompare.src

See Also

[dotfeq-dotfne](#)

feqmt,fgemt,fgtmt,flemt,fltmt,fnemt

Purpose

Fuzzy comparison functions. These functions use the *fcmtol* argument to fuzz the comparison operations to allow for roundoff error.

Format

```
y = feqmt(a, b, fcmtol);
y = fgemt(a, b, fcmtol);
y = fgtmt(a, b, fcmtol);
y = flemt(a, b, fcmtol);
y = fltmt(a, b, fcmtol);
y = fnemt(a, b, fcmtol);
```

Input

<i>a</i>	NxK matrix, first matrix.
<i>b</i>	LxM matrix, second matrix, ExE compatible with <i>a</i> .
<i>fcmtol</i>	scalar, comparison tolerance.

Output

<i>y</i>	scalar, 1 (TRUE) or 0 (FALSE).
----------	--------------------------------

Remarks

The return value is TRUE if every comparison is TRUE.

feqmt,fgemt,fgtmt,flemt,fltmt,fnemt

The statement:

```
y = feqmt(a,b,1e-15);
```

is equivalent to:

```
y = a eq b;
```

For the sake of efficiency, these functions are not written to handle missing values. If *a* and *b* contain missing values, use **missrv** to convert the missing values to something appropriate before calling a fuzzy comparison function.

Example

```
tol = 1e-12;

x = rndu(2,2);

y = x + 0.5*(tol);

if fgemt(x,y,tol);
    print "each element of x is greater than";
    print "or equal to each element of y";
else;
    print "at least one element of x is less";
    print "its corresponding element in y";
endif;
```

Source

fcomparemt.src

See Also

[dotfeqmt-dotfnemt](#)

fflush

Purpose

Flushes a file's output buffer.

Format

```
ret = fflush(f);
```

Input

f scalar, file handle of a file opened with **fopen**.

Output

ret scalar, 0 if successful, -1 if not.

Remarks

If **fflush** fails, you can call **fstrerror** to find out why.

If you pass **fflush** the handle of a file opened with **open** (i.e., a data set or matrix file), your program will terminate with a fatal error.

fft

Purpose

Computes a 1- or 2-D Fast Fourier transform.

Format

```
y = fft(x);
```

ffti

Input

x	$N \times K$ matrix.
-----	----------------------

Output

y	$L \times M$ matrix, where L and M are the smallest powers of 2 greater than or equal to N and K , respectively.
-----	--

Remarks

This computes the FFT of x , scaled by $1/N$.

This uses a Temperton Fast Fourier algorithm.

If N or K is not a power of 2, x will be padded out with zeros before computing the transform.

See Also

[ffti](#), [rfft](#), [rffti](#)

ffti

Purpose

Computes an inverse 1- or 2-D Fast Fourier transform.

Format

$y = \mathbf{ffti}(x);$

Input

x	$N \times K$ matrix.
-----	----------------------

Output

y	LxM matrix, where L and M are the smallest prime factor products greater than or equal to N and K, respectively.
-----	--

Remarks

Computes the inverse FFT of x , scaled by $1/N$.

This uses a Temperton prime factor Fast Fourier algorithm.

See Also

[fft](#), [rfft](#), [rffti](#)

fftm

Purpose

Computes a multi-dimensional FFT.

Format

```
 $y = \text{fftm}(x, \text{dim});$ 
```

Input

x	Mx1 vector, data.
dim	Kx1 vector, size of each dimension.

Output

y	Lx1 vector, FFT of x .
-----	--------------------------

Remarks

The multi-dimensional data are laid out in a recursive or heirarchical fashion in the vector x . That is to say, the elements of any given dimension are stored in sequence left to right within the vector, with each element containing a sequence of elements of the next smaller dimension. In abstract terms, a 4-dimensional $2 \times 2 \times 2 \times 2$ hypercubic x would consist of two cubes in sequence, each cube containing two matrices in sequence, each matrix containing two rows in sequence, and each row containing two columns in sequence. Visually, x would look something like this:

$$X_{hyper} = X_{cube1} | X_{cube2}$$

$$X_{cube1} = X_{mat1} | X_{mat2}$$

$$X_{mat1} = X_{row1} | X_{row2}$$

Or, in an extended **GAUSS** notation, x would be:

```
Xhyper = x[1, ., ., .] | x[2, ., ., .];
Xcube1 = x[1, 1, ., .] | x[1, 2, ., .];
Xmat1   = x[1, 1, 1, .] | x[1, 1, 2, .];
Xrow1   = x[1, 1, 1, 1] | x[1, 1, 1, 2];
```

To be explicit, x would be laid out like this:

```
x[1, 1, 1, 1] x[1, 1, 1, 2] x[1, 1, 2, 1] x[1, 1, 2, 2]
x[1, 2, 1, 1] x[1, 2, 1, 2] x[1, 2, 2, 1] x[1, 2, 2, 2]
x[2, 1, 1, 1] x[2, 1, 1, 2] x[2, 1, 2, 1] x[2, 1, 2, 2]
x[2, 2, 1, 1] x[2, 2, 1, 2] x[2, 2, 2, 1] x[2, 2, 2, 2]
```

If you look at the last diagram for the layout of x , you'll notice that each line actually constitutes the elements of an ordinary matrix in normal row-major order. This is easy to achieve with **vecr**. Further, each pair of lines or "matrices" constitutes one of the

desired cubes, again with all the elements in the correct order. And finally, the two cubes combine to form the hypercube. So, the process of construction is simply a sequence of concatenations of column vectors, with a **vecr** step if necessary to get started.

Here's an example, this time working with a 2x3x2x3 hypercube.

```
let dim = 2 3 2 3;
let x1[2,3] = 1 2 3 4 5 6;
let x2[2,3] = 6 5 4 3 2 1;
let x3[2,3] = 1 2 3 5 7 11;
xc1 = vecr(x1) | vecr(x2) | vecr(x3); /* cube 1 */
let x1 = 1 1 2 3 5 8;
let x2 = 1 2 6 24 120 720;
let x3 = 13 17 19 23 29 31;
xc2 = x1|x2|x3;                      /* cube 2 */

xh = xc1|xc2;                        /* hypercube */
xhfft = fftm(xh,dim);

let dimi = 2 4 2 4;
xhffti = fftmi(xhfft,dimi);
```

We left out the **vecr** step for the 2nd cube. It's not really necessary when you're constructing the matrices with **let** statements.

dim contains the dimensions of *x*, beginning with the highest dimension. The last element of *dim* is the number of columns, the next to the last element of *dim* is the number of rows, and so on. Thus

```
dim = { 2, 3, 3 };
```

indicates that the data in *x* is a 2x3x3 three-dimensional array, i.e., two 3x3 matrices of data. Suppose that *x1* is the first 3x3 matrix and *x2* the second 3x3 matrix, then:

fftm

```
x = vecr (x1) | vecr (x2)
```

The size of *dim* tells you how many dimensions *x* has.

The arrays have to be padded in each dimension to the nearest power of two. Thus the output array can be larger than the input array. In the 2x3x2x3 hypercube example, *x* would be padded from 2x3x2x3 out to 2x4x2x4. The input vector would contain 36 elements, while the output vector would contain 64 elements. You may have noticed that we used a *dim* with padded values at the end of the example to check our answer.

Source

fftm.src

See Also

[fftm](#), [fft](#), [ffti](#), [fftn](#)

fftm

Purpose

Computes a multi-dimensional inverse FFT.

Format

```
y = fftm(x, dim);
```

Input

<i>x</i>	Mx1 vector, data.
<i>dim</i>	Kx1 vector, size of each dimension.

Output

 y Lx1 vector, inverse FFT of x .

Remarks

The multi-dimensional data are laid out in a recursive or heirarchical fashion in the vector x . That is to say, the elements of any given dimension are stored in sequence left to right within the vector, with each element containing a sequence of elements of the next smaller dimension. In abstract terms, a 4-dimensional 2x2x2x2 hypercubic x would consist of two cubes in sequence, each cube containing two matrices in sequence, each matrix containing two rows in sequence, and each row containing two columns in sequence. Visually, x would look something like this:

$$X_{hyper} = X_{cube1} | X_{cube2}$$

$$X_{cube1} = X_{mat1} | X_{mat2}$$

$$X_{mat1} = X_{row1} | X_{row2}$$

Or, in an extended **GAUSS** notation, x would be:

```
Xhyper = x[1, ., ., .] | x[2, ., ., .];
Xcube1 = x[1, 1, ., .] | x[1, 2, ., .];
Xmat1 = x[1, 1, 1, .] | x[1, 1, 2, .];
Xrow1 = x[1, 1, 1, 1] | x[1, 1, 1, 2];
```

To be explicit, x would be laid out like this:

```
x[1, 1, 1, 1] x[1, 1, 1, 2] x[1, 1, 2, 1] x[1, 1, 2, 2]
x[1, 2, 1, 1] x[1, 2, 1, 2] x[1, 2, 2, 1] x[1, 2, 2, 2]
x[2, 1, 1, 1] x[2, 1, 1, 2] x[2, 1, 2, 1] x[2, 1, 2, 2]
x[2, 2, 1, 1] x[2, 2, 1, 2] x[2, 2, 2, 1] x[2, 2, 2, 2]
```

If you look at the last diagram for the layout of x , you'll notice that each line actually constitutes the elements of an ordinary matrix in normal row-major order. This is easy to achieve with **vecr**. Further, each pair of lines or "matrices" constitutes one of the desired cubes, again with all the elements in the correct order. And finally, the two cubes combine to form the hypercube. So, the process of construction is simply a sequence of concatenations of column vectors, with a **vecr** step if necessary to get started.

Here's an example, this time working with a 2x3x2x3 hypercube.

```
let dim = 2 3 2 3;
let x1[2,3] = 1 2 3 4 5 6;
let x2[2,3] = 6 5 4 3 2 1;
let x3[2,3] = 1 2 3 5 7 11;
xc1 = vecr(x1) | vecr(x2) | vecr(x3); /* cube 1 */
let x1 = 1 1 2 3 5 8;
let x2 = 1 2 6 24 120 720;
let x3 = 13 17 19 23 29 31;
xc2 = x1|x2|x3; /* cube 2 */

xh = xc1|xc2; /* hypercube */
xhfft1 = fftmi(xh,dim);
```

We left out the **vecr** step for the 2nd cube. It's not really necessary when you're constructing the matrices with **let** statements.

dim contains the dimensions of x , beginning with the highest dimension. The last element of *dim* is the number of columns, the next to the last element of *dim* is the number of rows, and so on. Thus

```
dim = { 2, 3, 3 };
```

indicates that the data in x is a 2x3x3 three-dimensional array, i.e., two 3x3 matrices of data. Suppose that $x1$ is the first 3x3 matrix and $x2$ the second 3x3 matrix, then

```
x = vecr(x1) | vecr(x2)
```

The size of `dim` tells you how many dimensions `x` has.

The arrays have to be padded in each dimension to the nearest power of two. Thus the output array can be larger than the input array. In the 2x3x2x3 hypercube example, `x` would be padded from 2x3x2x3 out to 2x4x2x4. The input vector would contain 36 elements, while the output vector would contain 64 elements.

Source

`fftm.src`

See Also

[fft](#), [ffti](#), [fftn](#)

fftn

Purpose

Computes a complex 1- or 2-D FFT.

Format

```
y = fftn(x);
```

Input

<code>x</code>	NxK matrix.
----------------	-------------

Output

<code>y</code>	LxM matrix, where L and M are the smallest prime factor products greater than or equal to N and K,
----------------	--

respectively.

Remarks

fftn uses the Temperton prime factor FFT algorithm. This algorithm can compute the FFT of any vector or matrix whose dimensions can be expressed as the product of selected prime number factors. **GAUSS** implements the Temperton algorithm for any power of 2, 3, and 5, and one factor of 7. Thus, **fftn** can handle any matrix whose dimensions can be expressed as

$$2^p \times 3^q \times 5^r \times 7^s$$

where p , q and r are nonnegative integers and s is equal to 0 or 1.

If a dimension of x does not meet this requirement, it will be padded with zeros to the next allowable size before the FFT is computed.

fftn pads matrices to the next allowable dimensions; however, it generally runs faster for matrices whose dimensions are highly composite numbers, i.e., products of several factors (to various powers), rather than powers of a single factor. For example, even though it is bigger, a 33600x1 vector can compute as much as 20% faster than a 32768x1 vector, because 33600 is a highly composite number, $2^6 \times 3 \times 5^2 \times 7$, whereas 32768 is a simple power of 2, 2^{15} . For this reason, you may want to hand-pad matrices to optimum dimensions before passing them to **fftn**. The **Run-Time Library** includes a routine, **optn**, for determining optimum dimensions.

The **Run-Time Library** also includes the **nextn** routine, for determining allowable dimensions for a matrix. (You can use this to see the dimensions to which **fftn** would pad a matrix.)

fftn scales the computed FFT by $1/(L \times M)$.

See Also

[fft](#), [ffti](#), [fftm](#), [fftni](#), [rfft](#), [rffti](#), [rfftip](#), [rfftn](#), [rfftnp](#), [rfftp](#)

fgets

Purpose

Reads a line of text from a file.

Format

```
str = fgets(f, maxsize);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
<i>maxsize</i>	scalar, maximum size of string to read in, including the terminating null byte.

Output

<i>str</i>	string.
------------	---------

Remarks

fgets reads text from a file into a string. It reads up to a newline, the end of the file, or *maxsize*-1 characters. The result is placed in *str*, which is then terminated with a null byte. The newline, if present, is retained.

If the file is already at end-of-file when you call **fgets**, your program will terminate with an error. Use **eof** in conjunction with **fgets** to avoid this.

If the file was opened for update (see **fopen**) and you are switching from writing to reading, don't forget to call **fseek** or **fflush** first, to flush the file's buffer.

If you pass **fgets** the handle of a file opened with [open](#) (i.e., a data set or matrix file), your program will terminate with a fatal error.

fgetsa

See Also

[fgetst](#), [fgetsa](#), [fopen](#)

fgetsa

Purpose

Reads lines of text from a file into a string array.

Format

```
sa = fgetsa(f, numl);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
<i>numl</i>	scalar, number of lines to read.

Output

<i>sa</i>	Nx1 string array, N <= <i>numl</i> .
-----------	--------------------------------------

Remarks

fgetsa reads up to *numl* lines of text. If **fgetsa** reaches the end of the file before reading *numl* lines, *sa* will be shortened. Lines are read in the same manner as **fgets**, except that no limit is placed on the size of a line. Thus, **fgetsa** always returns complete lines of text. Newlines are retained. If *numl* is 1, **fgetsa** returns a string. (This is one way to read a line from a file without placing a limit on the length of the line.)

If the file is already at end-of-file when you call **fgetsat**, your program will terminate with an error. Use **eof** in conjunction with **fgetsat** to avoid this. If the file was opened for update (see **fopen**) and you are switching from writing to reading, don't forget to call **fseek** or **fflush** first, to flush the file's buffer.

If you pass **fgetsat** the handle of a file opened with [open](#) (i.e., a data set or matrix file), your program will terminate with a fatal error.

See Also

[fgetsat](#), [fgets](#), [fopen](#)

fgetsat

Purpose

Reads lines of text from a file into a string array.

Format

```
sa = fgetsat(f, numl);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
<i>numl</i>	scalar, number of lines to read.

Output

<i>sa</i>	Nx1 string array, N <= <i>numl</i> .
-----------	--------------------------------------

Remarks

fgetsat operates identically to **fgetsat**, except that newlines are not retained as

fgetst

text is read into *sa*.

In general, you don't want to use **fgetsat** on files opened in binary mode (see **fopen**). **fgetsat** drops the newlines, but it does NOT drop the carriage returns that precede them on some platforms. Printing out such a string array can produce unexpected results.

See Also

[fgetsa](#), [fgetst](#), [fopen](#)

fgetst

Purpose

Reads a line of text from a file.

Format

```
str = fgetst(f, maxsize);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
<i>maxsize</i>	scalar, maximum size of string to read in, including the null terminating byte.

Output

<i>str</i>	string.
------------	---------

Remarks

fgetst operates identically to **fgets**, except that the newline is not retained in the

string.

In general, you don't want to use **fgetst** on files opened in binary mode (see **fopen**). **fgetst** drops the newline, but it does NOT drop the preceding carriage return used on some platforms. Printing out such a string can produce unexpected results.

See Also

[fgets](#), [fgetsat](#), [fopen](#)

fileinfo

Purpose

Returns names and information for files that match a specification.

Format

```
{ fnames, finfo } = fileinfo(fspec);
```

Input

<i>fspec</i>	string, file specification. Can include path. Wildcards are allowed in <i>fspec</i> .
--------------	---

Output

<i>fnames</i>	Nx1 string array of all file names that match, null string if none are found.
<i>finfo</i>	Nx13 matrix, information about matching files.
	Linux
	<i>[N, 1]</i> filesystem ID

fileinfo

<i>[N, 2]</i>	inode number
<i>[N, 3]</i>	mode bit mask
<i>[N, 4]</i>	number of links
<i>[N, 5]</i>	user ID
<i>[N, 6]</i>	group ID
<i>[N, 7]</i>	device ID (char/block special files only)
<i>[N, 8]</i>	size in bytes
<i>[N, 9]</i>	last access time
<i>[N, 10]</i>	last data modification time
<i>[N, 11]</i>	last file status change time
<i>[N, 12]</i>	preferred I/O block size
<i>[N, 13]</i>	number of 512-byte blocks allocated

Windows

<i>[N, 1]</i>	drive number (A = 0, B = 1, etc.)
<i>[N, 2]</i>	n/a, 0
<i>[N, 3]</i>	mode bit mask
<i>[N, 4]</i>	number of links, always 1
<i>[N, 5]</i>	n/a, 0
<i>[N, 6]</i>	n/a, 0
<i>[N, 7]</i>	n/a, 0
<i>[N, 8]</i>	size in bytes
<i>[N, 9]</i>	last access time
<i>[N, 10]</i>	last data modification time
<i>[N, 11]</i>	creation time
<i>[N, 12]</i>	n/a, 0
<i>[N, 13]</i>	n/a, 0

fileinfo will be a scalar zero if no matches are found.

Remarks

fnames will contain file names only; any path information that was passed is dropped.

The time stamp fields (*finfo*[N,9:11]) are expressed as the number of seconds since midnight, Jan. 1, 1970, Coordinated Universal Time (UTC).

See Also

[filesa](#)

filesa

Purpose

Returns a string array of file names.

Format

y = **filesa**(*n*);

Input

<i>n</i>	string, file specification to search for. Can include path. Wildcards are allowed in <i>n</i> .
----------	---

Output

<i>y</i>	Nx1 string array of all file names that match, or null string if none are found.
----------	--

Remarks

y will contain file names only; any path information that was passed is dropped.

floor

Example

```
y = filesa("ch*");
```

In this example all files listed in the current directory that begin with "ch" will be returned.

```
proc  
  exist(filename);  
  retp(not filesa(filename)  $\$==$  "");  
endp;
```

This procedure will return 1 if the file exists or 0 if not.

See Also

[fileinfo](#), [shell](#)

floor

Purpose

Round down toward $-\infty$.

Format

```
y = floor(x);
```

Input

x

NxK matrix or N-dimensional array.

Output

y

NxK matrix or N-dimensional array containing the

elements of x rounded down.

Remarks

This rounds every element in x down to the nearest integer.

Example

```
//Set the seed for repeatable random numbers
rndseed 9072345;

//Create random normal numbers with a standard
//deviation of 100
x = 100*rndn(2,2);

//Round the numbers down
f = floor(x);

//Format so numbers will print in decimal form rather than
//scientific notation) and will show 2 digits after the
//decimal point
format /rd 8,2;

print "*****";
print "After running this code:";
print "*****\n";
print "x = " x;
print "";
print "and, f = " f;
```

produces:

```
*****
After running this code:
*****
```

fmod

```
x =
    0.11    314.05
   -80.87    103.73

and, f =
    0.00    314.00
   -81.00    103.00
```

Notice in the code above, how the `\n` at the end of the statement printing the line of asterisks, inserts a newline.

See Also

[ceil](#), [round](#), [trunc](#)

fmod

Purpose

Computes the floating-point remainder of x/y .

Format

```
r = fmod(x, y);
```

Input

x	$N \times K$ matrix.
y	$L \times M$ matrix, $E \times E$ conformable with x .

Output

r	$\max(N,L)$ by $\max(K,M)$ matrix.
-----	------------------------------------

Remarks

Returns the floating-point remainder r of x/y such that $x = iy + r$, where i is an integer, r has the same sign as x and $|r| < |y|$.

Compare this with `%`, the modulo division operator. (See **OPERATORS**, CHAPTER 1.)

Example

This example extracts all of the years which are evenly divisible by four, from a vector with all of the years between 1900 and 2000.

```
//Create a vector with all years from 1900 to 2000
//i.e. 1900, 1901, 1902...2000
yrs = seqa(1900, 1, 101);

//Create an empty matrix into which we can put our output
y4 = {};

//Loop through each element in yrs
for i(1, rows(yrs), 1);
    //If the 'i'th element of 'yrs' is evenly divisible by
    //4, vertically concatenate it on to the bottom of 'y4'
    if not fmod(yrs[i], 4);
        y4 = y4|yrs[i];
    endif;
endfor;

//No digits after the decimal place
format /rd 8,0;

//Split 'y4' into two columns, each with half of the data
//and print the columns next to each other
print y4[1:13]~y4[14:26];
```

produces:

fn

1900	1952
1904	1956
1908	1960
1912	1964
1916	1968
1920	1972
1924	1976
1928	1980
1932	1984
1936	1988
1940	1992
1944	1996
1948	2000

fn

Purpose

Allows user to create one-line functions.

Format

```
fn fn_name(args) = code_for_function;
```

Remarks

Functions can be called in the same way as other procedures.

Example

```
fn area(r) = pi*r*r;  
a = area(4);
```

After the code above:

```
a = 50.2625
```

fonts

Purpose

Loads fonts to be used in the graph. Note: this function is for the deprecated PQG graphics.

Library

pgraph

Format

```
fonts(str);
```

Input

<i>str</i>	string or character vector containing the names of fonts to be used in the plot. The following fonts are available:
Simplex	standard sans serif font.
Simgrma	Simplex greek, math.
Microb	bold and boxy.
Complex	standard font with serif.

Remarks

The first font specified will be used for the axes numbers.

If *str* is a null string, or **fonts** is not called, Simplex is loaded by default.

fopen

For more information on how to select fonts within a text string, see **PUBLICATION QUALITY GRAPHICS**, CHAPTER 1.

Source

`pgraph.src`

See Also

[title](#), [xlabel](#), [ylabel](#), [zlabel](#)

fopen

Purpose

Opens a file.

Format

```
f = fopen(filename, omode);
```

Input

<i>filename</i>	string, name of file to open.
<i>omode</i>	string, file I/O mode. (See Remarks, below.)

Output

<i>f</i>	scalar, file handle.
----------	----------------------

Portability

Linux/Mac

Carriage return-linefeed conversion for files opened in text mode is unnecessary, because in Linux/Mac a newline is simply a linefeed.

Remarks

filename can contain a path specification.

omode is a sequence of characters that specify the mode in which to open the file. The first character must be one of:

- r* Open an existing file for reading. If the file does not exist, **fopen** fails.
- w* Open or create a file for writing. If the file already exists, its current contents will be destroyed.
- a* Open or create a file for appending. All output is appended to the end of the file.

To this can be appended a *+* and/or a *b*. The *+* indicates the file is to be opened for reading and writing, or update, as follows:

- r+* Open an existing file for update. You can read from or write to any location in the file. If the file does not exist, **fopen** fails.
- w+* Open or create a file for update. You can read from or write to any location in the file. If the file already exists, its current contents will be destroyed.
- a+* Open or create a file for update. You can read from any location in the file, but all output will be appended to the end of the file.

Finally, the *b* indicates whether the file is to be opened in text or binary mode. If the file is opened in binary mode, the contents of the file are read verbatim; likewise, anything output to the file is written verbatim. In text mode (the default), carriage return-

for

linefeed sequences are converted on input to linefeeds, or newlines. Likewise on output, newlines are converted to carriage return-linefeeds. Also in text mode, if a CTRL+Z (char 26) is encountered during a read, it is interpreted as an end-of-file character, and reading ceases. In binary mode, CTRL+Z is read in uninterpreted.

The order of *+* and *b* is not significant; *rb+* and *r+b* mean the same thing.

You can both read from and write to a file opened for update. However, before switching from one to the other, you must make an **fseek** or **fflush** call, to flush the file's buffer.

If **fopen** fails, it returns a 0.

Use **close** and **closeall** to close files opened with **fopen**.

See Also

[fseek](#), [close](#), [closeall](#)

for

Purpose

Begins a **for** loop.

Format

```
for i(start, stop, step);  
.  
.  
.  
endfor;
```

Input

i literal, the name of the counter variable.

<i>start</i>	scalar expression, the initial value of the counter.
<i>stop</i>	scalar expression, the final value of the counter.
<i>step</i>	scalar expression, the increment value.

Remarks

The counter is strictly local to the loop. The expressions, *start*, *stop* and *step* are evaluated only once when the loop initializes and are stored local to the loop.

The `for` loop is optimized for speed and much faster than a `do` loop.

The commands `break` and `continue` are supported. The `continue` command steps the counter and jumps to the top of the loop. The `break` command terminates the current loop.

The loop terminates when the value of *i* exceeds *stop*. If `break` is used to terminate the loop and you want the final value of the counter, you need to assign it to a variable before the `break` statement (see the third example, following).

Example

Example 1

```
//A basic 'for' loop
for i (1, 4, 1);
    print i;
endfor;
```

The code above, will print out:

```
1.000
2.000
3.000
4.000
```

Example 2

for

```
x = zeros(10,5);
for i (1, rows(x), 1);
    for j (1, cols(x), 1);
        x[i,j] = i*j;
    endfor;
endfor;
```

Example 3

```
x = randn(3,3);
y = randn(3,3);

for i (1, rows(x), 1);
    for j (1, cols(x), 1);
        if x[i,j] >= y[i,j];
            continue;
        endif;
        temp = x[i,j];
        x[i,j] = y[i,j];
        y[i,j] = temp;
    endfor;
endfor;
```

Example 4

```
li = 0;
x = randn(100,1);
y = randn(100,1);

for i (1, rows(x), 1);
    if x[i] != y[i];
        li = i;
        break;
    endif;
endfor;

if li;
```



```
print "Compare failed on row " li;
endif;
```

format

Purpose

Controls the format of matrices and numbers printed out with `print` statements.

Format

```
format [[/typ]] [[/fmted]] [[/mf]] [[/jnt]] [[f,p]]
```

Input

<code>/typ</code>	literal, symbol type flag(s). Indicate which symbol types you are setting the output format for. <code>/mat, /sa, /str</code>	Formatting parameters are maintained separately for matrices and arrays (<code>/mat</code>), string arrays (<code>/sa</code>), and strings (<code>/str</code>). You can specify more than one <code>/typ</code> flag; the format will be set for all types indicated. If no <code>/typ</code> flag is listed, <code>format</code> assumes <code>/mat</code> .
<code>/fmted</code>	literal, enable formatting flag. <code>/on, /off</code>	Enable/disable formatting.

format

f

/mf literal, matrix row format flag.
 /m0

/m1 or /mb1

/m2 or /mb2

/m3 or /mb3

When formatting is disabled, the contents of a variable are dumped to the screen in a "raw" format. */off* is currently supported only for strings. "Raw" format for strings means that the entire string is printed, starting at the current cursor position. When formatting is enabled for strings, they are handled the same as string arrays. This shouldn't be too surprising, since a string is actually a 1x1 string array.

no delimiters before or after rows when printing out matrices.

print 1 carriage return/line feed pair before each row of a matrix with more than 1 row.

print 2 carriage return/line feed pairs before each row of a matrix with more than 1 row.

print "Row 1", "Row 2"... before each row of a matrix with more than one row.

<code>/ma1</code>	print 1 carriage return/line feed pair after each row of a matrix with more than 1 row.
<code>/ma2</code>	print 2 carriage return/line feed pairs after each row of a matrix with more than 1 row.
<code>/a1</code>	print 1 carriage return/line feed pair after each row of a matrix.
<code>/a2</code>	print 2 carriage return/line feed pairs after each row of a matrix.
<code>/b1</code>	print 1 carriage return/line feed pair before each row of a matrix.
<code>/b2</code>	print 2 carriage return/line feed pairs before each row of a matrix.
<code>/b3</code>	print "Row 1", "Row 2"... before each row of a matrix.
<code>/jnt</code>	literal, matrix element format flag - controls justification, notation and trailing character.
Right-Justified	
<code>/rd</code>	Signed decimal number in the form #####.####, where ##### is one or more decimal digits. The number of digits before the decimal point depends on the

format

f

/re

/ro

/rz

magnitude of the number, and the number of digits after the decimal point depends on the precision. If the precision is 0, no decimal point will be printed.

Signed number in the form $\# . \# \# E \pm \# \# \#$, where $\#$ is one decimal digit, $\# \#$ is one or more decimal digits depending on the precision, and $\# \# \#$ is three decimal digits. If precision is 0, the form will be $[-] \# E \pm \# \# \#$ with no decimal point printed.

This will give a format like */rd* or */re* depending on which is most compact for the number being printed. A format like */re* will be used only if the exponent value is less than -4 or greater than the precision. If a */re* format is used, a decimal point will always appear. The precision signifies the number of significant digits displayed.

This will give a format like

Left-Justified*/ld*

/rd or */re* depending on which is most compact for the number being printed. A format like */re* will be used only if the exponent value is less than -4 or greater than the precision. If a */re* format is used, trailing zeros will be suppressed and a decimal point will appear only if one or more digits follow it. The precision signifies the number of significant digits displayed.

Signed decimal number in the form $[-]####.####$, where $####$ is one or more decimal digits. The number of digits before the decimal point depends on the magnitude of the number, and the number of digits after the decimal point depends on the precision. If the precision is 0, no decimal point will be printed. If the number is positive, a space character will replace the leading minus sign.

format

f

`/le`

Signed number in the form `[-] # . ##E±###`, where `#` is one decimal digit, `##` is one or more decimal digits depending on the precision, and `###` is three decimal digits. If precision is 0, the form will be `[-] #E±###` with no decimal point printed. If the number is positive, a space character will replace the leading minus sign.

`/lo`

This will give a format like `/ld` or `/le` depending on which is most compact for the number being printed. A format like `/le` will be used only if the exponent value is less than -4 or greater than the precision. If a `/le` format is used, a decimal point will always appear. If the number is positive, a space character will replace the leading minus sign. The precision specifies the number of significant digits displayed.

`/lz`

This will give a format like

`/ld` or `/le` depending on which is most compact for the number being printed. A format like `/le` will be used only if the exponent value is less than -4 or greater than the precision. If a `/le` format is used, trailing zeros will be suppressed and a decimal point will appear only if one or more digits follow it. If the number is positive, a space character will replace the leading minus sign. The precision specifies the number of significant digits displayed.

Trailing Character

The following characters can be added to the `/jnt` parameters above to control the trailing character if any:

```
format /rdn 1,3;
```

<i>s</i>	The number will be followed immediately by a space character. This is the default.
<i>c</i>	The number will be followed immediately by a comma.
<i>t</i>	The number will be followed immediately by a tab character.

format

<i>n</i>	No trailing character.
<i>f</i>	scalar expression, controls the field width.
<i>p</i>	scalar expression, controls the precision.

Remarks

If character elements are to be printed, the precision should be at least 8 or the elements will be truncated. This does not affect the string data type.

For numeric values in matrices, *p* sets the number of significant digits to be printed. For string arrays, strings, and character elements in matrices, *p* sets the number of characters to be printed. If a string is shorter than the specified precision, the entire string is printed. For string arrays and strings, *p* = -1 means print the entire string, regardless of its length *p* = -1 is illegal for matrices; setting *p* >= 8 means the same thing for character elements.

The */xxx* slash parameters are optional. Field and precision are optional also, but if one is included, then both must be included.

Slash parameters, if present, must precede the field and precision parameters.

A `format` statement stays in effect until it is overridden by a new `format` statement. The slash parameters may be used in a `print` statement to override the current default.

f and *p* may be any legal expressions that return scalars. Nonintegers will be truncated to integers.

The total width of field will be overridden if the number is too big to fit into the space allotted. For instance, `format /rds 1,0` can be used to print integers with a single space between them, regardless of the magnitudes of the integers.

Complex numbers are printed with the sign of the imaginary half separating them and an "i" appended to the imaginary half. Also, the field parameter refers to the width of field for each half of the number, so a complex number printed with a field of 8 will actually take (at least) 20 spaces to print. The character printed after the imaginary part can be changed (for example, to a "j") with the **sysstate** function, case 9.

The default when **GAUSS** is first started is:

```
format /mb1 /ros 16,8;
```

Example

This code:

```
x = rndn(3,3);

format /m1 /rd 16,8;
print x;
```

produces:

2.25240104	0.53724423	-0.67744907
-0.16183998	1.57152099	1.33836836
0.00666162	-1.24948147	-0.77987532

This code:

```
format /m1 /rzs 1,10;
print x;
```

produces:

2.252401038	0.5372442301	-0.6774490661
-0.1618399808	1.571520994	1.338368355
0.00666161784	-1.24948147	-0.7798753222

This code:

```
format /m3 /rdn 16,4;
print x;
```

format

produces:

```
print x;
```

Row 1	2.2524	0.5372	-0.6774
Row 2	-0.1618	1.5715	1.3384
Row 3	0.0067	-1.2495	-0.7799

This code:

```
format /m1 /ldn 16,4;  
print x;
```

produces:

2.2524	0.5372	-0.6774
-0.1618	1.5715	1.3384
0.0067	-1.2495	-0.7799

This code:

```
format /m1 /res 12,4;  
print x;
```

produces:

2.2524e+000	5.3724e-001	-6.7745e-001
-1.6184e-001	1.5715e+000	1.3384e+000
6.6616e-003	-1.2495e+000	-7.7988e-001

See Also

[formatcv](#), [formatnv](#), [print](#), [output](#)

formatcv

Purpose

Sets the character data format used by **printfmt**.

Format

```
oldfmt = formatcv(newfmt);
```

Input

<i>newfmt</i>	1x3 vector, the new format specification.
---------------	---

Output

<i>oldfmt</i>	1x3 vector, the old format specification.
---------------	---

Remarks

See **printfm** for details on the format vector.

Example

This example saves the old format, sets the format desired for printing *x*, prints *x*, then restores the old format. This code:

```
x = { A 1, B 2, C 3 };  
oldfmt = formatcv("*.*s" ~ 3 ~ 3);  
call printfmt(x,0~1);  
call formatcv(oldfmt);
```

produces:

formatnv

```
A 1
B 2
C 3
```

Source

gauss.src

Globals

__fmtcv

See Also

[formatnv](#), [printfm](#), [printfmt](#)

formatnv

Purpose

Sets the numeric data format used by **printfmt**.

Format

```
oldfmt = formatnv(newfmt);
```

Input

<i>newfmt</i>	1x3 vector, the new format specification.
---------------	---

Output

<i>oldfmt</i>	1x3 vector, the old format specification.
---------------	---

Remarks

See [printfm](#) for details on the format vector.

Example

This example saves the old format, sets the format desired for printing `x`, prints `x`, then restores the old format. This code:

```
x = { A 1, B 2, C 3 };
oldfmt = formatnv("%.1f" ~ 8 ~ 4);
call printfmt(x, 0~1);
call formatnv(oldfmt);
```

produces:

```
A 1.0000
B 2.0000
C 3.0000
```

Source

gauss.src

Globals

`__fmtnv`

See Also

[formatev](#), [printfm](#), [printfmt](#)

fputs

Purpose

Writes strings to a file.

fputs

Format

```
numl = fputs(f, sa);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
<i>sa</i>	string or string array.

Output

<i>numl</i>	scalar, the number of lines written to the file.
-------------	--

Examples

Example 1: Write string to text file

```
//Create string
quote = "There is nothing either good or bad, but thinking
makes it so.";

//Open file for writing
fh = fopen("hamlet.txt", "w");

//Write the string to the first line of the file
call fputs(fh, quote);

//Close the file
call close(fh);
```

After the code above, you should have a file named `hamlet.txt` in your current working directory, containing the contents of the `quote` string.

Example 2: Write CSV data to text file

```
//Create string containing a comma separated list
//of variable names and an ending newline
text = "alpha,beta,gamma,delta\n";

fh = fopen("temp.csv", "w");

//Write the string to the first line of the file
call fputs(fh, text);

//Create some numeric data
x = { 1 2 3 4,
      5 6 7 8 };

//Convert numeric data to 2x4 string array
x_str = ntos(x);

//Combine each row of 'x_str' into
//a single comma separated string
x_str = strjoin(x_str, ",");

//Add newlines to the end of each line
x_str = x_str $+ "\n";

//Write the comma separated data to the file
call fputs(fh, x_str);

//Close the file
call close(fh);
```

After the above code, you should have a file named `temp.csv` with the following contents:

```
alpha,beta,gamma,delta
1,2,3,4
5,6,7,8
```

fputst

Remarks

- To write to the standard output stream or the standard error stream, pass in `__STDOUT` or `__STDERR` as the file handle argument.

```
str = "sample string";
num = fputs(__STDOUT, str);
```

- **fputs** writes the contents of each string in `sa`, minus the null terminating byte, to the file specified. If the file was opened in text mode (see **fopen**), any newlines present in the strings are converted to carriage return-linefeed sequences on output. If `num1` is not equal to the number of elements in `sa`, there may have been an I/O error while writing the file. You can use **fcheckerr** or **fclearerr** to check this. If there was an error, you can call **fstrerror** to find out what it was. If the file was opened for update (see **fopen**) and you are switching from reading to writing, don't forget to call **fseek** or **fflush** first, to flush the file's buffer. If you pass **fputs** the handle of a file opened with [open](#) (i.e., a data set or matrix file), your program will terminate with a fatal error.

Portability

Linux/Mac

Carriage return-linefeed conversion for files opened in text mode is unnecessary, because in Linux/Mac a newline is simply a linefeed.

See Also

[fputst](#), [fopen](#)

fputst

Purpose

Writes strings followed by a newline to a file.

Format

```
numl = fputst(f, sa);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
<i>sa</i>	string or string array.

Output

<i>numl</i>	scalar, the number of lines written to the file.
-------------	--

Examples

Example 1: Write string to text file

```
//Create string
quote = "A horse! a horse! my kingdom for a horse!.";

//Open file for writing
fh = fopen("king_richard_III.txt", "w");

//Write the string to the first line of the file
call fputst(fh, quote);

//Close the file
call close(fh);
```

After the code above, you should have a file named `king_richard_III.txt` in your current working directory, containing the contents of the quote string followed by an empty line. To avoid the final empty line, use **fputs**.

Example 2: Write CSV data to text file

fputst

```
//Create string containing a comma separated list
//of variable names (fputst will add an ending newline)
text = "alpha,beta,gamma,delta";

fh = fopen("temp.csv", "w");

//Write the string to the first line of the file
call fputst(fh, text);

//Create some numeric data
x = { 1 2 3 4,
      5 6 7 8 };

//Convert numeric data to 2x4 string array
x_str = ntos(x);

//Combine each row of 'x_str' into
//a single comma separated string
x_str = strjoin(x_str, ",");

//Write the comma separated data to the file
call fputst(fh, x_str);

//Close the file
call close(fh);
```

After the above code, you should have a file named `temp.csv` with the following contents:

```
alpha,beta,gamma,delta
1,2,3,4
5,6,7,8
```

Portability

Linux/Mac

Carriage return-linefeed conversion for files opened in text mode is unnecessary, because in Linux/Mac a newline is simply a linefeed.

Remarks

- To write to the standard output stream or the standard error stream, pass in `__STDOUT` or `__STDERR` as the file handle argument.

```
str = "sample string";
num = fputst(__STDOUT, str);
```

- **fputst** works identically to **fputs**, except that a newline is appended to each string that is written to the file. If the file was opened in text mode (see **fopen**), these newlines are also converted to carriage return-linefeed sequences on output.

See Also

[fputs](#), [fopen](#)

fseek

Purpose

Positions the file pointer in a file.

Format

```
ret = fseek(f, offs, base);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
<i>offs</i>	scalar, offset (in bytes).
<i>base</i>	scalar, base position.

fseek

0	beginning of file.
1	current position of file pointer.
2	end of file.

Output

ret scalar, 0 if successful, 1 if not.

Portability

Linux/Mac

Carriage return-linefeed conversion for files opened in text mode is unnecessary, because in Linux/Mac a newline is simply a linefeed.

Remarks

fseek moves the file pointer *offs* bytes from the specified *base* position. *offs* can be positive or negative. The call may fail if the file buffer needs to be flushed (see **fflush**).

If **fseek** fails, you can call **fstrerror** to find out why.

For files opened for update (see **fopen**), the next operation can be a read or a write.

fseek is not reliable when used on files opened in text mode (see **fopen**). This has to do with the conversion of carriage return-linefeed sequences to newlines. In particular, an **fseek** that follows one of the **fgetxxx** or **fputxxx** commands may not produce the expected result. For example:

```
p = ftell(f);  
s = fgetsa(f, 7);  
call fseek(f, p, 0);
```

is not reliable. We have found that the best results are obtained by **fseek**'ing to the beginning of the file and then **fseek**'ing to the desired location, as in

```
p = ftell(f);  
s = fgetsa(f, 7);  
call fseek(f, 0, 0);  
call fseek(f, p, 0);
```

If you pass **fseek** the handle of a file opened with **open** (i.e., a data set or matrix file), your program will terminate with a fatal error.

See Also

[fopen](#)

fstrerror

Purpose

Returns an error message explaining the cause of the most recent file I/O error.

Format

```
s = fstrerror;
```

Output

<i>s</i>	string, error message.
----------	------------------------

Remarks

Any time an I/O error occurs on a file opened with **fopen**, an internal error flag is updated. (This flag, unlike those accessed by **fcheckerr** and **fclearerr**, is not specific to a given file; rather, it is system-wide.) **fstrerror** returns an error message based on the value of this flag, clearing it in the process. If no error has occurred, a null string is returned.

ftell

Since **fstrerror** clears the error flag, if you call it twice in a row, it will always return a null string the second time.

The Windows system command called by **ftell** does not set the internal error flag accessed by **fstrerror**. Therefore, calling **fstrerror** after **ftell** on Windows will not produce any error information.

See Also

[fopen](#), [ftell](#)

ftell

Purpose

Gets the position of the file pointer in a file.

Format

```
pos = ftell(f);
```

Input

<i>f</i>	scalar, file handle of a file opened with fopen .
----------	--

Output

<i>pos</i>	scalar, current position of the file pointer in a file.
------------	---

Remarks

ftell returns the position of the file pointer in terms of bytes from the beginning of the file. The call may fail if the file buffer needs to be flushed (see **fflush**).

If an error occurs, **ftell** returns -1. You can call **fstrerror** to find out what the error was.

If you pass **ftell** the handle of a file opened with [open](#) (i.e., a data set or matrix file), your program will terminate with a fatal error.

See Also

[fopen](#), [fseek](#)

ftocv

Purpose

Converts a matrix containing floating point numbers into a matrix containing the decimal character representation of each element.

Format

```
y = ftocv(x, field, prec);
```

Input

<i>x</i>	NxK matrix containing numeric data to be converted.
<i>field</i>	scalar, minimum field width.
<i>prec</i>	scalar, the numbers created will have <i>prec</i> places after the decimal point.

Output

<i>y</i>	NxK matrix containing the decimal character equivalent of the corresponding elements in <i>x</i> in the format defined by <i>field</i> and <i>prec</i> .
----------	--

ftos

Remarks

If a number is narrower than *field*, it will be padded on the left with zeros.

If *prec* = 0, the decimal point will be suppressed.

Example

```
y = seqa(6,1,5);  
x = 0 $+ "beta" $+ ftocv(y,2,0);  
print $x;
```

results in the following output:

```
beta06  
beta07  
beta08  
beta09  
beta10
```

Notice that the (0 \$+) above was necessary to force the type of the result to matrix because the string constant "beta" would be of type string. The left operand in an expression containing a \$+ operator controls the type of the result.

See Also

[ftos](#)

ftos

Purpose

Converts a scalar into a string containing the decimal character representation of that number.

Format

```
y = ftos(x, fmat, field, prec);
```

Input

<i>x</i>	scalar, the number to be converted.
<i>fmat</i>	string, the format string to control the conversion.
<i>field</i>	scalar or 2x1 vector, the minimum field width. If <i>field</i> is 2x1, it specifies separate field widths for the real and imaginary parts of <i>x</i> .
<i>prec</i>	scalar or 2x1 vector, the number of places following the decimal point. If <i>prec</i> is 2x1, it specifies separate precisions for the real and imaginary parts of <i>x</i> .

Output

<i>y</i>	string containing the decimal character equivalent of <i>x</i> in the format specified.
----------	---

Remarks

The format string corresponds to the `format/jnt` (justification, notation, trailing character) slash parameter as follows:

<code>/rdn</code>	<code>"%*.*lf"</code>
<code>/ren</code>	<code>"%*.*le"</code>
<code>/ron</code>	<code>"%#*.*lG"</code>

```
/rzn      "%*.*lG"  
/ldn      "%- *.*lf"  
/len      "%- *.*lE"  
/lon      "%-# *.*lG"  
/lzn      "%- *.*lG"
```

If x is complex, you can specify separate formats for the real and imaginary parts by putting two format specifications in the format string. You can also specify separate fields and precisions. You can position the sign of the imaginary part by placing a "+" between the two format specifications. If you use two formats, no "*i*" is appended to the imaginary part. This is so you can use an alternate format if you prefer, for example, prefacing the imaginary part with a "j".

The format string can be a maximum of 80 characters.

If you want special characters to be printed after x , include them as the last characters of the format string. For example:

`"%*.*lf, "` right-justified decimal followed by a comma.

`"%-*.*s "` left-justified string followed by a space.

`"%*.*lf"` right-justified decimal followed by nothing.

You can embed the format specification in the middle of other text:

```
"Time: %*.*lf seconds."
```

If you want the beginning of the field padded with zeros, then put a "0" before the first "*" in the format string:

`"%0*.*lf"` right-justified decimal.

If `prec = 0`, the decimal point will be suppressed.

Example

You can create custom formats for complex numbers with **ftos**. For example,

```
let c = 24.56124+6.3224e-2i;

field = 1;
prec = 3|5;
fmat = "%lf + j%le is a complex number.";
cc = ftos(c,fmat,field,prec);
```

results in

```
cc = "24.561 + j6.32240e-02 is a complex number."
```

Some other things you can do with **ftos**:

```
let x = 929.857435324123;
let y = 5.46;
let z = 5;

field = 1;
prec = 0;
fmat = "%*.*lf";
zz = ftos(z,fmat,field,prec);

field = 1;
prec = 10;
fmat = "%*.*1E";
xx = ftos(x,fmat,field,prec);

field = 7;
prec = 2;
fmat = "%*.*lf seconds";
s1 = ftos(x,fmat,field,prec);
s2 = ftos(y,fmat,field,prec);
```

ftostrC

```
field = 1;
prec = 2;
fmt = "The maximum resistance is %*.*lf ohms.";
om = ftos(x,fmt,field,prec);
```

The results:

```
zz = "5"

xx = "9.2985743532E+002"

s1 = "929.86 seconds"

s2 = "5.46 seconds"

om = "The maximum resistance is 929.86 ohms."
```

See Also

[ftocv](#), [stof](#), [format](#)

ftostrC

Purpose

Converts a matrix to a string array using a C language format specification.

Format

```
sa = ftostrC(x, fmt);
```

Input

x	$N \times K$ matrix, real or complex.
fmt	$K \times 1$, $1 \times K$ or 1×1 string array containing format information.

Output

sa	$N \times K$ string array.
------	----------------------------

Remarks

If fmt has K elements, each column of sa can be formatted separately. If x is complex, there must be two format specifications in each element of fmt .

Example

```
declare string fmtr = { "%6.3lf",  
                        "%11.8lf" };  
  
declare string fmtc = { "(%6.3lf, %6.3lf)",  
                        "(%11.8lf, %11.8lf)" };  
  
xr = rndn(4, 2);  
xc = sqrt(xr)';  
  
sar = ftostrC(xr, fmtr);  
sac = ftostrC(xc, fmtc);  
  
print sar;  
print sac;
```

produces:

ftostrC

```
-0.166 1.05565441
-1.590 -0.79283296
 0.130 -1.84886957
 0.789 0.86089687

( 0.000, -0.407) ( 1.02745044, 0.00000000)
( 0.000, -1.261) ( 0.00000000, -0.89041168)
( 0.361, 0.000) ( 0.00000000, -1.35973143)
( 0.888, 0.000) ( 0.92784529, 0.00000000)
```

See Also

[strtof](#), [strtocfplx](#)

g

gamma

Purpose

Returns the value of the gamma function.

Format

```
y = gamma(x);
```

Input

x NxK matrix or N-dimensional array.

Output

y NxK matrix or N-dimensional array.

Remarks

For each element of x this function returns the integral

$$\int_0^{\infty} t^{(x-1)} e^{-t} dt$$

gammacplx

All elements of x must be positive and less than or equal to 169. Values of x greater than 169 will cause an overflow.

The natural log of **gamma** is often what is required and it can be computed without the overflow problems of **gamma** using **lnfact**.

Example

```
y = gamma(2.5);
```

After the code above:

```
y = 1.329340
```

See Also

[cdfchic](#), [cdfbeta](#), [cdfFc](#), [cdfnc](#), [cdfFc](#), [erf](#), [erfc](#), [lnfact](#)

gammacplx

Purpose

Computes the Gamma function for complex inputs.

Format

```
f = gammacplx(z);
```

Input

z NxK matrix; z may be complex.

Output

f NxK matrix; f may be complex.

Technical Notes

Accuracy is 15 significant digits along the real axis and 13 significant digits elsewhere. This routine uses the Lanczos series approximation for the complex Gamma function.

References

1. C. Lanczos, SIAM JNA 1, 1964, pp. 86-96.
2. Y. Luke, "The Special ... approximations," 1969, pp. 29-31.
3. Y. Luke, "Algorithms ... functions," 1977.
4. J. Spouge, SIAM JNA 31, 1994, pp. 931-944.
5. W. Press, "Numerical Recipes."
6. S. Chang, "Computation of special functions," 1996.
7. W. J. Cody "An Overview of Software Development for Special Functions," 1975.
8. P. Godfrey "A note on the computation of the convergent Lanczos complex Gamma approximation."
9. Original code by Paul Godfrey

gammai

Purpose

Computes the inverse incomplete gamma function.

gausset

Format

```
x = gammaaii(a, p);
```

Input

<i>a</i>	MxN matrix, exponents.
<i>p</i>	KxL matrix, ExE conformable with <i>a</i> , incomplete gamma values.

Output

<i>x</i>	max(M,K) by max(N,L) matrix, abscissae.
----------	---

Source

`cdfchii.src`

Globals

`__ginvinc`, `__macheps`

gausset

Purpose

Resets the global control variables declared in `gauss.dec`.

Format

```
gausset;
```

Source

`gauss.src`

Globals

__altnam, __con, __ff, __fmtcv, __fmtnv, __header, __miss, __output, __row, __rowfac, __sort, __title, __tol, __vpad, __vtype, __weight

gdaAppend

Purpose

Appends data to a variable in a **GAUSS** Data Archive.

Format

ret = **gdaAppend**(*filename*, *x*, *varname*);

Input

<i>filename</i>	string, name of data file.
<i>x</i>	matrix, array, string or string array, data to append.
<i>varname</i>	string, variable name.

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes:
1	Null file name.
2	File open error.
3	File write error.
4	File read error.

gdaAppend

5	Invalid data file type.
8	Variable not found.
10	File contains no variables.
14	File too large to be read on current platform.
17	Type mismatch.
18	Argument wrong size.
19	Data must be real.
20	Data must be complex.

Remarks

This command appends the data contained in *x* to the variable *varname* in *filename*. Both *x* and the variable referenced by *varname* must be the same data type, and they must both contain the same number of columns.

Because **gdaAppend** increases the size of the variable, it moves the variable to just after the last variable in the data file to make room for the added data, leaving empty bytes in the variable's old location. It also moves the variable descriptor table, so it is not overwritten by the variable data. This does not change the index of the variable because variable indices are determined NOT by the order of the variable data in a GDA, but by the order of the variable descriptors. Call **gdaPack** to pack the data in a GDA, so it contains no empty bytes.

Example

```
x = rndn(100,50);
ret = gdaCreate("myfile.gda",1);
ret = gdaWrite("myfile.gda",x,"x1");

y = rndn(25,50);
ret = gdaAppend("myfile.gda",y,"x1");
```

This example adds 25*50=1250 elements to *x1*, making it a 125x50 matrix.

See Also

[gdaWriteSome](#), [gdaUpdate](#), [gdaWrite](#)

gdaCreate

Purpose

Creates a GAUSS Data Archive.

Format

```
ret = gdaCreate(filename, overwrite);
```

Input

<i>filename</i>	string, name of data file to create.
<i>overwrite</i>	scalar, one of the following: 0 error out if file already exists. 1 overwrite file if it already exists.

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes: 1 Null file name. 3 File write error. 6 File already exists. 7 Cannot create file.
------------	--

gdaDStat

Remarks

This command creates a **GAUSS** Data Archive containing only a header. To add data to the GDA, call **gdaWrite**.

It is recommended that you include a `.gda` extension in `filename`. However, **gdaCreate** will not force an extension.

Example

```
ret = gdaCreate("myfile.gda",1);
```

See Also

[gdaWrite](#)

gdaDStat

Purpose

Computes descriptive statistics on multiple Nx1 variables in a **GAUSS** Data Archive.

Format

```
dout = gdaDStat(dc0, filename, vars);
```

Input

<code>dc0</code>	an instance of a dstatmtControl structure with the following members:
<code>dc0.altnames</code>	Kx1 string array of alternate variable names for the output. Default = "".
<code>dc0.maxbytes</code>	scalar, the maximum number of bytes

	to be read per iteration of the read loop. Default = 1e9.						
<i>dc0.maxvec</i>	scalar, the largest number of elements allowed in any one matrix. Default = 20000.						
<i>dc0.miss</i>	scalar, one of the following: <table> <tr> <td>0</td><td>There are no missing values (fastest).</td></tr> <tr> <td>1</td><td>Listwise deletion, drop a row if any missings occur in it.</td></tr> <tr> <td>2</td><td>Pairwise deletion.</td></tr> </table> Default = 0.	0	There are no missing values (fastest).	1	Listwise deletion, drop a row if any missings occur in it.	2	Pairwise deletion.
0	There are no missing values (fastest).						
1	Listwise deletion, drop a row if any missings occur in it.						
2	Pairwise deletion.						
<i>dc0.output</i>	scalar, one of the following: <table> <tr> <td>0</td><td>Do not print output table.</td></tr> <tr> <td>1</td><td>Print output table.</td></tr> </table> Default = 1.	0	Do not print output table.	1	Print output table.		
0	Do not print output table.						
1	Print output table.						
<i>dc0.row</i>	scalar, the number of rows of <i>var</i> to be read per iteration of the read loop. If 0, (default) the number of rows will be calculated using <i>dc0.maxbytes</i> and <i>dc0.maxvec</i> .						
<i>filename</i>	string, name of data file.						
<i>vars</i>	Kx1 string array, names of variables						
	- or -						

Kx1 vector, indices of variables.

Output

<i>dout</i>	an instance of a dstatmtOut structure with the following members:	
<i>dout.vnames</i>	Kx1 string array,	the names of the variables used in the statistics.
<i>dout.mean</i>	Kx1 vector,	means.
<i>dout.var</i>	Kx1 vector,	variance.
<i>dout.std</i>	Kx1 vector,	standard deviation.
<i>dout.min</i>	Kx1 vector,	minima.
<i>dout.max</i>	Kx1 vector,	maxima.
<i>dout.valid</i>	Kx1 vector,	the number of valid cases.
<i>dout.missing</i>	Kx1 vector,	the number of missing cases.
<i>dout.errcode</i>	scalar, error code, 0 if successful, or one of the following:	
	1	No GDA indicated.
	4	Not implemented for complex data.
	5	Variable must be type matrix.
	6	Too many variables specified.
	7	Too many missings - no data left after packing.
	8	Name variable wrong size.
	9	<i>altnames</i> member of

	dstatmtControl structure wrong size.
11	Data read error.

Remarks

The variables referenced by *vars* must all be Nx1.

The names of the variables in the GDA will be used for the output by default. To use alternate names, set the *altnames* member of the **dstatmtControl** structure.

If pairwise deletion is used, the minima and maxima will be the true values for the valid data. The means and standard deviations will be computed using the correct number of valid observations for each variable.

Example

```
struct dstatmtControl dc0;
struct dstatmtOut dout;

//Set structure to default values
dc0 = dstatmtControlCreate();

vars = { 1,4,5,8 };
dout = gdaDStat(dc0,"myfile.gda",vars);
```

This example computes descriptive statistics on the first, fourth, fifth and eighth variables in `myfile.gda`.

Source

gdadstat.src

gdaDStatMat

See Also

[gdaDStatMat](#), [dstatmtControlCreate](#)

gdaDStatMat

Purpose

Computes descriptive statistics on a selection of columns from a matrix located in a **GAUSS** Data Archive.

Format

```
dout = gdaDStatMat(dc0, filename, gmat, colind, vnamevar);
```

Input

<i>dc0</i>	an instance of a dstatmtControl structure with the following members:
<i>dc0.altnames</i>	Kx1 string array of alternate variable names for the output. Default = "". If set, it must have the same number of rows as <i>colind</i> .
<i>dc0.maxbytes</i>	scalar, the maximum number of bytes to be read per iteration of the read loop. Default = 1e9.
<i>dc0.maxvec</i>	scalar, the largest number of elements allowed in any one matrix. Default = 20000.
<i>dc0.miss</i>	scalar, one of the following: 0 There are no missing

		values (fastest).
	1	Listwise deletion, drop a row if any missings occur in it.
	2	Pairwise deletion.
	Default = 0.	
<i>dc0.output</i>	scalar, one of the following:	
	0	Do not print output table.
	1	Print output table.
	Default = 1.	
<i>dc0.row</i>	scalar, the number of rows of <i>var</i> to be read per iteration of the read loop.	
	If 0, (default) the number of rows will be calculated using <i>dc0.maxbytes</i> and <i>dc0.maxvec</i> .	
<i>filename</i>	string, name of data file.	
<i>gmat</i>	string, name of matrix	
	- or -	
	scalar, index of matrix.	
<i>colind</i>	Kx1 vector, indices of columns in variable to use.	
<i>vnamevar</i>	string, name of the string containing the variable names in the matrix	
	- or -	
	scalar, index of the string containing the variable names in	

the matrix.

Output

<i>dout</i>	an instance of a dstatmtOut structure with the following members:												
<i>dout.vnames</i>	Kx1 string array, the names of the variables used in the statistics.												
<i>dout.mean</i>	Kx1 vector, means.												
<i>dout.var</i>	Kx1 vector, variance.												
<i>dout.std</i>	Kx1 vector, standard deviation.												
<i>dout.min</i>	Kx1 vector, minima.												
<i>dout.max</i>	Kx1 vector, maxima.												
<i>dout.valid</i>	Kx1 vector, the number of valid cases.												
<i>dout.missing</i>	Kx1 vector, the number of missing cases.												
<i>dout.errcode</i>	scalar, error code, 0 if successful, otherwise one of the following: <table><tr><td>1</td><td>No GDA indicated.</td></tr><tr><td>3</td><td>Variable must be Nx1.</td></tr><tr><td>4</td><td>Not implemented for complex data.</td></tr><tr><td>5</td><td>Variable must be type matrix.</td></tr><tr><td>7</td><td>Too many missings, no data left after packing.</td></tr><tr><td>9</td><td><i>altnames</i> member of dstatmtControl structure wrong size.</td></tr></table>	1	No GDA indicated.	3	Variable must be Nx1.	4	Not implemented for complex data.	5	Variable must be type matrix.	7	Too many missings, no data left after packing.	9	<i>altnames</i> member of dstatmtControl structure wrong size.
1	No GDA indicated.												
3	Variable must be Nx1.												
4	Not implemented for complex data.												
5	Variable must be type matrix.												
7	Too many missings, no data left after packing.												
9	<i>altnames</i> member of dstatmtControl structure wrong size.												

Remarks

Set *colind* to a scalar 0 to use all of the columns in *var*.

vnamevar must either reference an Mx1 string array variable containing variable names, where M is the number of columns in the data set variable, or be set to a scalar 0. If *vnamevar* references an Mx1 string array variable, then only the elements indicated by *colind* will be used. Otherwise, if *vnamevar* is set to a scalar 0, then the variable names for the output will be generated automatically ("X1,X2,...,XK") unless the alternate variable names are set explicitly in the *alt-names* member of the **dstatmtControl** structure.

If pairwise deletion is used, the minima and maxima will be the true values for the valid data. The means and standard deviations will be computed using the correct number of valid observations for each variable.

Example

In order to create a real, working example that you can use, you must first create a sample **GAUSS Data Archive** with the code below.

```
//Create an example GAUSS Data Archive
ret = gdaCreate("myfile.gda",1);

//Add a variable 'A' which is a 10x5 random normal matrix
ret = gdaWrite("myfile.gda",rndn(10,5),"A");

//Add a variable 'COLS' which is a 5x1 string array
string vnames = { "X1", "X2", "X3", "X4", "X5" };
ret = gdaWrite("myfile.gda", vnames, "COLS");
```

gdaDStatMat

This code above will create a **GAUSS Data Archive** containing two variables, the **GAUSS** matrix *A* containing the data and *COLS* which contains the names for the columns of the matrix *A* which are the model variables (*X1*, *X2*,...).

The code below computes the statistics on each of the columns of the matrix *A*.

```
#include dstatmt.sdf
struct dstatmtControl dc0;
struct dstatmtout dout;

dc0 = dstatmtControlCreate;
colind = { 1, 2, 3, 4, 5 };
dout = gdaDStatMat(dc0, "myfile.gda", "A", colind, "COLS"
);
```

The final input to **gdaDStatMat** above tells the function the names to use for the columns of *A*. In this example, you can reference the *COLS* variable by name as you see in the example below. Alternatively, you can access this variable by index. Since *COLS* is the second variable in the **GAUSS Data Archive** created at the start of this example, the following is equivalent to the last line above:

```
dout = gdaDStatMat(dc0, "myfile.gda", "A", colind, 2 );
```

If you wanted to calculate the statistics on just the first, third and fifth columns of *A*:

```
colind = { 1, 3, 5 };
dout = gdaDStatMat(dc0, "myfile.gda", "A", colind, "COLS"
);
```

Notice in these lines above that *COLS* still contains all of the variable names i.e. *X1*, *X2*, *X3*, *X4* and *X5*. *COLS* should always contain the full list of all variables in the matrix *A*.

Source

gdadstat.src

See Also

[gdaDStat](#), [dstatmtControlCreate](#)

gdaGetIndex

Purpose

Gets the index of a variable in a **GAUSS** Data Archive.

Format

```
ind = gdaGetIndex(filename, varname);
```

Input

<i>filename</i>	string, name of data file.
<i>varname</i>	string, name of variable in the GDA.

Output

<i>ind</i>	scalar, index of variable in the GDA.
------------	---------------------------------------

Remarks

If **gdaGetIndex** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

- | | |
|---|------------------|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |

gdaGetName

5	Invalid file type.
8	Variable not found.
10	File contains no variables.
14	File too large to be read on current platform.

Example

```
ind = gdaGetIndex("myfile.gda", "observed");
```

See Also

[gdaGetName](#), [gdaReadByIndex](#)

gdaGetName

Purpose

Gets the name of a variable in a **GAUSS** Data Archive.

Format

```
varname = gdaGetName(filename, varind);
```

Input

<i>filename</i>	string, name of data file.
<i>varind</i>	scalar, index of variable in the GDA.

Output

<i>varname</i>	string, name of variable in the GDA.
----------------	--------------------------------------

Remarks

If **gdaGetName** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

- | | |
|---|---------------------|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |
| 5 | Invalid file type. |
| 8 | Variable not found. |

Example

```
varname = gdaGetName("myfile.gda",5);
```

See Also

[gdaGetIndex](#), [gdaRead](#), [gdaGetNames](#)

gdaGetNames

Purpose

Gets the names of all the variables in a **GAUSS** Data Archive.

Format

```
varnames = gdaGetNames(filename);
```

Input

<i>filename</i>	string, name of data file.
-----------------	----------------------------

gdaGetOrders

Output

<i>varnames</i>	Nx1 string array, names of all the variables in the GDA.
-----------------	--

Remarks

If **gdaGetNames** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

- | | |
|----|--|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |
| 5 | Invalid file type. |
| 10 | File contains no variables. |
| 13 | Result too large for current platform. |
| 14 | File too large to be read on current platform. |

Example

```
varnames = gdaGetNames("myfile.gda");
```

See Also

[gdaGetTypes](#), [gdaGetName](#)

gdaGetOrders

Purpose

Gets the orders of a variable in a **GAUSS** Data Archive.

Format

```
ord = gdaGetOrders(filename, varname);
```

Input

<i>filename</i>	string, name of data file.
<i>varname</i>	string, name of variable in the GDA.

Output

<i>ord</i>	Mx1 vector, orders of the variable in the GDA.
------------	--

Remarks

If the specified variable is a matrix or string array, then *ord* will be a 2x1 vector containing the rows and columns of the variable respectively. If the variable is a string, then *ord* will be a scalar containing the length of the string. If the variable is an N-dimensional array, then *ord* will be an Nx1 vector containing the sizes of each dimension.

If **gdaGetOrders** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

- | | |
|----|--|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |
| 5 | Invalid file type. |
| 8 | Variable not found. |
| 10 | File contains no variables. |
| 14 | File too large to be read on current platform. |

gdaGetType

Example

```
ord = gdaGetOrders ("myfile.gda", "x5");
```

See Also

[gdaGetName](#), [gdaGetIndex](#)

gdaGetType

Purpose

Gets the type of a variable in a **GAUSS** Data Archive.

Format

```
vartype = gdaGetType(filename, varname);
```

Input

<i>filename</i>	string, name of data file.
<i>varname</i>	string, name of variable in the GDA.

Output

<i>vartype</i>	scalar, type of the variable in the GDA.
----------------	--

Remarks

vartype may contain any of the following:

6	Matrix
---	--------

13	String
15	String array
21	Array

If **gdaGetType** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

1	Null file name.
2	File open error.
4	File read error.
5	Invalid file type.
8	Variable not found.
10	File contains no variables.
14	File too large to be read on current platform.

Example

```
vartype = gdaGetType("myfile.gda", "x1");
```

See Also

[gdaGetTypes](#)

gdaGetTypes

Purpose

Gets the types of all the variables in a **GAUSS** Data Archive.

gdaGetTypes

Format

```
vartypes = gdaGetTypes(filename);
```

Input

<i>filename</i>	string, name of data file.
-----------------	----------------------------

Output

<i>vartypes</i>	Nx1 vector, types of all the variables in the GDA.
-----------------	--

Remarks

vartypes may contain any of the following:

6	Matrix
13	String
15	String array
21	Array

If **gdaGetTypes** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. Valid error codes for this command include:

1	Null file name.
2	File open error.
4	File read error.
5	Invalid file type.
10	File contains no variables.
14	File too large to be read on current platform.

Example

```
vartypes = gdaGetTypes("myfile.gda");
```

See Also

[gdaGetNames](#), [gdaRead](#)

gdaGetVarInfo

Purpose

Gets information about all of the variables in a **GAUSS** Data Archive and returns it in an array of **gdavartable** structures.

Include

gdafns.sdf

Format

```
vtab = gdaGetVarInfo(filename);
```

Input

<i>filename</i>	string, name of data file.
-----------------	----------------------------

Output

<i>vtab</i>	Nx1 array of gdavartable structures, where N is the number of variables in <i>filename</i> , containing the following members: <i>vtab</i> string, name of variable.
-------------	--

gdaGetVarInfo

<i>[i].name</i>	
<i>vtab</i>	scalar, type of variable.
<i>[i].type</i>	
<i>vtab</i>	Mx1 vector or scalar, orders of the
<i>[i].orders</i>	variable.

Remarks

The size of *vtab.orders* is dependent on the type of the variable as follows:

Variable Type	<i>vtab.orders</i>
array	Mx1 vector, where M is the number of dimensions in the array, containing the sizes of each dimension, from the slowest-moving dimension to the fastest-moving dimension.
matrix	2x1 vector containing the rows and columns of the matrix, respectively.
string	scalar containing the length of string, excluding the null terminating byte.
string array	2x1 vector containing the rows and columns of the string array, respectively.
<i>vtab.type</i> may contain any of the following:	
6	matrix

13	string
15	string array
21	array

Example

```
//Execute structure definition
#include gdafns.sdf
struct gdavartable vtab;

vtab = gdaGetVarInfo("myfile.gda");
```

Source

gdafns.src

See Also

[gdaReportVarInfo](#), [gdaGetNames](#), [gdaGetTypes](#), [gdaGetOrders](#)

gdaIsCplx

Purpose

Checks to see if a variable in a **GAUSS** Data Archive is complex.

Format

```
y = gdaIsCplx(filename, varname);
```

gdaLoad

Input

<i>filename</i>	string, name of data file.
<i>varname</i>	string, name of variable in the GDA.

Output

<i>y</i>	scalar, 1 if variable is complex; 0 if real.
----------	--

Remarks

If **gdaIsCplx** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. Valid error codes for this command include:

- | | |
|----|--|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |
| 5 | Invalid file type. |
| 8 | Variable not found. |
| 10 | File contains no variables. |
| 14 | File too large to be read on current platform. |

Example

```
cplx = gdaIsCplx("myfile.gda","x1");
```

gdaLoad

Purpose

Loads variables in a GDA into the workspace.

Format

```
ret = gdaLoad(filename, create, modify, rename, ftypes,  
errh, report);
```

Input

<i>filename</i>	string, name of data file.
<i>create</i>	scalar, create flag: <i>0</i> do not create any new variables in the workspace. <i>1</i> create new variables in the workspace.
<i>modify</i>	scalar, modify flag: <i>0</i> do not modify any variables in the workspace. <i>1</i> if the name of a variable in the data file matches the name of a variable already in the workspace, modify that variable.
<i>rename</i>	scalar, rename flag: <i>0</i> do not rename a variable retrieved from the data file when copying it into the workspace. <i>1</i> rename variables retrieved from the data file when copying them into the workspace if there are name conflicts with existing variables, which may not be modified.
<i>ftypes</i>	scalar, type force flag: <i>0</i> do not force a type change on any

gdaLoad

g

		variables in the workspace when modifying.
	1	force a type change on a variable in the workspace when modifying it with the data in a variable of the same name in the data file. Note that if <i>ftypes</i> is set to 1, gdaLoad will follow regular type change rules. The types of sparse matrix and structure variables will NOT be changed.
<i>errh</i>		scalar, controls the error handling of gdaLoad :
	0	skip operations that cannot be performed, without setting an error return.
	1	return an error code if operations are skipped.
	2	terminate program if operations are skipped.
<i>report</i>		scalar, controls reporting:
	0	no reporting.
	1	report only name changes and operations that could not be performed.
	2	report type changes, name changes, and operations that could not be performed.
	3	report everything.

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one
------------	---

of the following error codes:

4	File read error.
5	Invalid file type.
10	File contains no variables.
14	File too large to be read on current platform.
24	Variables skipped.
26	Cannot add structure definition.
27	Structure definition does not match.

Remarks

For each variable in *filename*, **gdaLoad** will first compare the name of the variable against the names of the variables already resident in the **GAUSS** workspace to see if there is a match. If there is not a match, and *create* is set to 1, it will create a new variable. Otherwise if *create* is set to 0, it will skip that variable.

If the variable name does match that of a variable already resident in the **GAUSS** workspace, and *modify* is set to 1, it will attempt to modify that variable. If the types of the two variables are different, and *ftype* is set to 1, it will force the type change if possible and modify the existing variable.

If it cannot modify the variable or *modify* is set to 0, it will check to see if *rename* is set to 1, and if so, attempt to rename the variable, appending an *_ num* to the variable name, beginning with *num* = 1 and counting upward until it finds a name with which there are no conflicts. If the variable cannot be modified and *rename* is set to 0, then the variable will be skipped.

The *rename* argument also controls the handling of structure definitions. If a structure variable is encountered in the GDA file, and no variable of the same name exists in the workspace (or the variable is renamed), **gdaLoad** will attempt to find a

gdaPack

structure definition in the workspace that matches the one in the GDA. Note that in order for structure definitions to match, the structure definition names must be the same as well as the number, order, names, and types of their members.

If no matching structure definition is found, the definition in the file will be loaded into the workspace. If there is already a non-matching structure definition with the same name in the workspace and *rename* is set to 1, then **gdaLoad** will attempt to rename the structure definition, using the same method as it does for variable names.

If a structure variable is encountered in the GDA file, a structure variable of the same name already exists in the workspace, and *modify* is set to 1, then **gdaLoad** will modify the existing variable, providing that the structure definitions of the two variables match.

Example

```
ret = gdaLoad("myfile.gda",1,1,1,1,1,3);
```

This example loads the variables in `myfile.gda` into the workspace, creating a new variable if a variable of the same name does not already exist, modifying an existing variable if a variable of the same name does already exist and the modification does not result in an impossible type change, and renaming the variable if none of the above is possible. The example returns an error code if any variables in `myfile.gda` are skipped and reports all activity.

See Also

[gdaSave](#)

gdaPack

Purpose

Packs the data in a **GAUSS** Data Archive, removing all empty bytes and truncating the file.

Format

```
ret = gdaPack(filename);
```

Input

<i>filename</i>	string, name of data file.
-----------------	----------------------------

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes:
1	Null file name.
2	File open error.
3	File write error.
4	File read error.
5	Invalid data file type.
10	File contains no variables.
12	File truncate error.
14	File too large to be read on current platform.

Remarks

You may want to call **gdaPack** after several calls to **gdaUpdate** to remove all of the empty bytes from a GDA.

Example

```
ret = gdaPack("myfile.gda");
```

gdaRead

See Also

[gdaUpdate](#), [gdaWrite](#)

gdaRead

Purpose

Gets a variable from a **GAUSS** Data Archive.

Format

```
y = gdaRead(filename, varname);
```

Input

<i>filename</i>	string, name of data file.
<i>varname</i>	string, name of variable in the GDA.

Output

<i>y</i>	matrix, array, string or string array, variable data.
----------	---

Remarks

If **gdaRead** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

- | | |
|---|--------------------|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |
| 5 | Invalid file type. |

8	Variable not found.
10	File contains no variables.
14	File too large to be read on current platform.

Example

```
y = gdaRead("myfile.gda", "x1");
```

See Also

[gdaReadByIndex](#), [gdaGetName](#)

gdaReadByIndex

Purpose

Gets a variable from a **GAUSS** Data Archive given a variable index.

Format

```
y = gdaReadByIndex(filename, varind);
```

Input

<i>filename</i>	string, name of data file.
<i>varind</i>	scalar, index of variable in the GDA.

Output

<i>y</i>	matrix, array, string or string array, variable data.
----------	---

gdaReadSome

Remarks

If **gdaReadByIndex** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

- | | |
|----|-----------------------------|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |
| 5 | Invalid file type. |
| 8 | Variable not found. |
| 10 | File contains no variables. |

Example

```
y = gdaReadByIndex("myfile.gda",3);
```

See Also

[gdaRead](#), [gdaGetIndex](#)

gdaReadSome

Purpose

Reads part of a variable from a **GAUSS** Data Archive.

Format

```
y = gdaReadSome(filename, varname, index, orders);
```

Input

<i>filename</i>	string, name of data file.
<i>varname</i>	string, name of variable in the GDA.
<i>index</i>	scalar or Nx1 vector, index into variable where read is to begin.
<i>orders</i>	scalar or Kx1 vector, orders of object to output.

Output

<i>y</i>	matrix, array, string or string array, variable data.
----------	---

Remarks

This command reads part of the variable *varname* in *filename*, beginning at the position indicated by *index*. The *orders* argument determines the size and shape of the object outputted by **gdaReadSome**. The number of elements read equals the product of all of the elements in *orders*.

If *index* is a scalar, it will be interpreted as the *index*th element of the variable. Thus if *varname* references a 10x5 matrix, an *index* of 42 would indicate the 42nd element, which is equivalent to the [8,2] element of the matrix (remember that **GAUSS** matrices are stored in row major order). If *index* is an Nx1 vector, then N must equal the number of dimensions in the variable referenced by *varname*.

If *orders* is a Kx1 vector, then *y* will be a K-dimensional object. If *orders* is a scalar *r*, then *y* will be an rx1 column vector. To specify a 1xr row vector, set *output* = { 1, *r* }.

If the variable referenced by *varname* is numeric (a matrix or array) and *orders* is a scalar or 2x1 vector, then *y* will be of type matrix. If the variable is numeric and *orders* is an Nx1 vector where N>2, then *y* will be of type array.

gdaReadSome

If *varname* references a string, then both *index* and *orders* must be scalars, and *index* must contain an index into the string in characters.

If **gdaReadSome** fails, it will return a scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

- | | |
|----|--|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |
| 5 | Invalid file type. |
| 8 | Variable not found. |
| 10 | File contains no variables. |
| 13 | Result too large for current platform. |
| 14 | File too large to be read on current platform. |
| 15 | Argument out of range. |
| 18 | Argument wrong size. |

Example

```
x = rndn(100,50);
ret = gdaCreate("myfile.gda",1);
ret = gdaWrite("myfile.gda",x,"x1");

index = { 35,20 };
orders = { 25,5 };
y = gdaReadSome("myfile.gda","x1",index,orders);
```

This example reads $25 \times 5 = 125$ elements from *x1*, beginning with the [35,20] element. The 125 elements are returned as a 25×5 matrix, *y*.

See Also

[gdaWriteSome](#), [gdaRead](#)

gdaReadSparse

Purpose

Gets a sparse matrix from a **GAUSS** Data Archive.

Format

```
sm = gdaReadSparse(filename, varname);
```

Input

<i>filename</i>	string, name of data file.
<i>varname</i>	string, name of sparse matrix variable in the GDA.

Output

<i>sm</i>	sparse matrix.
-----------	----------------

Remarks

If **gdaReadSparse** fails, it will return a sparse scalar error code. Call **scalerr** to get the value of the error code. The error code may be any of the following:

- | | |
|----|--|
| 1 | Null file name. |
| 2 | File open error. |
| 4 | File read error. |
| 5 | Invalid file type. |
| 8 | Variable not found. |
| 10 | File contains no variables. |
| 14 | File too large to be read on current platform. |

gdaReadStruct

Example

```
sparse matrix sm1;  
sm1 = gdaReadSparse("myfile.gda", "sm");
```

See Also

[gdaRead](#), [gdaReadStruct](#), [gdaWrite](#)

gdaReadStruct

Purpose

Gets a structure from a GAUSS Data Archive.

Format

```
{ instance, retcode } = gdaReadStruct(filename, varname,  
structure_type);
```

Input

<i>filename</i>	string, name of data file.
<i>varname</i>	string, name of structure instance in the GDA.
<i>structure_type</i>	string, structure type.

Output

<i>instance</i>	instance of the structure.
<i>retcode</i>	scalar, 0 if successful, otherwise, any of the following error codes:

1	Null file name.
2	File open error.
4	File read error.
5	Invalid file type.
8	Variable not found.
10	File contains no variables.
14	File too large to be read on current platform.

Remarks

instance can be an array of structures.

Example

```
struct mystruct {  
    matrix x;  
    array a;  
};  
  
struct mystruct msw;  
msw.x = rdn(500,25);  
msw.a = areshape(rdn(5000,100),10|500|100);  
ret = gdaCreate("myfile.gda",1);  
ret = gdaWrite("myfile.gda",msw,"ms");  
struct mystruct msr;  
{ msr, ret } = gdaReadStruct("myfile.gda","ms","mystruct");
```

See Also

[gdaRead](#), [gdaReadSparse](#), [gdaWrite](#)

gdaReportVarInfo

gdaReportVarInfo

Purpose

Gets information about all of the variables in a **GAUSS** Data Archive and returns it in a string array formatted for printing.

Format

```
vinfo = gdaReportVarInfo(filename);
```

Input

<i>filename</i>	string, name of data file.
-----------------	----------------------------

Output

<i>vinfo</i>	Nx1 string array containing variable information.
--------------	---

Remarks

If you just want to print the information to the window, call **gdaReportVarInfo** without assigning the output to a symbol name:

```
gdaReportVarInfo(filename);
```

Example

```
x1 = rndn(100,50);  
x2 = rndn(75,5);  
a = areshape(rndn(10000,1),10|100|10);  
fname = "myfile.gda";  
ret = gdaCreate(fname,1);
```



```
ret = gdaWrite(fname,x1,"x1");  
ret = gdaWrite(fname,x2,"x2");  
ret = gdaWrite(fname,a,"a1");  
gdaReportVarInfo(fname);
```

produces:

Index	Name	Type	cOrders
1	x1	matrix	100x50
2	x2	matrix	75x5
3	a1	array	10x100x10

Source

gdafns.src

See Also

[gdaGetVarInfo](#), [gdaGetNames](#), [gdaGetTypes](#), [gdaGetOrders](#)

gdaSave

Purpose

Writes variables in a workspace to a GDA.

Format

```
ret = gdaSave(filename, varnames, exclude, overwrite,  
report);
```

Input

<i>filename</i>	string, name of data file.
<i>varnames</i>	string or NxK string array, names of variables in the workspace to include or exclude.
<i>exclude</i>	scalar, include/exclude flag: <div><div>0</div>include all variables contained in <i>varnames</i>. <div>1</div>exclude all variables contained in <i>varnames</i>.</div>
<i>overwrite</i>	scalar, controls the overwriting of the file and variables in the file: <div><div>0</div>if file exists, return with an error code. <div>1</div>if file exists, overwrite completely. <div>2</div>if file exists, append to file, appending to variable names if necessary to avoid name conflicts. <div>3</div>if file exists, update file. When a name conflict occurs, update the existing variable in the file with the new variable.</div>
<i>report</i>	scalar, controls reporting: <div><div>0</div>no reporting. <div>1</div>report only name changes (note that name changes occur only when <i>overwrite</i> is set to 2). <div>3</div>report everything.</div>

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes:
1	Null file name.
3	File write error.
4	File read error.
5	Invalid file type.
6	File exists and <i>overwrite</i> set to 0.
7	Cannot create file.
14	File too large to be read on current platform.
16	Cannot write to GDA - version outdated.
17	Type mismatch.

Remarks

Only initialized variables are written to the GDA with **gdaSave**.

If *varnames* is a null string and *exclude* is set to 0, it will be interpreted as indicating all of the variables in the workspace.

You may add an asterisk (*) to the end of a variable name in *varnames* to indicate that all variables beginning with the specified text are to be selected. For example, setting *varnames* to the string "*_* *" and setting *exclude* to 1 indicates that all variables EXCEPT those starting with an underscore should be written to the GDA.

The names of the variables in the workspace are the names that are given to the variables when they are written to the GDA, with the exception of names that are changed to avoid conflicts.

gdaUpdate

If you set `overwrite` to 2, and variable name conflicts are encountered, **gdaSave** will append an underscore and a number to the name of the variable it is adding. It will first try changing the name to `name_1`. If there is a conflict with that name, it will change it to `name_2`, and so on until it finds a name that does not conflict with any of the variables already in the GDA.

Example

```
run -r myfile.gau;  
ret = gdaSave("myfile.gda", "x*", 0, 2, 3);
```

This example runs a **GAUSS** program called `myfile.gau` and then writes all initialized variables in the workspace beginning with 'x' to the file `myfile.gda`. If `myfile.gda` already exists, this example appends to it, changing the names of the variables that it writes to the file if necessary to avoid name conflicts. All writing and variable name changing is reported.

See Also

[gdaLoad](#)

gdaUpdate

Purpose

Updates a variable in a **GAUSS** Data Archive.

Format

```
ret = gdaUpdate(filename, x, varname);
```

Input

<i>filename</i>	string, name of data file.
-----------------	----------------------------

<i>x</i>	matrix, array, string or string array, data.
<i>varname</i>	string, variable name.

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes:
1	Null file name.
2	File open error.
3	File write error.
4	File read error.
5	Invalid data file type.
8	Variable not found.
10	File contains no variables.
14	File too large to be read on current platform.

Remarks

This command updates the variable *varname* in *filename* with the data contained in *x*.

If *x* is larger than the specified variable in the file, then **gdaUpdate** writes the new variable data after the last variable in the data file, moving the variable descriptor table to make room for the data and leaving empty bytes in the place of the old variable. This does not change the index of the variable because variable indices are determined NOT by the order of the variable data in a GDA, but by the order of the variable descriptors.

gdaUpdateAndPack

If *x* is the same size or smaller than the specified variable in the file, then **gdaUpdate** writes the data in *x* over the specified variable. If *x* is smaller, then **gdaUpdate** leaves empty bytes between the end of the updated variable and the beginning of the next variable in the data file.

This command updates variables quickly by not moving data in the file unnecessarily. However, calling **gdaUpdate** several times for one file may result in a file with a large number of empty bytes. To pack the data in a GDA, so it contains no empty bytes, call **gdaPack**. Or to update a variable without leaving empty bytes in the file, call **gdaUpdateAndPack**.

Example

```
x = rndn(100,50);  
ret = gdaCreate("myfile.gda",1);  
ret = gdaWrite("myfile.gda",x,"x1");  
  
y = rndn(75,5);  
ret = gdaUpdate("myfile.gda",y,"x1");
```

See Also

[gdaUpdateAndPack](#), [gdaPack](#), [gdaWrite](#)

gdaUpdateAndPack

Purpose

Updates a variable in a **GAUSS** Data Archive, leaving no empty bytes if the updated variable is smaller or larger than the variable it is replacing.

Format

```
ret = gdaUpdateAndPack(filename, x, varname);
```

Input

<i>filename</i>	string, name of data file.
<i>x</i>	matrix, array, string or string array, data.
<i>varname</i>	string, variable name.

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes: <div><div>1</div><div>Null file name.</div></div> <div><div>2</div><div>File open error.</div></div> <div><div>3</div><div>File write error.</div></div> <div><div>4</div><div>File read error.</div></div> <div><div>5</div><div>Invalid data file type.</div></div> <div><div>8</div><div>Variable not found.</div></div> <div><div>10</div><div>File contains no variables.</div></div> <div><div>12</div><div>File truncate error.</div></div> <div><div>14</div><div>File too large to be read on current platform.</div></div>
------------	---

Remarks

This command updates the variable *varname* in *filename* with the data contained in *x*. **gdaUpdateAndPack** always writes the data in *x* over the specified variable in the file. If *x* is larger than the specified variable, then it first moves all subsequent data in the file to make room for the new data. If *x* is smaller, then **gdaUpdateAndPack** writes the data, packs all of the subsequent data, leaving no empty bytes after the updated variable, and truncates the file.

gdaVars

This command uses disk space efficiently; however, it may be slow for large files (especially if the variable to be updated is one of the first variables in the file). If speed is a concern, you may want to use **gdaUpdate** instead.

Example

```
x = rndn(100,50);  
ret = gdaCreate("myfile.gda",1);  
ret = gdaWrite("myfile.gda",x,"x1");  
  
y = rndn(75,5);  
ret = gdaUpdateAndPack("myfile.gda",y,"x1");
```

See Also

[gdaUpdate](#), [gdaWrite](#)

gdaVars

Purpose

Gets the number of variables in a **GAUSS** Data Archive.

Format

```
nvars = gdaVars(filename);
```

Input

<i>filename</i>	string, name of data file.
-----------------	----------------------------

Output

<i>nvars</i>	scalar, the number of variables in <i>filename</i> .
--------------	--

Example

```
nvars = gdaVars("myfile.gda");
```

Source

gdafns.src

See Also

[gdaReportVarInfo](#), [gdaGetNames](#)

gdaWrite

Purpose

Writes a variable to a **GAUSS** Data Archive.

Format

```
ret = gdaWrite(filename, x, varname);
```

Input

<i>filename</i>	string, name of data file.
<i>x</i>	matrix, array, string or string array, data to write to the GDA.
<i>varname</i>	string, variable name.

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes:
------------	---

gdaWrite32

1	Null file name.
2	File open error.
3	File write error.
4	File read error.
5	Invalid data file type.
9	Variable name too long.
11	Variable name must be unique.
14	File too large to be read on current platform.

Remarks

gdaWrite adds the data in *x* to the end of the variable data in *filename*, and gives the variable the name contained in *varname*.

Example

```
x = rndn(100,50);
ret = gdaCreate("myfile.gda",1);
ret = gdaWrite("myfile.gda",x,"x1");
```

See Also

[gdaWrite32](#), [gdaCreate](#)

gdaWrite32

Purpose

Writes a variable to a **GAUSS** Data Archive using 32-bit system file write commands.

Format

```
ret = gdaWrite32(filename, x, varname);
```

Input

<i>filename</i>	string, name of data file.
<i>x</i>	matrix, array, string or string array, data to write to the GDA.
<i>varname</i>	string, variable name.

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes:
1	Null file name.
2	File open error.
3	File write error.
4	File read error.
5	Invalid data file type.
9	Variable name too long.
11	Variable name must be unique.
14	File too large to be read on current platform.
25	Not supported for use with a file created on a machine with a different byte order.

Remarks

```
gdaWrite32 adds the data in x to the end of the variable data in filename, and
```

gdaWriteSome

gives the variable the name contained in *varname*.

This command is a speed optimization command for Windows. On all other platforms, this function is identical to **gdaWrite**. **gdaWrite** uses system file write commands that support 64-bit file sizes. These commands are slower on Windows XP than the 32-bit file write commands that were used for binary writes in **GAUSS** 6.0 and earlier.

gdaWrite32 uses the 32-bit Windows system write commands, which will be faster on Windows XP. Note, however, that **gdaWrite32** does not support 64-bit file sizes.

This command does not support writing to a GDA that was created on a platform with a different byte order than the current machine. **gdaWrite** supports full cross-platform writing to GDA's.

Example

```
x = rndn(100,50);  
ret = gdaCreate("myfile.gda",1);  
ret = gdaWrite32("myfile.gda",x,"x1");
```

See Also

[gdaWrite](#), [gdaCreate](#)

gdaWriteSome

Purpose

Overwrites part of a variable in a **GAUSS** Data Archive.

Format

```
ret = gdaWriteSome(filename, x, varname, index);
```

Input

<i>filename</i>	string, name of data file.
<i>x</i>	matrix, array, string or string array, data.
<i>varname</i>	string, variable name.
<i>index</i>	scalar or Nx1 vector, index into variable where new data is to be written.

Output

<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes:
1	Null file name.
2	File open error.
3	File write error.
4	File read error.
5	Invalid data file type.
8	Variable not found.
10	File contains no variables.
14	File too large to be read on current platform.
15	Argument out of range.
17	Type mismatch.
18	Argument wrong size.
19	Data must be real.
20	Data must be complex.

Remarks

This command overwrites part of the variable *varname* in *filename* with the

gdaWriteSome

data contained in *x*. The new data is written to *varname* beginning at the position indicated by *index*.

If *index* is a scalar, it will be interpreted as the *index*th element of the variable. Thus if *varname* references a 10x5 matrix, an *index* of 42 would indicate the 42nd element, which is equivalent to the [8,2] element of the matrix (remember that **GAUSS** matrices are stored in row major order). If *index* is an Nx1 vector, then N must equal the number of dimensions in the variable referenced by *varname*.

If *varname* references a string, then *index* must be a scalar containing an index into the string in characters.

gdaWriteSome may not be used to extend the size of a variable in a GDA. If there are more elements (or characters for strings) in *x* than there are from the indexed position of the specified variable to the end of that variable, then **gdaWriteSome** will fail. Call **gdaAppend** to append data to an existing variable.

The shape of *x* need not match the shape of the variable referenced by *varname*. If *varnum* references an NxK matrix, then *x* may be any LxM matrix (or P-dimensional array) that satisfies the size limitations described above. If *x* contains R elements, then the elements in *x* will simply replace the indexed element of the specified variable and the subsequent R-1 elements (as they are laid out in memory).

If *varname* references a string array, then the size of the overall variable will change if the sum of the length of the string array elements in *x* is different than the sum of the length of the elements that they are replacing.

In this case, if the variable increases in size, then the variable data will be rewritten after the last variable in the data file, moving the variable descriptor table to make room for the data and leaving empty bytes in its old location. This does not change the index of the variable because variable indices are determined NOT by the order of the variable data in a GDA, but by the order of the variable descriptors. If the variable decreases in size, then **gdaWriteSome** leaves empty bytes between the end of the variable and the beginning of the next variable in the data file. Call **gdaPack** to pack the data in a GDA, so it contains no empty bytes.

Example

```
x = rndn(100,50);
ret = gdaCreate("myfile.gda",1);
ret = gdaWrite("myfile.gda",x,"x1");

y = rndn(75,5);
index = { 52, 4 };
ret = gdaWriteSome("myfile.gda",y,"x1",index);
```

This example replaces $75 \times 5 = 375$ elements in $x1$, beginning with the [52,4] element, with the elements in y .

See Also

[gdaReadSome](#), [gdaUpdate](#), [gdaWrite](#)

getarray

Purpose

Gets a contiguous subarray from an N-dimensional array.

Format

```
y = getarray(a, loc);
```

Input

<i>a</i>	N-dimensional array.
<i>loc</i>	Mx1 vector of indices into the array to locate the subarray of interest, where $1 \leq M \leq N$.

getdims

Output

y [N-M]-dimensional subarray or scalar.

Remarks

If N-M>0, **getarray** will return an array of [N-M] dimensions, otherwise, if N-M=0, it will return a scalar.

Example

```
a = seqa (1,1,720) ;  
a = areshape (a,2|3|4|5|6) ;  
loc = { 2,1 } ;  
y = getarray (a,loc) ;
```

y will be a 4x5x6 array of sequential values, beginning at [1,1,1] with 361, and ending at [4,5,6] with 480.

See Also

[getmatrix](#)

getdims

Purpose

Gets the number of dimensions in an array, matrix, string array or other symbol.

Format

```
ndims = getdims(a);
```


Input

<i>a</i>	N-dimensional array, matrix, string, string array or sparse matrix.
----------	---

Output

<i>ndims</i>	scalar, the number of dimensions in the symbol.
--------------	---

Example

```
//Create a 2x120x12 random normal array
nelems = 2 * 120 * 12;
a = areshape(rndn(nelems, 1), 2|120|12);

//Find ut the number of dimensions in 'a'
ndims = getdims(a);
```

The code above, assigns *ndims* to be equal to 3.

See Also

[getorders](#)

getf

Purpose

Loads an ASCII or binary file into a string.

getf

Format

```
y = getf(filename, mode);
```

Input

<i>filename</i>	string, any valid file name.
<i>mode</i>	scalar 1 or 0 which determines if the file is to be loaded in ASCII mode (0) or binary mode (1).

Output

<i>y</i>	string containing the file.
----------	-----------------------------

Remarks

If the file is loaded in ASCII mode, it will be tested to see if it contains any end of file characters. These are ^Z (ASCII 26). The file will be truncated before the first ^Z, and there will be no ^Z's in the string. This is the correct way to load most text files because the ^Z's can cause problems when trying to print the string to a printer.

If the file is loaded in binary mode, it will be loaded just like it is with no changes.

Example

Suppose you have a file which writes the results of its calculations to a file in a report format. For this example, we will use the code snippet below:

```
x1 = rndn(100,5);  
y1 = rndu(100,1);  
  
output file = regression_results.txt reset;  
call ols("", y1, x1);  
output off;
```

```
x2 = rndn(100,5);  
y2 = rndu(100,1);  
  
output file = ols_results.txt reset;  
call ols("", y2, x2);  
output off;
```

Running the code above will create a file named "regression_results.txt" and a file named "ols_results.txt" in your current working directory. You can retrieve the output from either of these files with the **getf** command.

```
str = getf("regression_results.txt",1);  
print str;
```

You can take this further and create a procedure that will load a list of output files for you. It can then print the output from each file as you are ready to read it.

```
declare string array fileList = { "regression_results.txt",  
"ols_results.txt" };  
showOutput(fileList);  
  
proc (0) = showOutput(fileList);  
    local k;  
    for i(1, rows(fileList), 1);  
        print "Press any key to view the next file:";  
        //wait for user input and assign the first key stroke  
        //to 'k'  
        k = keyw;  
        print getf(fileList[i],1);  
    endfor;  
endp;
```

getGAUSSHome

See Also

[load](#), [save](#), [let](#), [con](#)

getGAUSSHome

Purpose

Returns the full path to the GAUSS home directory..

Format

```
g_home = getGAUSSHome();
```

Output

g_home string, full path to **GAUSS** home directory.

Remarks

getGAUSSHome can be used to load and save files in directories that are relative to the **GAUSS** installation directory without knowing its exact location ahead of time. This can help to make programs that will run on multiple computers, using different paths without any changes to the code.

Example

Example 1: locate the GAUSS home directory

If you installed GAUSS in the directory, C:\gauss:

```
g_home = getGAUSSHome ();  
print g_home;
```

produces:

```
C:\gauss
```

Example 2: loading a file from the GAUSS examples directory

```
//Create full path to dataset
f_name = getGAUSSHome() $+ "examples/fueleconomy.dat";

//Load the dataset
fuel_economy = loadadd(f_name);
```

getmatrix

Purpose

Gets a contiguous matrix from an N-dimensional array.

Format

```
y=getmatrix(a, loc);
```

Input

<i>a</i>	N-dimensional array.
<i>loc</i>	Mx1 vector of indices into the array to locate the matrix of interest, where M equals N, N-1 or N-2.

Output

<i>y</i>	KxL or 1xL matrix or scalar, where L is the size of the fastest moving dimension of the array and K is the size of the second fastest moving dimension.
----------	---

getmatrix

Remarks

Inputting an $N \times 1$ locator vector will return a scalar, an $(N-1) \times 1$ locator vector will return a $1 \times L$ matrix, and an $(N-2) \times 1$ locator vector will return a $K \times L$ matrix.

Example

```
//Create the sequence 1, 2, 3...20
a = seqa(1, 1, 20);

//Reshape the column vector 'a' into a 3x3x2 dimensional
//array
a = areshape(a, 3|3|2);

//Extract the second 3x2 array
mat = getmatrix(a, 2);
```

After code above *a* is equal to:

```
Plane [1,...]

      1.0000000      2.0000000
      3.0000000      4.0000000
      5.0000000      6.0000000

Plane [2,...]

      7.0000000      8.0000000
      9.0000000     10.000000
     11.000000     12.000000

Plane [3,...]

     13.000000     14.000000
```

15.000000	16.000000
17.000000	18.000000

and *mat* is equal to:

7.0000000	8.0000000
9.0000000	10.000000
11.000000	12.000000

See Also

[getarray](#), [getmatrix4D](#)

getmatrix4D

Purpose

Gets a contiguous matrix from a 4-dimensional array.

Format

y = **getmatrix4D**(*a*, *i1*, *i2*);

Input

- | | |
|-----------|--|
| <i>a</i> | 4-dimensional array. |
| <i>i1</i> | scalar, index into the slowest moving dimension of the array. |
| <i>i2</i> | scalar, index into the second slowest moving dimension of the array. |

getmatrix4D

Output

y

KxL matrix, where L is the size of the fastest moving dimension of the array and K is the size of the second fastest moving dimension.

Remarks

getmatrix4D returns the contiguous matrix that begins at the [*i1*,*i2*,1,1] position in array *a* and ends at the [*i1*, *i2*,K,L] position.

A call to **getmatrix4D** is faster than using the more general **getmatrix** function to get a matrix from a 4-dimensional array, especially when *i1* and *i2* are the counters from nested **for** loops.

Example

```
//Create a column vector 1, 2, 3...120
a = seqa(1,1,120);

//Reshape the column vector into a 2x3x4x5 dimensional
//array
a = areshape(a,2|3|4|5);

//Extract a submatrix
y = getmatrix4D(a,2,3);
```

After the code above:

	101	102	103	104	105
y =	106	107	108	109	110
	111	112	113	114	115
	116	117	118	119	120

See Also

[getmatrix](#), [getscalar4D](#), [getarray](#)

getname

Purpose

Returns a column vector containing the names of the variables in a **GAUSS** data set.

Format

```
y = getname(dset);
```

Input

<i>dset</i>	string specifying the name of the data set from which the function will obtain the variable names.
-------------	--

Output

<i>y</i>	Nx1 vector containing the names of all of the variables in the specified data set.
----------	--

Remarks

The output, *y*, will have as many rows as there are variables in the data set.

Example

```
y = getname(getGaussHome $+ "examples/freqdata.dat");
```

getnamef

```
format 8,8;  
print $y;
```

produces:

```
AGE  
PAY  
sex  
WT
```

The above example assumes that the data set `olsdat` contains the variables: *TIME*, *DIST*, *TEMP*, *FRICT*.

Note that the extension is not included in the filename passed to the **getname** function.

See Also

[getnamef](#), [indcv](#)

getnamef

Purpose

Returns a string array containing the names of the variables in a **GAUSS** data set.

Format

```
y = getnamef(f);
```

Input

<i>f</i>	scalar, file handle of an open data set
----------	---

Output

y	Nx1 string array containing the names of all of the variables in the specified data set.
-----	--

Remarks

The output, y , will have as many rows as there are variables in the data set.

Example

```
file = getgausshome()$+ "examples/freqdata.dat";  
//Open the dataset  
open f = ^file for read;  
  
//Create a string array with the variable names from the  
//dataset  
y = getnamef(f);  
  
//Check which variables are character and which are numeric  
t = vartypef(f);  
  
print y;
```

produces:

```
AGE  
PAY  
sex  
WT
```

getNextTradingDay

The above example assumes that the data set `freqdata` contains the variables: *AGE*, *PAY*, *sex*, *WT*.

Note the use of **vartypef** to determine the types of these variables.

See Also

[getname](#), [indcv](#), [vartypef](#)

getNextTradingDay

Purpose

Returns the next trading day.

Format

```
n = getNextTradingDay(a);
```

Input

<i>a</i>	scalar, date in DT scalar format.
----------	-----------------------------------

Output

<i>n</i>	scalar, next trading day in DT scalar format.
----------	---

Remarks

A trading day is a weekday that is not a holiday as defined by the New York Stock Exchange from 1888 through 2006. Holidays are defined in `holidays.asc`. You may edit that file to modify or add holidays.

Source

`finutils.src`

Globals

`_fin_holidays`

See Also

[getPreviousTradingDay](#), [annualTradingDays](#)

getNextWeekDay**Purpose**

Returns the next day that is not on a weekend.

Format

```
n = getNextWeekDay(a);
```

Input

<code>a</code>	scalar, date in DT scalar format.
----------------	-----------------------------------

Output

<code>n</code>	scalar, next week day in DT scalar format.
----------------	--

Source

`finutils.src`

getnr

See Also

[getPreviousWeekDay](#)

getnr

Purpose

Computes number of rows to read per iteration for a program that reads data from a disk file in a loop.

Format

```
nr = getnr(nsets, ncols);
```

Input

<i>nsets</i>	scalar, estimate of the maximum number of duplicate copies of the data matrix read by readr to be kept in memory during each iteration of the loop.
<i>ncols</i>	scalar, columns in the data file.

Output

<i>nr</i>	scalar, number of rows readr should read per iteration of the read loop.
-----------	---

Remarks

If `__row` is greater than 0, *nr* will be set to `__row`.

If an insufficient memory error is encountered, change `__rowfac` to a number less than 1.0 (e.g., 0.75). The number of rows read will be reduced in size by this factor.

Source

gauss.src

Globals

__row, __rowfac, __maxvec

getnrmt

Purpose

Computes number of rows to read per iteration for a program that reads data from a disk file in a loop.

Format

```
nr=getnr(nsets, ncols, row, rowfac, maxv);
```

Input

<i>nsets</i>	scalar, estimate of the maximum number of duplicate copies of the data matrix read by readr to be kept in memory during each iteration of the loop.
<i>ncols</i>	scalar, columns in the data file.
<i>row</i>	scalar, if row is greater than 0, <i>nr</i> will be set to <i>row</i> .
<i>rowfac</i>	scalar, <i>nr</i> will be reduced in size by this factor. If insufficient memory error is encountered, change this to a number less than one (e.g., 0.9).
<i>maxv</i>	scalar, the largest number of elements allowed in any one matrix.

getorders

Output

nr

scalar, number of rows **readr** should read per iteration of the read loop.

Source

`gaussmt.src`

getorders

Purpose

Returns a vector containing the size of the dimensions of an array, matrix, or other symbol. sss

Format

```
orders = getorders(a);
```

Input

a

N-dimensional array, matrix, sparse matrix, string or string array.

Output

orders

Nx1 vector of orders, the sizes of the dimensions of the array.

Example

Example 1

```
//Allocate a 5x100x3 dimensional array with each element
```



```
equal to 0.  
a = arrayinit(5|100|3,0);  
  
//Find the size of the dimensions in 'a'  
orders = getorders(a);
```

After the code above:

```
orders =    5  
          100  
          3
```

Example 2

```
//Create 121x4 random matrix  
x = rndn(121, 4);  
  
//Get the number of rows and columns of the matrix  
dims = getorders(x);
```

After the above code, **dims** should equal:

```
121  
4
```

See Also

[getdims](#), [rows](#), [cols](#)

getpath

Purpose

Returns an expanded filename including the drive and path.

getpath

Format

```
fname = getpath(pfname);
```

Input

<i>p</i> fname	string, partial filename with only partial or missing path information.
----------------	---

Output

<i>f</i> name	string, filename with full drive and path.
---------------	--

Remarks

This function handles relative path references.

Example

```
y = getpath("temp.e");  
print y;
```

produces:

```
C:\gauss\temp.e
```

assuming that C:\gauss is the current directory.

Source

getpath.src

getPreviousTradingDay

Purpose

Returns the previous trading day.

Format

```
n = getPreviousTradingDay(a);
```

Input

<i>a</i>	scalar, date in DT scalar format.
----------	-----------------------------------

Output

<i>n</i>	scalar, previous trading day in DT scalar format.
----------	---

Remarks

A trading day is a weekday that is not a holiday as defined by the New York Stock Exchange from 1888 through 2006. Holidays are defined in `holidays.asc`. You may edit that file to modify or add holidays.

Source

`finutils.src`

Globals

`_fin_holidays`

See Also

[getNextTradingDay](#)

getPreviousWeekDay

getPreviousWeekDay

Purpose

Returns the previous day that is not on a weekend.

Format

```
n = getPreviousWeekDay(a);
```

Input

<i>a</i>	scalar, date in DT scalar format.
----------	-----------------------------------

Output

<i>n</i>	scalar, previous week day in DT scalar format.
----------	--

Source

finutils.src

See Also

[getNextWeekDay](#)

getRow

Purpose

Returns a specified row from a matrix.

Format

```
y = getRow(a, row);
```

Input

<i>a</i>	NxK matrix
<i>row</i>	The row of the matrix to extract.

Output

<i>y</i>	A 1xK row vector.
----------	-------------------

Remarks

getRow is designed to give an alternative access to rows in a matrix than indexing the matrix by brackets.

Example

First create a matrix, *a*:

```
a = randn(10,10);
```

Now you can assign a variable *y* to be equal the third row of *a* with either of the following statements.

```
y = getRow(a,3);
```

or

```
y = a[3,:];
```

While both statements will produce the same result, the first may make for code that is easier to read and interpret.

getscalar3D

See Also

[CR-getTrRow](#)

getscalar3D

Purpose

Gets a scalar from a 3-dimensional array.

Format

```
y = getscalar3D(a, i1, i2, i3);
```

Input

<i>a</i>	3-dimensional array.
<i>i1</i>	scalar, index into the slowest moving dimension of the array.
<i>i2</i>	scalar, index into the second slowest moving dimension of the array.
<i>i3</i>	scalar, index into the fastest moving dimension of the array.

Output

<i>y</i>	scalar, the element of the array indicated by the indices.
----------	--

Remarks

getscalar3D returns the scalar that is located in the [*i1*, *i2*, *i3*] position of array *a*.

A call to **getscalar3D** is faster than using the more general **getmatrix** function to get a scalar from a 3-dimensional array.

Example

```
//Create a column vector 1, 2, 3,...24
a = seqa(1,1,24);

//Reshape the column vector into a 2x3x4 dimensional array
a = areshape(a,2|3|4);

y = getscalar3D(a,1,3,2);
```

A 2x3x4 dimensional array can be thought of as two 3x4 dimensional matrices. The call to **getScalar3D** above, returns the [3,2] element of the first of these matrices. The value of which is:

```
y = 10
```

See Also

[getmatrix](#), [getscalar4D](#), [getarray](#)

getscalar4D

Purpose

Gets a scalar from a 4-dimensional array.

Format

```
y=getscalar4D(a, i1, i2, i3, i4);
```

getscalar4D

Input

<i>a</i>	4-dimensional array.
<i>i1</i>	scalar, index into the slowest moving dimension of the array.
<i>i2</i>	scalar, index into the second slowest moving dimension of the array.
<i>i3</i>	scalar, index into the second fastest moving dimension of the array.
<i>i4</i>	scalar, index into the fastest moving dimension of the array.

Output

<i>y</i>	scalar, the element of the array indicated by the indices.
----------	--

Remarks

getscalar4D returns the scalar that is located in the [*i1*, *i2*, *i3*, *i4*] position of array *a*.

A call to **getscalar4D** is faster than using the more general **getmatrix** function to get a scalar from a 4-dimensional array.

Example

```
a = seqa(1,1,120);  
a = areshape(a,2|3|4|5);  
y = getscalar4D(a,1,3,2,5);
```

The code above assigns *y* equal to 50.

See Also

[getmatrix](#), [getscalar3D](#), [getarray](#)

getTrRow

Purpose

Transposes a matrix and then returns a single row from it.

Format

```
y = getTrRow(a, row);
```

Input

<i>a</i>	NxK matrix
<i>row</i>	The row of the matrix to extract.

Output

<i>y</i>	A 1xK row vector.
----------	-------------------

Remarks

getRow is designed to give an alternative access to rows in a matrix than indexing the matrix by brackets.

Example

```
a = rndn(10,10);
```

getwind

```
y = getTrRow(a, 3);
```

See Also

[getRow](#)

getwind

Purpose

Retrieve the current graphic panel number. Note: This function is for use with the deprecated PQG graphics.

9

Library

pgraph

Format

```
n = getwind;
```

Output

n scalar, graphic panel number of current graphic panel.

Remarks

The current graphic panel is the graphic panel in which the next graph will be drawn.

Source

pwindow.src

See Also

[endwind](#), [begwind](#), [window](#), [setwind](#), [nextwind](#)

glm

Purpose

Solves the generalized linear model problems

Format

```
out = glm(y, x, family);
out = glm(y, x, family, var_names);
out = glm(y, x, family, var_names, categoryIdx);
out = glm(y, x, family, var_names, categoryIdx, link);
out = glm(y, x, family, ctl);
or
out = glm(dataset_name, formula, family);
out = glm(dataset_name, formula, family, ctl);
```

Input

y	$n \times 1$ vector, the dependent, or response, variable. n is the number of the observations used in the analysis.
x	$n \times k$ matrix, the independent, or explanatory, variables. k is the number of the independent variables.
$dataset_name$	String, the name of data set. E.g. "credit.dat" or "example.fmt".
$formula$	String, formula string of the model. E.g "y ~ X1 + X2", 'y' is the name of dependent

<i>family</i>	<p>variable, 'X1' and 'X2' are names of independent variables;</p> <p>E.g. $y \sim .$, '.' means including all variables except dependent variable 'y';</p> <p>E.g. $y \sim -1 + X1 + X2$, '-1' means no intercept model.</p> <p>String, the distribution of the dependent variable. Options include:</p> <ul style="list-style-type: none"> • "binomial" • "gamma" • "normal" • "poisson" • "inverse gaussian"
<i>var_names</i>	<p>Optional argument, $(k + 1) \times 1$ string array or character matrix, the names of the variables. The first element must be the name of the dependent variable.</p> <p>e.g., <code>var_names = "admit" \$ "gre" \$ "gpa" \$ "rank"</code>, then "admit" will be the label of the response variable, "gre", "gpa", "rank" are the labels of the independent variables corresponding to the order in the X matrix.</p>
<i>categoryIdx</i>	<p>Optional argument, $1 \times k_d$ matrix, $k_d \leq k$. k_d is the categorical variable index of X matrix.</p> <p><i>categoryIdx</i> specifies the categorical variable columns to be used in the analysis.</p> <p>e.g. If <i>categoryIdx</i> = 0, then it means the independent variable does not contain any categorical</p>

variables;

if $categoryIdx = \{ 1\ 4 \}$, then it means that column 1 and column 4 in the X matrix are categorical variables.

Note: The function **glm** uses the smallest number as the reference category in each categorical variable.

link

String, the link function. Options include:

- "identity"
- "inverse"
- "inverse squared"
- "ln"
- "logit"
- "probit"
- "cloglog"
- "canonical"

The default link of each distribution is the canonical link function: Normal -- identity; Binomial -- logit; Gamma -- inverse; Poisson -- nature log.

ctl

An instance of a **glmControl** structure. For an instance named *ctl*, the members are:

ctl.varNames

$(k + 1) \times 1$

string array or character matrix, the names of the variables. The first

ctl.categoryIdx

element must be the name of the dependent variable.

$1 \times k_d$ matrix,
 $k_d \leq k$.

ctl.categoryIdx specifies the categorical variable columns to be used in the analysis.

e.g. If

ctl.categoryIdx = 0, then it means no categorical variable; if

ctl.categoryIdx = { 1 4}, then it means that column 1 and column 4 in x matrix are categorical variables.

Note: glm function uses the smallest number as the reference

ctl.link

category in each categorical variable.

String, the link function. Options include:

- "identity"
- "inverse"
- "inverse squared"
- "ln"
- "logit"
- "probit"
- "cloglog"
- "canonical"

The default link is the canonical link for each distribution.

ctl.constantFlag

Scalar, flag of constant term. The negative number means no intercept model, e.g. "-1". This member will be ignored if a [formula string](#) is

<code>ctl.printFlag</code>	used. String, "Y" or "N", flag of print to screen. The "N" means no printing.
<code>ctl.maxIters</code>	Scalar, maximum iterations. The default <code>ctl.maxIters</code> is 25.
<code>ctl.eps</code>	Scalar, convergence precision. The default is 1e-8.

Output

<code>out</code>	An instance of a glmOut structure. For an instance named <code>out</code> , the members are:	
<code>out.modelInfo</code>	An instance of a glmModelInfo structure. The members are:	
	<code>out.modelInfo.distribution</code>	string, the distribution of dependent variable
	<code>out.modelInfo.link</code>	string, the

	link function used in the procedure
<i>out.modelInfo.yName</i>	string, the label of dependent variable
<i>out.modelInfo.xNames</i>	string array, the label of independent variables with intercept and dummy variables for each categorical variable
<i>out.modelInfo.varNames</i>	string array, the label of variables

<code>out.modelInfo.n</code>	scalar, the number of valid cases used in the analysis
<code>out.modelInfo.df</code>	scalar, degree of freedom
<code>out.modelSelect</code>	An instance of a glmModelSelection structure. The members are:
<code>out.modelSelect.deviance</code>	scalar, the residual deviance from the fit model. The greater the deviance, the poorer the fit.
<code>out.modelSelect.pearson</code>	scalar, the Pearson Chi-square Statistics. Pearson statistic is an

alternative to the deviance for testing the fit of certain GLMs.

`out.modelSelect.LL`

scalar, the log likelihood of the fit model

`out.modelSelect.dispersion`

scalar, the estimate of the dispersion parameter by Pearson statistic and degree of freedom. It is fixed at 1 when the distribution is "poisson" or

6

		"binomial".
	<i>out.modelSelect.aic</i>	scalar, Akaike information criterion (AIC)
	<i>out.modelSelect.bic</i>	scalar, Bayesian information criterion (BIC)
<i>out.coef</i>	An instance of a glmParameters structure. The members are:	
	<i>out.coef.estimates</i>	matrix, the estimate value of parameters
	<i>out.coef.se</i>	matrix, the standard error of parameters
	<i>out.coef.testStat</i>	matrix, the statistic

		value of parameters
	<i>out.coef.testStatName</i>	string, the name of test statistic
	<i>out.coef.pvalue</i>	scalar, the p_value of parameters
<i>out.yhat</i>	scalar, the fitted mean values for response variable	
<i>out.residuals</i>	matrix, residuals on the linear predictor scale, equal to the adjusted response value minus the fitted linear predictors	
<i>out.covmat</i>	matrix, the covariance matrix for the parameters	
<i>out.corrmat</i>	matrix, the correlation matrix for the parameters	
<i>out.constantFlag</i>	String, flag of constant term.	
<i>out.iteration</i>	scalar, the number of iterations of IWLS used	

glm

<code>out.maxIters</code>	scalar, the maximum iterations
<code>out.eps</code>	scalar, convergence precision

Examples

Example 1

Ordinary linear regression with simulated data sets.

```
new;
cls;

//Set random number seed for repeatable random numbers
rndseed 86;

//Simulate data using rndn function
x = rndn(100,4);
y = rndn(100,1);

//Call glm function with the minimum inputs
call glm(y, x, "normal");
```

This example will compute a least squares regression of y on x . The results will be shown in the program input / output window. The return values are discarded by using a `call` statement.

Generalized Linear Model		
Valid cases:	100	Dependent Variable:
	y	
Degrees of freedom:	95	Distribution:

```

              normal
Deviance:              99.37      Link function:
              identity
Pearson Chi-square:    99.37      AIC:
              295.2
Log likelihood:        -141.6     BIC:
              310.8
Dispersion:            1.046     Iterations:
              2

```

Variable		Estimate	Standard Error
t-value	> t		
-----	-----	-----	----
CONSTANT		0.067084	0.10233
0.65556	0.513692		
x1		-0.027278	0.097162
-0.28074	0.779517		
x2		-0.10747	0.090888
-1.1825	0.239963		
x3		0.27659	0.093397
2.9615	0.00386701		
x4		0.067915	0.11099
0.6119	0.542062		

Example 2

Ordinary linear regression with categorical variables.

```

new;
cls;

//Create filename with full path

```

```

dataset = getGAUSSHome() $+ "examples/credit.dat";

//Import data
data = loadadd(dataset);

//Select the independent variables by index
x = data[:, 1 7 9] ;

//Select the dependent variable by index
y = data[:, 11];

//Get the labels of the variables with getname function
vnames = getname(dataset);
label = vnames[ 11 1 7 9, 1 ];

//Categorical variable index
categoryIdx = { 2 3 };

//Call glm function with three necessary inputs and two
optional inputs
call glm(y, x, "normal", label, categoryIdx);

```

In this example, the data set "**credit.dat**" is used to compute a ordinary linear regression.

The dependent variable is "**Balance**". The independent variables are: "**Married**", "**Gender**" and "**Income**".

vnames is a character vector from the "**credit.dat**" by using **getname** function. *label* is the labels used in the regression, the first element must be the label of dependent variable, the rest members are the labels for the independent variables corresponding to the order in the *x* matrix.

"**Gender**" and "**Married**" are categorical variables. The **glm** choose the smallest number(1) as the base category in each categorical variable. The following shows the output:

Generalized Linear Model

Valid cases:	400	Dependent Variable:
Balance		
Degrees of freedom:	396	Distribution:
normal		
Deviance:	6.611e+007	Link function:
identity		
Pearson Chi-square:	6.611e+007	AIC:
5951		
Log likelihood:	-2971	BIC:
5971		
Dispersion:	1.669e+005	Iterations:
2		

Variable		Estimate	Standard Error
t-value	> t		
-----	-----	-----	----
CONSTANT		246.19	46.535
5.2903	< 0.0001		
Gender	2	24.577	40.889
0.60108	0.548134		
Married	2	-21.279	41.963
-0.50708	0.612383		
Income		6.0626	0.58077
10.439	< 0.0001		

Example 3

Logistic regression with categorical variables.

```
new;
cls;
```

```
// Import data with csvReadM and csvReadSA functions
// The second input, '2', means that read the data from the
second row
file = getGAUSShome() $+ "examples/binary.csv";
data = csvReadM(file, 2);

// Import labels of variables
// The second input specify the range of rows, '1|1' means
from Row 1 to Row 1, only the first row.
vnames = csvReadSA(file, 1|1) ;

// Row vector -> column vector
vnames = vnames';

// Index dependent variable
y = data[:,1];

// Index independent variables
x = data[:,2:4];

// Index categorical variable in x matrix
categoryIdx = 3;

// Call glm function with three necessary inputs and two
dynamic inputs
call glm(y,x, "binomial",vnames, categoryIdx);
```

In this example, the data set "binary.csv" is used to compute a logistic regression.

The dependent variable is "admit". The independent variables are: "gre", "gpa" , and "rank".

data is a numeric matrix from the "binary.csv" by using **csvReadM** function with starting at the second row.

`vnames` is a string array from the "binary.csv" by using `csvReadSA` function. `vnames` is used in the regression as the labels for all variables, the first element must be the label of dependent variable, the rest members are the labels for the independent variables corresponding to the order in the `x` matrix.

`categoryIdx` specify the category column in the `x` matrix, "rank". "rank" is a categorical variable with 4 levels(1, 2, 3, 4). 1 is chosen as the base category automatically since 1 is the smallest number in "rank".

`link` specifies the link function to use in the model. If you specify the `link = ""` or didn't specify a link function, then the default canonical link function is used correspond to the "binomial" distribution. The followings are the output:

Generalized Linear Model

Valid cases:	admit	400	Dependent Variable:
Degrees of freedom:	binomial	394	Distribution:
Deviance:	logit	458.5	Link function:
Pearson Chi-square:		397.5	AIC:
		470.5	
Log likelihood:		-229.3	BIC:
		494.5	
Dispersion:		1	Iterations:
		4	

Variable	Estimate	Standard Error
z-value	> z	
-----	-----	-----
-----	-----	---
CONSTANT	-3.99	1.14

glm

```
-3.5001      0.000465027
rank         2      -0.67544      0.31649
-2.1342      0.0328288
          3      -1.3402      0.34531      -3.8812
0.000103942
          4      -1.5515      0.41783      -3.7131
0.000204711
gre              0.0022644      0.001094
  2.0699      0.0384651
gpa              0.80404      0.33182
  2.4231      0.0153879
```

Note: Dispersion parameter for BINOMIAL distribution taken to be 1

Example 4

Using **glmControl** and **glmOut** structure to run a Logistic regression with categorical variables.

```
new;
cls ;
//Create file name with full path
file = getGAUSShome() %+ "examples/binary.csv";

data = csvReadM(file, 2);

// Import labels of variables
// The second input specify the range of rows, '1|1' means
// from Row 1 to Row 1, only the first row.
vnames = csvReadSA(file, 1|1) ;

// Index dependent variable
y = data[:,1];
```

```

// Index independent variables
x = data[:,2:4];

// Declare 'binary_ctl' as a glmControl structure
struct glmControl binary_ctl;

// Specify the variable names
binary_ctl.varNames = vnames';

// Index categorical variable in x matrix
binary_ctl.categoryIdx = 3;

// Specify the link function
binary_ctl.link = "probit";

// Save out the results in glmOut structure
struct glmOut out1;
out1 = glm(y, x, "binomial", binary_ctl);

```

In this example, the data set "binary.csv" is used to compute a logistic regression. The dependent variable is "admit". The independent variables are: "gre", "gpa", and "rank"

binary_ctl is a **glmControl** structure. *binary_ctl.varNames* must be a string array. In this example, the first element is the "insurance", the rest members are the "gre", "gpa", and "rank" correspond to the variable order in the *x* matrix.

binary_ctl.categoryIdx specify the categorical variable in *x* matrix. "rank" is a categorical variable with 3 levels(1, 2, 3). The number 1 is chosen as the base category automatically since 1 is the smallest number in "rank".

binary_ctl.link specifies the link function to use in the model. In this model, we use "probit" as the link function. If you specify the *binary_ctl.link* = "" or

glm

didn't specify a link function, then the default canonical link function is used correspond to the "binomial" distribution.

`out1` is a **glmOut** structure. All the results are saved in the `out1`.

After running above code, the output is:

```
Generalized Linear Model

Valid cases:          400      Dependent Variable:
                        admit
Degrees of freedom:   394      Distribution:
                        binomial
Deviance:             458.4    Link function:
                        probit
Pearson Chi-square:   397.7    AIC:
                        470.4
Log likelihood:       -229.2   BIC:
                        494.4
Dispersion:           1       Iterations:
                        4

Variable              Estimate      Standard
z-value              >|z|          Error
-----
-----
CONSTANT              -2.3868      0.67395
-3.5416      0.000397733
rank              2      -0.4154      0.19498
-2.1305      0.0331297
              3      -0.81214      0.20836      -
3.8978      < 0.0001
              4      -0.9359      0.24527      -
3.8158      0.000135764
```

```

gre                0.0013756      0.00065003
  2.1162           0.0343292
gpa                0.47773        0.1972
  2.4226           0.0154097

```

Note: Dispersion parameter for BINOMIAL distribution taken to be 1

Example 5

A Poisson regression model with categorical variables.

```

new;
cls;

// Import data with loadm function

fname = getGAUSShome() $+ "examples/poisson_sim.fmt";
loadm data = ^fname;

// Index dependent variable, 'num_award'
y = data[:,2];

// Index independent variable, 'prog' and 'math'
x = data[:, 3 4];

// Specify the variable names
string var_names = {"num_award", "prog", "math"};

// Index categorical variable in x matrix
categoryIdx = 1;

// specify the link function, 'ln'

```

glm

```
link = "ln";

// Declare the glmOut structure
// All the results are saved in the out_poi
struct glmOut out_poi;
out_poi = glm(y, x, "poisson", var_names,categoryIdx ,link)
;
```

After running above code, the output is:

Generalized Linear Model			
Valid cases:	200	Dependent Variable:	
num_award			
Degrees of freedom:	196	Distribution:	
poisson			
Deviance:	189.4	Link function:	
ln			
Pearson Chi-square:	212.1	AIC:	
373.5			
Log likelihood:	-182.8	BIC:	
386.7			
Dispersion:	1	Iterations:	
6			
		Standard	
Variable	Estimate	Error	
z-value	> z		
-----	-----	-----	---
-----	-----		
CONSTANT	-5.2471	0.65845	
-7.9689	< 0.0001		
prog	1.0839	0.35825	
3.0254	0.00248303		

	3	0.36981	0.44107	0.83844
0.401786				
math		0.070152	0.010599	
6.6186	< 0.0001			

Note: Dispersion parameter for POISSON distribution taken to be 1

Example 6

Using **glmOut** structure to save result for a Gamma regression with categorical variables.

```
new;
cls;

// File name with full path
file = getGAUSShome() $+ "examples/yarn.xlsx";

//Read 4th column as a numeric matrix
y = xlsReadM(file, "D2:D28");

//Read columns 1, 2 and 3 as character data
x = xlsReadSA(file, "A2:C28");

//Find unique categorical levels
from = uniquesa(x[:,1]);

//Numeric categorical levels
to = {1, -1, 0};

//Reclassify the character to number
x = reclassify(x, from, to);
```

glm

```
//Declare 'ctl_gamma' as a glmControl struct
struct glmControl ctl_gamma;

//Read variable names and transpose
//to a column vector
ctl_gamma.varNames = xlsReadSA(file, "A1:D1" );

//Specify categorical columns
ctl_gamma.categoryIdx = { 1 2 3 };

//Specify link function
ctl_gamma.link = "ln";

//Declare 'out_gamma' to be a glmOut structure
struct glmOut out_gamma;

//Call 'glm' and fill 'out_gamma' with results
out_gamma = glm(y,x,"gamma",ctl_gamma );
```

In this example, the data set "yarn.xlsx" is used to perform a Gamma regression.

After running the code above, the output is :

Generalized Linear Model

Valid cases:	27	Dependent Variable:
yarn_length		
Degrees of freedom:	20	Distribution:
gamma		
Deviance:	0.7089	Link function:
ln		
Pearson Chi-square:	0.6917	AIC:
336.5		
Log likelihood:	-160.3	BIC:

```

Dispersion:      346.9      0.03458      Iterations:
                  5

```

Variable		Estimate	Standard Error	
t-value	> t			
-----	-----	-----	-----	---
CONSTANT		6.4841	0.09469	
68.477	< 0.0001			
amplitude	0	0.9136	0.087666	
10.421	< 0.0001			
	1	1.6791	0.087666	19.153
< 0.0001				6
load	0	-0.64738	0.087666	
-7.3846	< 0.0001			
	1	-1.2654	0.087666	-14.435
< 0.0001				
cycles	0	-0.31872	0.087666	
-3.6356	0.00164628			
	1	-0.7701	0.087666	-8.7844
< 0.0001				

Example 7

Using a **"*.dat"** file directly in **glm** for a Inverse Gaussian distribution.

```

new;
cls;

// File name with full path
fname = getGAUSShome() $+ "examples/clotting_time.dat";

```

glm

```
//Declare 'fit_inv' to be a glmOut structure
struct glmOut fit_inv;

//Call 'glm' and fill 'fit_inv' with results
fit_inv = glm(fname, "plasma ~ lot1", "inverse gaussian");
```

After running the code above, the output is :

9

Generalized Linear Model			
Valid cases:		9	Dependent Variable:
plasma			
Degrees of freedom:		7	Distribution:
inverse gaussian			
Deviance:		0.03557	Link function:
inverse squared			
Pearson Chi-square:		0.03511	AIC:
71.1			
Log likelihood:		-32.55	BIC:
71.69			
Dispersion:		0.005016	Iterations:
6			
Variable	Estimate	Standard Error	
t-value	> t		
-----	-----	-----	-----
-----	-----		
CONSTANT	-0.0034177	0.00074729	
-4.5735	0.00256355		

lot1	0.00019223	4.0768e-05
4.7154	0.00216923	

Example 8

Running a no intercept model.

```
new;
cls;

// File name with full path
fname = getGAUSShome() $+ "examples/credit.dat";

//Declare 'fit' to be a glmOut structure
struct glmOut fit;

//Call 'glm' with no intercept model
fit = glm(fname, "Balance ~ -1 + Income", "normal");
```

After running the code above, the output is :

Generalized Linear Model		
Valid cases:	400	Dependent Variable:
Balance		
Degrees of freedom:	399	Distribution:
normal		
Deviance:	7.538e+07	Link function:
identity		
Pearson Chi-square:	7.538e+07	AIC:
5998		

```

Log likelihood:          -2997      BIC:
                        6006
Dispersion:             1.889e+05    Iterations:
                        2

```

Variable	Estimate	Standard Error
t-value	> t	
-----	-----	-----
Income	9.4429	0.37925
24.899	< 0.0001	

Example 9

Running a linear regression model with HDF5 file.

```

new;
cls;

// Give a fully pathed HDF5 file name
file_name = getGAUSShome() $+ "examples/nba_data.h5";

// Given a data set name in above file
dname = "/nba_data";

// Give the name of attributes -- "headers"
// All variable names must be saved as "headers" in the
// data set for hdf5 file
attname = "headers";

// Read variable names
vname = h5readAttribute(file_name, dname, attname);

// Print variable names of the "nba_data" data set

```

```
print vname;

// Define the formula for the linear model
formula = "Weight ~ Height + Age";

// Specify data set name with schema "h5://"
datasetname = "h5://" $+ file_name $+ dname;

//Call 'glm'
call glm(datasetname, formula, "normal");
```

After running the code above, the output is :

```
Height
Weight
Age

Generalized Linear Model

Valid cases:          505      Dependent Variable:
                        Weight
Degrees of freedom:    502      Distribution:
                        normal
Deviance:              1.128e+05  Link function:
                        identity
Pearson Chi-square:    1.128e+05  AIC:
                        4173
Log likelihood:        -2082     BIC:
                        4189
Dispersion:           224.7     Iterations:
                        2
```

glm

Variable		Estimate	Standard Error
t-value	> t		
-----	-----	-----	----
-----	-----		
CONSTANT		-296.6	15.804
-18.768	< 0.0001		
Height		6.3267	0.1933
32.729	< 0.0001		
Age		0.6511	0.15436
4.2181	< 0.0001		

Remarks

1. The **glmControl** structure stores the user defined options.
2. The **glmOut** structure stores all the results after running **glm** function.
3. For the categorical variables, **glm** chooses the smallest value as the base category. You can change the base category by using the **reclassify** or **recode** functions to change the base category with the smallest value in the variable.
4. The dispersion parameter is calculated based on Pearson Chi-square Statistics.
5. The **glm** function cannot handle missing values. You can use **packr** function to delete the rows of a matrix that contain any missing values.
6. The weights for each observation are equal.
7. The supported data set types are [CSV](#), [XLS](#), [XLSX](#), [HDF5](#), [FMT](#), [DAT](#).
For HDF5 file, the *dataset* must include [file schema](#) and both file name and data set name must be provided, e.g. **glm**
("h5://C:/gauss17/examples/testdata.h5/mydata", *formula*, *family*).

Source

glm.src

See Also

[ols](#), [olsmt](#), [reclassify](#), [packr](#), [Formula String](#)

gosub

Purpose

Causes a branch to a subroutine. Note: This is an advanced function that gives extra flexibility for sophisticated users in some circumstances. In most cases, it is preferable to create a procedure ([proc](#)).

Format

```
gosub label;  
.  
.  
.  
label:  
.  
.  
.  
return;
```

Remarks

For multi-line recursive user-defined functions, see **PROCEDURES AND KEYWORDS**, CHAPTER 1.

When a [gosub](#) statement is encountered, the program will branch to the label and begin executing from there. When a [return](#) statement is encountered, the program will resume executing at the statement following the [gosub](#) statement. Labels are 1-

gosub

32 characters long and are followed by a colon. The characters can be A-Z or 0-9, but they must begin with an alphabetic character. Uppercase or lowercase is allowed.

It is possible to pass parameters to subroutines and receive parameters from them when they return. See the second example, following.

The only legal way to enter a subroutine is with a `gosub` statement.

If your subroutines are at the end of your program, you should have an `end` statement before the first one to prevent the program from running into a subroutine without using a `gosub`. This will result in a Return without gosub error message.

The variables used in subroutines are not local to the subroutine and can be accessed from other places in your program. (See **PROCEDURES AND KEYWORDS**, CHAPTER 1.)

9

Example

In the program below the name *mysub* is a label. When the `gosub` statement is executed, the program will jump to the label *mysub* and continue executing from there. When the `return` statement is executed, the program will resume executing at the statement following the `gosub`.

```
x = rndn(3,3);
z = 0;
gosub mysub;
print z;
end;

/* ----- Subroutines Follow ----- */

mysub:
  z = inv(x);
  return;
```

Parameters can be passed to subroutines in the following way (line numbers are added for clarity):

```

1. gosub mysub(x,y);
2. pop j; /* b will be in j */
3. pop k; /* a will be in k */
4. t = j*k;
5. print t;
6. end;
7.
8. /* ---- Subroutines Follow ---- */
9.
10. mysub:
11. pop b; /* y will be in b */
12. pop a; /* x will be in a */
13.
14. a = inv(b)*b+a;
15. b = a'b;
16. return(a,b);

```

In the above example, when the `gosub` statement is executed, the following sequence of events results (line numbers are included for clarity):

1. x and y are pushed on the stack and the program branches to the label *mysub* in line 10.
11. the second argument that was pushed, y , is `pop`'ped into b .
12. the first argument that was pushed, x , is `pop`'ped into a .
14. `inv(b)*b+a` is assigned to a .
15. $a'b$ is assigned to b .
16. a and b are pushed on the stack and the program branches to the statement following the `gosub`, which is line 2.
2. the second argument that was pushed, b , is `pop`'ped into j .
3. the first argument that was pushed, a , is `pop`'ped into k .
4. $j*k$ is assigned to t .

goto

5. t is printed.
6. the program is terminated with the `end` statement.

Matrices are pushed on a last-in/first-out stack in the `gosub()` and `return()` statements. They must be `pop`'ped off in the reverse order. No intervening statements are allowed between the label and the `pop` or the `gosub` and the `pop`. Only one matrix may be `pop`'ped per `pop` statement.

See Also

[goto](#), [proc](#), [pop](#), [return](#)

goto

Purpose

Causes a branch to a label.

Format

```
goto label;  
.  
.  
.  
label:
```

Remarks

Label names can be any legal **GAUSS** names up to 32 alphanumeric characters, beginning with an alphabetic character or an underscore, not a reserved word.

Labels are always followed immediately by a colon.

Labels do not have to be declared before they are used. **GAUSS** knows they are labels by the fact that they are followed immediately by a colon.

When **GAUSS** encounters a `goto` statement, it jumps to the specified label and continues execution of the program from there.

Parameters can be passed in a `goto` statement the same way as they can with a `gosub`.

Example

```
x = sega (.1, .1, 5);  
n = { 1 2 3 };  
goto fip;  
print x;  
end;  
  
fip:  
print n;
```

produces:

```
1.0000000 2.0000000 3.0000000
```

See Also

[gosub](#), [if](#)

gradMT

Purpose

Computes numerical gradient.

gradMT

Include

optim.sdf

Format

```
g = gradMT(&fct, par1, data1);
```

Input

<i>&fct</i>	scalar, pointer to procedure returning either Nx1 vector or 1x1 scalar.
<i>par1</i>	an instance of structure of type PV containing parameter vector at which gradient is to be evaluated.
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i> .

Output

<i>g</i>	NxK Jacobian or 1xK gradient.
----------	-------------------------------

Remarks

par1 must be created using the **pvPack** procedures.

Example

```
#include optim.sdf
struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1, 0.1|0.2, "P");

struct DS d0;
```

```
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);

proc fct(struct PV p0, struct DS d0);
    local p,y;
    p = pvUnpack(p0, "P");
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    retp(y);
endp;

g = gradMT(&fct,p1,d0);
```

Source

gradmt.src

gradMTm

Purpose

Computes numerical gradient with mask.

Include

optim.sdf

Format

```
g=gradMTm(&fct,par1,data1, mask);
```

Input

<i>&fct</i>	scalar, pointer to procedure returning either Nx1 vector or 1x1 scalar.
-----------------	---

gradMTm

<i>par1</i>	an instance of structure of type PV containing parameter vector at which gradient is to be evaluated.
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i> .
<i>mask</i>	Kx1 matrix, elements in <i>g</i> corresponding to elements of <i>mask</i> set to zero are not computed, otherwise they are computed.

Output

<i>g</i>	NxK Jacobian or 1xK gradient.
----------	-------------------------------

Remarks

par1 must be created using the **pvPack** procedures.

Example

```
#include optim.sdf
struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1, 0.1|0.2, "P");

struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1, 1, 15);

proc fct(struct PV p0, struct DS d0);
    local p, y;
    p = pvUnpack(p0, "P");
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    retp(y);
endp;
```



```
mask = { 0, 1 };  
g = gradMTm(&fct,p1,d0,mask);
```

Source

gradmt.src

gradMTT

Purpose

Computes numerical gradient using available threads.

Include

optim.sdf

Format

```
g = gradMTT(&fct,par1,data1);
```

Input

<i>fct</i>	scalar, pointer to procedure returning either Nx1 vector or 1x1 scalar.
<i>par1</i>	structure of type PV containing parameter vector at which gradient is to be evaluated
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i>

gradMTT

Output

g

NxK Jacobian or 1xK gradient

Remarks

par1 must be created using the **pvPack** procedures

Example

```
#include optim.sdf
struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1, 0.1|0.2, "P");

struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);

proc fct(struct PV p0, struct DS d0);
  local p,y;
  p = pvUnpack(p0, "P");
  y = p[1] * exp(-p[2] * d0.dataMatrix);
  retp(y);
endp;

g = gradMT(&fct,p1,d0);
```

Source

gradmtt.src

gradMTTm

Purpose

Computes numerical gradient with mask using threads.

Include

sqpsolvemt.sdf

Format

```
g = gradMTTm(&fct, par1, data1, mask);
```

Input

<i>&fct</i>	scalar, pointer to procedure returning either Nx1 vector or 1x1 scalar
<i>par1</i>	structure of type PV containing parameter vector at which gradient is to be evaluated
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i>
<i>mask</i>	Kx1 matrix, elements in <i>g</i> corresponding to elements of mask set to zero are not computed otherwise are computed.

Output

<i>g</i>	NxK Jacobian or 1xK gradient
----------	------------------------------

Remarks

par1 must be created using the **pvPack** procedures

gradp, gradcplx

Example

```
#include sqpsolvemt.sdf
struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1, 0.1|0.2, "P");

struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1, 1, 15);

proc fct(struct PV p0, struct DS d0);
    local p, y;
    p = pvUnpack(p0, "P");
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    retp(y);
endp;

mask = { 0, 1 };
g = gradMTTm(&fct, p1, d0, mask);
```

Source

gradmtt.src

gradp, gradcplx

Purpose

Computes the gradient vector or matrix (Jacobian) of a vector-valued function that has been defined in a procedure. Single-sided (forward difference) gradients are computed. **gradcplx** allows for complex arguments.

Format

```
 $g = \text{gradp}(\&f, x0);$   
 $g = \text{gradcplx}(\&f, x0);$ 
```

Input

$\&f$ a pointer to a vector-valued function ($f: K \times 1 \rightarrow N \times 1$) defined as a procedure. It is acceptable for $f(x)$ to have been defined in terms of global arguments in addition to x , and thus f can return an $N \times 1$ vector:

```
proc f(x);  
  ret( exp(x.*b) );  
endp;
```

$x0$ $K \times 1$ vector of points at which to compute gradient.

Output

g $N \times K$ matrix containing the gradients of f with respect to the variable x at $x0$.

Remarks

gradp will return a row for every row that is returned by f . For instance, if f returns a scalar result, then **gradp** will return a $1 \times K$ row vector. This allows the same function to be used regardless of N , where N is the number of rows in the result returned by f . Thus, for instance, **gradp** can be used to compute the Jacobian matrix of a set of equations.

graphprt

Example

```
proc myfunc(x);  
    retp(x .* 2 .* exp(x .* x ./ 3));  
endp;  
  
x0 = 2.5|3.0|3.5;  
y = gradp(&myfunc,x0);
```

After the code above, *y* is equal to:

82.989017	0.00000000	0.00000000
0.00000000	281.19753	0.00000000
0.00000000	0.00000000	1087.9541

It is a 3x3 matrix because we are passing it 3 arguments and **myfunc** returns 3 results when we do that; the off-diagonals are zeros because the cross-derivatives of 3 arguments are 0.

Source

gradp.src

See Also

[hessp](#), [hesseplx](#)

graphprt

Purpose

Controls automatic printer hardcopy and conversion file output. Note: This function is for use with the deprecated PQG graphics. Use the **plotSave** function instead.

Library

pgraph

Format

```
graphprt(str);
```

Input

<i>str</i>	string, control string.
------------	-------------------------

Portability

UNIX

Not supported.

Remarks

graphprt is used to create hardcopy output automatically without user intervention. The input string *str* can have any of the following items, separated by spaces. If *str* is a null string, the interactive mode is entered. This is the default.

<i>-p</i>	print graph.
<i>-po=c</i>	set print orientation:
<i>l</i>	landscape.
<i>p</i>	portrait.
<i>-c=n</i>	convert to another file format:
<i>1</i>	Encapsulated PostScript file.
<i>3</i>	HPGL Plotter file.
<i>5</i>	BMP (Windows Bitmap).

graphprt

	8	WMF (Windows Enhanced Metafile).
-cf=name		set converted output file name.
-i		minimize (iconize) the graphics window.
-q		close window after processing.
-w=n		display graph, wait <i>n</i> seconds, then continue.

If you are not using graphic panels, you can call **graphprt** anytime before the call to the graphics routine. If you are using graphic panels, call **graphprt** just before the **endwind** statement.

The print option default values are obtained from the viewer application. Any parameters passed through **graphprt** will override the default values. See **PUBLICATION QUALITY GRAPHICS, CHAPTER 1**.

Example

Automatic print using a single graphics call:

```
library pgraph;
graphset;

load x,y;

graphprt("-p"); /* tell "xy" to print */
xy(x,y);        /* create graph and print */
```

Automatic print using multiple graphic panels. Note **graphprt** is called once just before the **endwind** call:

```
library pgraph;
graphset;

load x,y;

begwind;
```



```
window(1,2,0); /* create two windows */
setwind(1);
xy(x,y); /* first graphics call */
nextwind;
xy(x,y); /* second graphics call */
graphprt("-p");
endwind; /* print page containing all graphs */
```

The next example shows how to build a string to be used with **graphprt**:

```
library pgraph;
graphset;
load x,y;

cvtnam = "mycvt.eps"; /* name of output file */
/* concatenate options into one string */
cmdstr = "-c=1" $+ " -cf=" $+ cvtnam;
cmdstr = cmdstr $+ " -q";

graphprt(cmdstr); /* tell "xy" to convert and */
/* close */
xy(x,y); /* create graph and convert */
```

The above string *cmdstr* will read as follows:

```
"-c=1 -cf=mycvt.eps -q"
```

Source

pgraph.src

graphset

graphset

Purpose

Reset graphics global variables to default values. Note: This function is for use with the deprecated PQG graphics.

Library

pgraph

Format

```
graphset;
```

Remarks

This procedure is used to reset the defaults between graphs.

graphset may be called between each graphic panel to be displayed.

To change the default values of the global control variables, make the appropriate changes in the file `pgraph.dec` and to the procedure **graphset**.

Source

`pgraph.src`

h

h5create

Purpose

Creates an HDF5 data file, or adds a new dataset to an existing HDF5 file.

Format

```
ret = h5create(fname, dname, dims);  
ret = h5create(fname, dname, dims, datatype);  
ret = h5create(fname, dname, dims, datatype, chunk_size);
```

Input

<i>fname</i>	String, name of the HDF5 file. The file extension, .h5 recommended, but not required.
<i>dname</i>	String, a name of the data set in HDF5 file. e.g. <code>"/mydata"</code> .
<i>dims</i>	Nx1 matrix, where N is the number of dimensions of the dataset, the size of each of the dimensions of the dataset.
<i>datatype</i>	String, data type. Valid options include: <ul style="list-style-type: none">• <code>"double"</code>• <code>"float"</code>

h5create

	<ul style="list-style-type: none">• "int64"• "int32"• "uint64"• "uint32"
	Default is "double".
<code>chunk_size</code>	Matrix or array, with the same dimensions as the data set, specifying the size of the chunks of data that will be created in the file.

Output

<code>ret</code>	Scalar, 0 if successful.
------------------	--------------------------

Remarks

- HDF5 files can hold more than one dataset. They are referenced in the same manner as a Linux or Mac file system. The base or root node is '/'. All datasets are relative to this root node.
- If a dataset name contains multiple intermediate groups, for example:

```
"/surveys/household/Washington"
```

- and the intermediate groups, **surveys** and **household** in the above string, do not yet exist, **h5create** will create them.
- By default, HDF5 datasets may not change size. To make one of the dimensions expandable, set it to `__INFP`.
 - All columns of an HDF5 dataset must be of the same data type. However, multiple datasets with different data types may be created in a single HDF5 file.
 - Information about a dataset, called an attribute, may be attached to a dataset in an HDF5 file with the function **h5writeAttribute**.

- Chunk size must be specified when users create a dataset with more than 2 dimensions and one of those dimensions is unlimited (`__INFP`).

Examples

Example 1: Create a fixed size 2-dimensional dataset

```
// Define a name of a HDF5 file
fname = "testdata.h5";

// Define a data set name under the HDF5 file
dname = "/mydata";

// Define the size of the data set, 100 rows and 5 columns

r = 100;
c = 5;

// Create a data set under the HDF5 file
call h5create(fname, dname, r|c);

// Fill dataset with random normal data
x = rndn(100, 5);
h5write(fname, dname, x);
```

Example 2: Create a 2-dimensional dataset with 5 columns and a flexible number of rows

```
// Define a size of flexible rows and 5 columns
dims = __INFP|5;

// Create a data set
call h5create("expandable_data.h5", "/data", dims);
```

Example 3: Create a 3-Dimensional dataset and one intermediate group

h5open

```
// Define a new data set name, including one intermediate
group
dname = "/household/Washington";

// Define a 3-dimensional dataset, containing 3 matrices
with
// 8 columns and an expandable number of rows
dims = 3|__INFP|8;

// Store the data in chunks of 1000x8 elements
chunk_size = { 1, 1000, 8 };

// Store data as 4 byte floating point (about 8 digits of
precision)
dtype = "float";

// Create the data set
call h5create("surveys.h5", dname, dims, dtype, chunk_
size);

// Create another data set of the same type inside the same
file
call h5create("surveys.h5", "/household/Oregon", dims,
dtype, chunk_size);
```

See Also

[h5read](#), [h5write](#), [open](#), [create](#), [writer](#), [seekr](#), [eof](#)

h5open

Purpose

Open an HDF5 dataset and returns a file handle.

Format

```
fh = h5open(fname, dname, mode);
```

Input

<i>fname</i>	string, a name of HDF5 file to open.
<i>dname</i>	string, a name of a data set (or group) in HDF5 file. e.g. "/mydata" .
<i>mode</i>	string, the mode with which to open the file. Valid options include: <ul style="list-style-type: none">• "open": open file for read, positioned at the first row.• "update": open file for write, positioned at the first row.• "append": open file for write, positioned at the end of the file.

Output

<i>fh</i>	scalar, file handle for use with readr , or writer .
-----------	--

Remarks

- The file handle must be closed with either the **close** function or **closeall** when you are finished using it.
- To read or write data to a file opened with **h5open**, use **readr**, and **writer**.
- To read and write data to an HDF5 dataset without opening a file handle, use **h5read** and **h5write**.
- The function **dataopen** can open file handles for reading and writing to HDF5 datasets and other file types.

Examples

Example 1: Create and write to an HDF5 dataset

```
rndseed 2344;

// Create a 4 row by 3 column HDF5 data set
call h5create("testdata.h5", "/mydata", 4 | 3);

// Create a 2x3 matrix
x = { 1.1 2.2 3.3,
      4.4 5.5 6.6 };

// Open a file handle
fh = h5open("testdata.h5", "/mydata", "update");

// Write the data in 'x' to the the first two rows
call writer(fh, x);

// Create a 2x3 matrix
y = { 10 20 30,
      40 50 60 };

// Write the data in 'y' to the the final two rows
call writer(fh, y);

// Close the file handle
close(fh);
```

Example 2: Read data written in Example 1

```
// Open a file handle
fh = h5open("testdata.h5", "/mydata", "read");

// Read the first row
```



```

a  = readr(fh,1);

// Read the second, third and fourth rows
b  = readr(fh,3);

// Close the file
call close(fh);

```

After the code above

```

a =  1.10  2.20  3.30

b =   4.4   5.5   6.6
    10.0  20.0  30.0
    40.0  50.0  60.0

```

See Also

[h5create](#), [h5read](#), [h5write](#), [open](#), [dataopen](#), [readr](#), [seekr](#)

h5read

Purpose

Reads data from an HDF5 dataset into a GAUSS matrix.

Format

```

y = h5read(fname, dname);
y = h5read(fname, dname, dims);
y = h5read(fname, dname, dims, offset);

```

h5read

Input

<i>fname</i>	String, a name of the HDF5 file.
<i>dname</i>	String, a name of the data set in HDF5 file. e.g. <code>"/mydata"</code> .
<i>dims</i>	Nx1 vector, where N is the number of dimensions in the dataset, the dimensions of data to read.
<i>offset</i>	Nx1 vector, where N is the number of dimensions in the dataset, the data to skip.

Output

<i>y</i>	Matrix, or multi-dimensional array, the data requested from the file.
----------	---

Examples

Example 1: Basic write then read entire contents of an HDF5 file

```
// Define a name of a HDF5 file
fname = "testdata.h5";

// Define a name of a dataset
dname = "/mydata";

// Create a 3x2 fixed dimension data set in a HDF5 file
call h5create(fname, dname, 3|2);

// Create a 3x2 matrix
x = { 9 4,
      2 1,
      0 7 };

// Write x to dataset
```

```
call h5write(fname, dname,x);

// Read all contents from dataset
y = h5read(fname, dname);
```

After the code above:

```
x = 9  4      y = 9  4
    2  1      2  1
    0  7      0  7
```

Example 2: Read HDF5 first two rows and first two columns

```
// Size of data to read
dims = 2|2;

// Read data from file created in Example 1, above
y2 = h5read("testdata.h5", "/mydata", dims);
```

After the code above:

```
y2 =      9  4
      2  1
```

Example 3: Read HDF5 first two rows and first two columns

```
// Size of data to read
dims = 2|2;

// Dimensions to skip: 1 row and 0 columns
offset = 1|0;
```

h

h5read

```
// Read data from file created in Example 1, above
y3 = h5read("testdata.h5", "/mydata", dims, offset);
```

After the code above:

```
y3 =      2 1
        0 7
```

Example 4: Read HDF5 file with offset at more than one dimension

```
// Define a new data set name
dname = "/highdimension";

// Create 3 dimensions, with __INFP indicating expandable
rows
dims = 3 | __INFP | 4;

// Define data type
datatype = "double";

// Define chunk size since the second dimension is infinite

chunk_size = 1|128|4;

// Create "highdimension" data set, inside file created in
Example 1 (above)
call h5create("testdata.h5", dname, dims, datatype, chunk_
size);

// Set seed for repeatable random numbers
rndseed 7672342;

// Create random normal data array, with dimensions 3x10x4

x = areshape(rndn(3 * 10 * 4, 1), 3 | 10 | 4);
```

```

// Write it into data set
call h5write("testdata.h5", dname, x);

// Skip first two rows and first column of each
// of the 3 matrices in the 3x10x4 array
offset = 0|2|1;

// Define the read size after removing offset
dims_read = 2|2|2;

// Run h5read function
y4 = h5read("testdata.h5", dname, dims_read, offset);

```

After the code above, we see that *y4* is a 2x2x2 (number of dimensions to read *dims_read*) array, containing the contents of the *x*, after skipping the dimensions { 0, 2, 1 } specified in *offset*:

```

y4 = Plane [1,...]

      -0.61557786      -0.76592868
      -0.30976522      0.11296623

      Plane [2,...]

      -0.23144975      -1.1369840
      -0.89682110      1.6684102

x =   Plane [1,...]

      -1.2045242      -1.0675179      -0.74403139      -
0.72860218
      0.20337032      -0.48451306      -0.0039387096
0.46361645

```

h5read

-0.57448560	-0.61557786	-0.76592868	-
0.032267807			
-0.88033211	-0.30976522	0.11296623	
1.2724183			
-1.4409872	-0.90939666	0.22487451	-
0.37188053			
-1.5478724	-0.43944280	0.010049938	
1.0196427			
1.3352024	1.0734150	-0.98373668	-
0.57590137			
-0.32428680	0.53099143	-0.71162764	-
2.1188409			
-0.22060808	-0.024172215	0.64942867	-
0.51276843			
0.12600180	-0.65155519	-2.2815720	
1.4961735			
Plane [2,...]			
-0.78337697	-0.52759501	-1.2322159	-
0.31936828			
-0.47552440	-1.0708763	0.43111378	
1.5146598			
0.54119533	-0.23144975	-1.1369840	-
0.11052318			
0.47963176	-0.89682110	1.6684102	-
0.43704128			
-0.27511827	-0.65207535	-0.17394561	-
0.84737201			
-0.14595989	-0.028056845	0.50018732	-
0.76191566			
-0.98846912	1.4389099	1.3716329	-
1.3419693			
-0.29630831	-1.2029618	-1.4958204	-
2.0829113			

```
-0.56764971    -0.53397186    -0.95002213    -
0.10182348
    -1.6156998    -1.5120152    0.013456774    -
0.037790884

    Plane [3,...]

    0.42346079    -0.61879151    0.062894922
0.43245351
    0.092322769    0.68876937    -0.61677358
2.8805431
    0.33204968    2.1878476    -1.1113500    -
0.38579652
    0.35943828    0.32172778    -0.25074937    -
0.34662609
    -0.95053031    0.0010335034    -0.12838005    -
1.2333248
    0.59991891    0.73834232    -0.23521782    -
0.63566653
    -1.5290045    -0.36202638    0.69077565
1.0898312
    0.71036599    0.086441099    -0.40250335
0.58997554
    -0.55612014    0.084524826    0.60194547
0.26031576
    0.49760949    0.070206385    0.65894867
0.73385573
```

See Also

[h5create](#), [h5write](#), [open](#), [create](#), [writer](#), [seekr](#), [eof](#)

h5readAttribute

h5readAttribute

Purpose

Read attributes from an HDF5 file into GAUSS.

Format

```
attr_read = h5readAttribute(fname, dname, attr_name);
```

Input

<i>fname</i>	String, a name of HDF5 file.
<i>dname</i>	String, a name of the HDF5 data set.
<i>attr_name</i>	String, the name of attribute.

Output

<i>attr_read</i>	Matrix, multi-dimensional array or string array, entire contents of the HDF5 dataset attribute.
------------------	---

Remarks

- HDF5 does not support partial read or write of dataset attributes. The entire contents of the attribute will be read.
- GAUSS functions that accept HDF5 datasets as a datasource, expect the dataset to have an attribute named "headers", containing the variable names of the dataset.

Example

Example 1: Create an HDF5 dataset and add headers

```
//Create an HDF5 dataset with room for 100 observations of
4 variables
call h5create("commodities.h5", "/energy", 100 | 4);

//Variable names for the dataset
attr = "Crude Oil"|$|"Gasoline"|$|"Heating Oil"|$|"Diesel";

//Define a name of the attributes
attr_name = "headers";

//Write attributes to a HDF5 file
call h5writeAttribute("commodities.h5", "/energy", attr_
name, attr);

//Read attributes from a HDF5 file
attr_read = h5readAttribute("commodities.h5", "/energy",
attr_name);
```

Example 1a: Add data and calculate descriptive statistics

```
//Set seed for repeatable random data
rndseed 54235;

//Create 100x4 random normal data
x = rndn(100, 4);

//Write data to dataset created in the example above
call h5write("commodities.h5", "/energy", x);

//Calculate descriptive statistics on some of the variables
//using an hdf5 file schema (h5://filename/dataset)
call dstatmt("h5://commodities.h5/energy", "Gasoline + Heat-
ing Oil");
```

h

h5write

Variable		Mean	Std Dev	Variance	Minimum
Maximum	Valid	Missing			

Gasoline		0.0212	1.0130	1.0261	-2.9943
2.3527	100	0			
Heating Oil		-0.1120	0.9263	0.8580	-2.7726
3.0910	100	0			

See Also

[h5create](#), [h5writeAttribute](#), [h5read](#), [h5write](#)

h5write

Purpose

Write a GAUSS matrix or N-dimensional array to an HDF5 dataset.

Format

```
ret = h5write(fname, dname, x);
```

Input

<i>fname</i>	String, name of the HDF5 file.
<i>dname</i>	String, name of the dataset in the HDF5 file.
<i>x</i>	Matrix or N-dimensional array, the data to write to the file.

Output

```
ret          Scalar, 0 if successful.
```

Examples

Example 1: Basic HDF5 file/dataset creation and write

```
//Create a 3x2 dataset
call h5create("testdata.h5", "/writetest", 3|2);

//Create a data matrix
x = { 1.1 2.2,
      3.3 4.4,
      5.5 6.6 };

//Write x to HDF5 dataset
call h5write("testdata.h5", "/writetest", x);

//Read data from a data set of a HDF5 file
y = h5read("testdata.h5", "/writetest");
```

After the code above:

```
y =  1.1  2.2
     3.3  4.4
     5.5  6.6
```

Example 2: Write over first two rows of dataset

```
x_new = { 1000 2000,
          3000 4000 };

//Write x_new over first 2 rows of HDF5 dataset
call h5write("testdata.h5", "/writetest", x_new);
```

h5writeAttribute

```
//Read data again
y_new = h5read("testdata.h5", "/writetest");
```

After the above code:

```
y_new = 1000    2000
        3000    4000
        5.5     6.6
```

Remarks

To write rows of data to an HDF5 dataset in a iterative manner, see **writer**.

See Also

[h5create](#), [h5read](#), [h5writeAttribute](#), [dataopen](#), [writer](#), [seekr](#)

h5writeAttribute

Purpose

Writes a GAUSS matrix, N-dimensional array or string array as an attribute of an HDF5 dataset.

Format

```
ret = h5writeAttribute(fname, dname, attr_name, attr);
```

Input

<i>fname</i>	String, a name of the HDF5 file.
<i>dname</i>	String, a name of the dataset in the HDF5 file.

<i>attr_name</i>	String, the name of attribute to write.
<i>attr</i>	Matrix, N-dimensional array or string array, the contents of the attribute.

Output

<i>ret</i>	Scalar, 0 if successful.
------------	--------------------------

Remarks

- Attributes in an HDF5 file cannot be read or written partially. The entire contents of the attribute must be read or written in one call.
- **GAUSS** functions that take in an HDF5 dataset as a datasource (see **dstatmt**, **glm**), expect the dataset to have an attribute called "headers", containing the variable names of the dataset.

Examples

Example 1: Create an HDF5 dataset and add headers

```
//Create an HDF5 dataset with room for 100 observations of
4 variables
call h5create("commodities.h5", "/energy", 100 | 4);

//Variable names for the dataset
attr = "Crude Oil"|"Gasoline"|"Heating Oil"|"Diesel";

//Define a name of the attributes
attr_name = "headers";

//Write attributes to a HDF5 file
call h5writeAttribute("commodities.h5", "/energy", attr_
```

h5writeAttribute

```
name, attr);

//Read attributes from a HDF5 file
attr_read = h5readAttribute("commodities.h5", "/energy",
attr_name);
```

Example 1a: Add data and calculate descriptive statistics

```
//Set seed for repeatable random data
rndseed 54235;

//Create 100x4 random normal data
x = rndn(100, 4);

//Write data to dataset created in the example above
call h5write("commodities.h5", "/energy", x);

//Calculate descriptive statistics on some of the variables
//using an hdf5 file schema (h5://filename/dataset)
call dstatmt("h5://commodities.h5/energy", "Gasoline + Heat-
ing Oil");
```

```
-----
-----
Variable          Mean   Std Dev   Variance   Minimum
Maximum   Valid   Missing
-----
-----

Gasoline          0.0212   1.0130     1.0261   -2.9943
2.3527          100         0
Heating Oil      -0.1120   0.9263     0.8580   -2.7726
3.0910          100         0
```

See Also

[h5readAttribute](#), [h5read](#), [h5write](#), [h5create](#), [h5write](#)

hasimag

Purpose

Tests whether the imaginary part of a complex matrix is negligible.

Format

```
y = hasimag(x);
```

Input

<i>x</i>	NxK matrix.
----------	-------------

Output

<i>y</i>	scalar, 1 if the imaginary part of <i>x</i> has any nonzero elements, 0 if it consists entirely of 0's.
----------	---

Remarks

The function **iscplx** tests whether *x* is a complex matrix or not, but it does not test the contents of the imaginary part of *x*. **hasimag** tests the contents of the imaginary part of *x* to see if it is zero.

hasimag actually tests the imaginary part of *x* against a tolerance to determine if it is negligible. The tolerance used is the imaginary tolerance set with the **sysstate** command, case 21.

Some functions are not defined for complex matrices. **iscplx** can be used to determine whether a matrix has no imaginary part and so can pass through those functions. **hasimag** can be used to determine whether a complex matrix has a negligible imaginary part and could thus be converted to a real matrix to pass through those functions.

header

iscplx is useful as a preliminary check because for large matrices it is much faster than **hasimag**.

Example

```
x = { 1    2 3i,  
      4-i 5 6i,  
      7    8i 9 };  
  
if hasimag(x);  
    //code path for complex case  
else;  
    //code path for real case  
endif;
```

See Also

[iscplx](#)

header

Purpose

Prints a header for a report.

Format

```
header(prcnm, dataset, ver);
```

Input

<i>prcnm</i>	string, name of procedure that calls header .
<i>dataset</i>	string, name of data set.

<code>ver</code>	2x1 numeric vector, the first element is the major version number of the program, the second element is the revision number. Normally this argument will be the version/revision global (<code>__??_ver</code>) associated with the module within which header is called. This argument will be ignored if set to 0.
------------------	--

Global Input

<code>__header</code>	string, containing one or more of the following letters: <div><div><code>t</code></div><div>title is to be printed</div></div> <div><div><code>l</code></div><div>lines are to bracket the title</div></div> <div><div><code>d</code></div><div>a date and time is to be printed</div></div> <div><div><code>v</code></div><div>version number of program is to be printed</div></div> <div><div><code>f</code></div><div>file name being analyzed is to be printed</div></div>
<code>__title</code>	string, title for header.

Source

gauss.src

headermt

Purpose

Prints a header for a report.

headermt

Format

```
headermt(prcnm, dataset, ver, header, title);
```

Input

<i>prcnm</i>	string, name of procedure that calls header .										
<i>dataset</i>	string, name of data set.										
<i>ver</i>	2x1 numeric vector, the first element is the major version number of the program, the second element is the revision number. Normally this argument will be the version/revision global (<code>__??_ver</code>) associated with the module within which header is called. This argument will be ignored if set to 0.										
<i>header</i>	string, containing one or more of the following letters: <table><tr><td><i>t</i></td><td>title is to be printed</td></tr><tr><td><i>l</i></td><td>lines are to bracket the title</td></tr><tr><td><i>d</i></td><td>a date and time is to be printed</td></tr><tr><td><i>v</i></td><td>version number of program is to be printed</td></tr><tr><td><i>f</i></td><td>file name being analyzed is to be printed</td></tr></table>	<i>t</i>	title is to be printed	<i>l</i>	lines are to bracket the title	<i>d</i>	a date and time is to be printed	<i>v</i>	version number of program is to be printed	<i>f</i>	file name being analyzed is to be printed
<i>t</i>	title is to be printed										
<i>l</i>	lines are to bracket the title										
<i>d</i>	a date and time is to be printed										
<i>v</i>	version number of program is to be printed										
<i>f</i>	file name being analyzed is to be printed										
<i>title</i>	string, title for header.										

Source

gaussmt.src

hess

Purpose

Computes the Hessenberg form of a square matrix.

Format

```
{ H, Z } = hess(A);
```

Input

A	$K \times K$ real or complex matrix.
-----	--------------------------------------

Output

H	$K \times K$ matrix, Hessenberg form.
Z	$K \times K$ matrix, transformation matrix.

Example

```
A = { 0.5 0.2 0.33,  
      1.4 0.5 0.6,  
      0.7 1.2 0.9 };
```

```
{ H, Z } = hess(A);
```

After the code above:

H =	0.500	-0.326	0.206	Z =	1.000	0.000
	0.000					
	-1.565	1.300	-0.400		0.000	-0.894
	0.447					-

hessMT

	0.000	-1.000	0.100	0.000	-0.447
0.894					

Remarks

hess computes the Hessenberg form of a square matrix. The Hessenberg form is an intermediate step in computing eigenvalues. It also is useful for solving certain matrix equations that occur in control theory (see Van Loan, Charles F. "Using the Hessenberg Decomposition in Control Theory". *Algorithms and Theory in Filtering and Control*. Sorenson, D.C. and R.J. Wets, eds., Mathematical Programming Study No. 18, North Holland, Amsterdam, 1982, 102-111).

Z is an orthogonal matrix that transforms A into H and vice versa. Thus:

$$H = Z' * A * Z$$

and since Z is orthogonal,

$$A = Z * H * Z'$$

A is reduced to upper Hessenberg form using orthogonal similiarity transformations. This preserves the Frobenious norm of the matrix and the condition numbers of the eigenvalues.

See Also

[eig](#), [qz](#), [schur](#)

hessMT

Purpose

Computes numerical Hessian.

Include

optim.sdf

Format

```
h = hessMT(&fct, par1, data1);
```

Input

<i>&fct</i>	scalar, pointer to procedure returning either Nx1 vector or 1x1 scalar.
<i>par1</i>	an instance of structure of type PV containing parameter vector at which Hessian is to be evaluated.
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i> .

Output

<i>h</i>	KxK matrix, Hessian.
----------	----------------------

Remarks

par1 must be created using the **pvPack** procedures.

Example

```
#include optim.sdf
struct PV p1;
struct DS d0;

p1 = pvCreate;
p1 = pvPack(p1, 0.1|0.2, "P");
```

hessMTg

```
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);

proc fct(struct PV p0, struct DS d0);
    local p,y;

    p = pvUnpack(p0, "P");
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    retp(y);
endp;

h = hessMT(&fct,p1,d0);
```

Source

hessmt.src

hessMTg

Purpose

Computes numerical Hessian using gradient procedure.

Include

optim.sdf

Format

h = **hessMTg**(*&gfct*, *par1*, *data1*);

Input

<i>&gfct</i>	scalar, pointer to procedure computing either 1xK gradient or NxK Jacobian.
------------------	---

<i>par1</i>	an instance of structure of type PV containing parameter vector at which Hessian is to be evaluated.
<i>data1</i>	structure of type DS containing any data needed by <i>gfct</i> .

Output

<i>h</i>	KxK matrix, Hessian.
----------	----------------------

Remarks

par1 must be created using the **pvPack** procedures.

Example

```
#include optim.sdf
struct PV p1;
struct DS d0;
p1 = pvCreate;
p1 = pvPack(p1,0.1|0.2, "P");
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);

proc gfct(&fct, struct PV p0, struct DS d0);
    local p,y,g1,g2;

    p = pvUnpack(p0, "P");
    g1 = exp(-p[2] * d0.dataMatrix);
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    g2 = -p[1] * d0.dataMatrix .* g1;
    retp(g1~g2);
endp;
```

h

hessMTgw

```
h = hessMTg(&gfct,p1,d0);
```

Source

hessmt.src

hessMTgw

Purpose

Computes numerical Hessian using gradient procedure with weights.

Include

optim.sdf

Format

```
h = hessMTgw(&gfct,par1,data1,wgts);
```

Input

<i>&gfct</i>	scalar, pointer to procedure computing either NxK Jacobian.
<i>par1</i>	an instance of structure of type PV containing parameter vector at which Hessian is to be evaluated.
<i>data1</i>	structure of type DS containing any data needed by <i>gfct</i> .
<i>wgts</i>	Nx1 vector.

Output

<i>h</i>	KxK matrix, Hessian.
----------	----------------------

Remarks

par1 must be created using the **pvPack** procedures.

Example

```
#includeoptim.sdf
struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1,0.1|0.2, "P");
struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);
wgts = zeros(5,1) | ones(10,1);

proc gfct(&fct, struct PV p0, struct DS d0);
    local p,y,g1,g2;

    p = pvUnpack(p0, "P");
    g1 = exp(-p[2] * d0.dataMatrix);
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    g2 = -p[1] * d0.dataMatrix .* g1;
    retp(g1~g2);
endp;

h = hessMTgw(&gfct,p1,d0,wgts);
```

Source

hessmt.src

hessMTm

Purpose

Computes numerical Hessian with mask.

hessMTm

Include

optim.sdf

Format

```
h = hessMTm(&fct, par1, data1, mask);
```

Input

<i>fct</i>	scalar, pointer to procedure returning either Nx1 vector or scalar.
<i>par1</i>	an instance of structure of type PV containing parameter vector at which Hessian is to be evaluated.
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i> .
<i>mask</i>	KxK matrix, elements in <i>h</i> corresponding to elements of mask set to zero are not computed, otherwise are computed.

Output

<i>h</i>	KxK matrix, Hessian.
----------	----------------------

Remarks

par1 must be created using the **pvPack** procedures. Only lower left part of mask looked at.

Example

```
#include optim.sdf
struct PV pl;
```

```

p1 = pvCreate;
p1 = pvPack(p1, 0.1|0.2, "P");
struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);

mask = { 1 1,
         1 0 };

proc fct(struct PV p0, struct DS d0);
    local p,y;

    p = pvUnpack(p0, "P");
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    retp(y);
endp;

h = hessMTm(&fct,p1,d0,mask);

```

Source

hessmt.src

hessMTmw

Purpose

Computes numerical Hessian with mask and weights.

Include

optim.sdf

Format

```
h = hessMTmw(&fct, par1, data1, mask, wgts);
```

Input

<i>&fct</i>	scalar, pointer to procedure returning Nx1 vector.
<i>par1</i>	an instance of structure of type PV containing parameter vector at which Hessian is to be evaluated.
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i> .
<i>mask</i>	KxK matrix, elements in <i>h</i> corresponding to elements of mask set to zero are not computed, otherwise are computed.
<i>wgts</i>	Nx1 vector, weights.

Output

<i>h</i>	KxK matrix, Hessian.
----------	----------------------

Remarks

fct must evaluate to an Nx1 vector conformable to the weight vector. *par1* must be created using the **pvPack** procedures.

Example

```
#include optim.sdf

struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1,0.1|0.2, "P");
struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);
wgts = zeros(5,1) | ones(10,1);
```

```

mask = { 1 1,
         1 0 };

proc fct(&fct, struct PV p0, struct DS d0, wgts);
    local p,y;

    p = pvUnpack(p0, "P");
    y = p[1] * exp( -p[2] * d0.dataMatrix);
    retp(y);
endp;

h = hessMTmw(&fct,p1,d0,mask,wt);

```

Source

hessmt.src

hessMTT

Purpose

Computes numerical Hessian using available threads.

Format

```
h = hessMTT(&fct,par1,data1);
```

Include

optim.sdf

Input

<i>fct</i>	scalar, pointer to procedure returning either Nx1 vector
------------	--

hessMTT

<i>par1</i>	or 1x1 scalar. structure of type PV containing parameter vector at which Hessian is to be evaluated
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i>

Output

<i>h</i>	KxK matrix, Hessian
----------	---------------------

Remarks

par1 must be created using the **pvpack** procedures

Example

```
#include optim.sdf
struct PV p1;
p1 = pvCreate;

p1 = pvPack(p1, 0.1|0.2, "P");
struct DS d0;
d0 = dsCreate;

d0.dataMatrix = seqa(1, 1, 15);

proc fct(struct PV p0, struct DS d0);
    local p, y;
    p = pvUnpack(p0, "P");
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    retp(y);
endp;
```

```
h = hessMTT (&fct, p1, d0);
```

Source

hessmtt.src

hessMTTg

Purpose

Computes numerical Hessian using gradient procedure with available threads.

Include

optim.sdf

Format

```
h = hessMTTg(&gfct, par1, data1);
```

Input

<i>&gfct</i>	scalar, pointer to procedure computing either 1xK gradient or NxK Jacobian
<i>par1</i>	structure of type PV containing parameter vector at which Hessian is to be evaluated
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i>

Output

<i>h</i>	KxK matrix, Hessian
----------	---------------------



hessMTTg

Remarks

par1 must be created using the **pvPack** procedures.

Example

```
#include optim.sdf

struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1,0.1|0.2, "P");

struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);

proc gfct(&fct, struct PV p0, struct DS d0, wgt);
    local p,y,g1,g2;
    p = pvUnpack(p0, "P");
    g1 = exp(-p[2] * d0.dataMatrix);
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    g2 = -p[1] * d0.dataMatrix .* g1;
    retp(g1~g2);
endp;

h = hessMTTg(&gfct,p1,d0);
```

Source

hessmtt.src

hessMTTgw

Purpose

Computes numerical Hessian using gradient procedure with weights and using available threads.

Include

optim.sdf

Format

$h = \text{hessMTTgw}(\&gfct, \text{par1}, \text{data1}, \text{wgts});$

Input

<i>gfct</i>	scalar, pointer to procedure computing either 1xK gradient or NxK Jacobian
<i>par1</i>	structure of type PV containing parameter vector at which Hessian is to be evaluated
<i>data1</i>	structure of type DS containing any data needed by fct
<i>wgts</i>	Nx1 vector, weights

Output

<i>h</i>	KxK matrix, Hessian
----------	---------------------

Remarks

par1 must be created using the **pvPack** procedures.

hessMTTm

Example

```
#include optim.sdf

struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1, 0.1|0.2, "P");

struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);
wgts = zeros(5,1) | ones(10,1);

proc gfct(&fct, struct PV p0, struct DS d0);
    local p,y,g1,g2;
    p = pvUnpack(p0, "P");
    g1 = exp(-p[2] * d0.dataMatrix);
    y = p[1] * exp(-p[2] * d0.dataMatrix);
    g2 = -p[1] * d0.dataMatrix .* g1;
    retp(g1~g2);
endp;

h = hessMTTg(&gfct,p1,d0,wgts);
```

Source

hessmtt.src

hessMTTm

Purpose

Computes numerical Hessian with mask using available threads.

Include

optim.sdf

Format

```
h = hessMTTm(&fct, par1, data1, mask);
```

Input

<i>fct</i>	scalar, pointer to procedure returning either Nx1 vector or 1x1 scalar.
<i>par1</i>	structure of type PV containing parameter vector at which Hessian is to be evaluated
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i>
<i>mask</i>	KxK matrix, elements in <i>h</i> corresponding to elements of mask set to zero are not computed otherwise are computed

Output

<i>h</i>	KxK matrix, Hessian
----------	---------------------

Remarks

par1 must be created using the **pvpPack** procedures. Only lower left part of mask looked at.

Example

```
#include optim.sdf
struct PV pl;
```

hessMTw

```
p1 = pvCreate;
p1 = pvPack(p1,0.1|0.2, "P");
struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);

mask = { 1 1
         1 0 };

proc fct(struct PV p0, struct DS d0);
    local p,y;
    p = pvUnpack(p0, "P");
    y = p[1] * exp( -p[2] * d0.dataMatrix);
    retp(y);
endp;

h = hessMTTm(&fct,p1,d0,mask);
```

Source

hessmtt.src

hessMTw

Purpose

Computes numerical Hessian with weights.

Include

optim.sdf

Format

```
h = hessMTw(&fct, parl, data1, wgts);
```

Input

<i>fct</i>	scalar, pointer to procedure returning Nx1 vector.
<i>par1</i>	an instance of structure of type PV containing parameter vector at which Hessian is to be evaluated.
<i>data1</i>	structure of type DS containing any data needed by <i>fct</i> .
<i>wgts</i>	Nx1 vector, weights.

Output

<i>h</i>	KxK matrix, Hessian.
----------	----------------------

Remarks

fct must evaluate to an Nx1 vector conformable to the weight vector. *par1* must be created using the **pvPack** procedures.

Example

```
#include optim.sdf
struct PV p1;
p1 = pvCreate;
p1 = pvPack(p1, 0.1|0.2, "P");

struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1,1,15);
wgt = zeros(5,1) | ones(10,1);

proc fct(&fct, struct PV p0, struct DS d0, wgt);
    local p,y;
```

h

hessp, hesscplx

```
p = pvUnpack(p0, "P");
y = p[1] * exp(-p[2] * d0.dataMatrix);
retp(y);
endp;

h = hessMTw(&fct,p1,d0,wgt);
```

Source

hessmt.src

hessp, hesscplx

Purpose

Computes the matrix of second partial derivatives (Hessian matrix) of a function defined as a procedure. **hesscplx** allows for complex arguments.

Format

```
h = hessp(&f, x0);
```

Input

$\&f$	pointer to a single-valued function $f(x)$, defined as a procedure, taking a single $K \times 1$ vector argument ($f: K \times 1 \rightarrow 1 \times 1$); $f(x)$ may be defined in terms of global arguments in addition to x .
$x0$	$K \times 1$ vector specifying the point at which the Hessian of $f(x)$ is to be computed.

Output

h $K \times K$ matrix of second derivatives of f with respect to x at x_0 ; this matrix will be symmetric.

Remarks

This procedure requires $K*(K+1)/2$ function evaluations. Thus if K is large, it may take a long time to compute the Hessian matrix.

No more than 3-4 digit accuracy should be expected from this function, though it is possible for greater accuracy to be achieved with some functions.

It is important that the function be properly scaled, in order to obtain greatest possible accuracy. Specifically, scale it so that the first derivatives are approximately the same size. If these derivatives differ by more than a factor of 100 or so, the results can be meaningless.

Example

```
x = { 1, 2, 3 };  
  
proc g(b);  
  ret( exp(x'b) );  
endp;  
  
b0 = { 3, 2, 1 };  
h = hessp(&g,b0);
```

The resulting matrix of second partial derivatives of $g(b)$ evaluated at $b=b_0$ is:

```
      22026.865   44053.686   66080.596  
h =  44053.686   88107.753  132161.059  
      66080.596  132161.059  198240.695
```

h

hist

Source

hessp.src

See Also

[gradp](#), [gradcplx](#)

hist

Purpose

Computes and graphs a frequency histogram for a vector. The actual frequencies are plotted for each category. Note: this function is for use with the deprecated PQG graphics. **plotHist** instead.

Library

pgraph

Format

```
{ b, m, freq } = hist(x, v);
```

Input

x	Mx1 vector of data.
v	Nx1 vector, the breakpoints to be used to compute the frequencies
	- or -
	scalar, the number of categories.

Output

<i>b</i>	Px1 vector, the breakpoints used for each category.
<i>m</i>	Px1 vector, the midpoints of each category.
<i>freq</i>	Px1 vector of computed frequency counts.

Remarks

If a vector of breakpoints is specified, a final breakpoint equal to the maximum value of *x* will be added if the maximum breakpoint value is smaller.

If a number of categories is specified, the data will be divided into *v* evenly spaced categories.

Each time an element falls into one of the categories specified in *b*, the corresponding element of *freq* will be incremented by one. The categories are interpreted as follows:

```
freq[1] =          x < b[1]
freq[2] = b[1]    < x < b[2]
freq[3] = b[2]    < x < b[3]
.
.
.
freq[P] = b[P-1]  < x < b[P]
```

Example

```
library pgraph;
x = rndn(5000,1);
{ b,m,f } = hist(x,20);
```

Source

phist.src

histf

See Also

[histp](#), [histf](#), [bar](#)

histf

Purpose

Graphs a histogram given a vector of frequency counts. Note: This function is for use with the deprecated PQG graphics. Use **plotSetHistF** instead.

Library

pgraph

Format

```
histf(f, c);
```

Input

<i>f</i>	Nx1 vector, frequencies to be graphed.
<i>c</i>	Nx1 vector, numeric labels for categories. If this is a scalar 0, a sequence from 1 to rows (<i>f</i>) will be created.

Remarks

The axes are not automatically labeled. Use **xlabel** for the category axis and **ylabel** for the frequency axis.

Source

phist.src

See Also

[hist](#), [bar](#), [xlabel](#), [ylabel](#)

histp

Purpose

Computes and graphs a percent frequency histogram of a vector. The percentages in each category are plotted.

Library

pgraph

Format

$\{ b, m, freq \} = \text{histp}(x, v);$

Input

x	Mx1 vector of data.
v	Nx1 vector, the breakpoints to be used to compute the frequencies
- or -	
	scalar, the number of categories.

Output

b	Px1 vector, the breakpoints used for each category.
m	Px1 vector, the midpoints of each category.
$freq$	Px1 vector of computed frequency counts. This is the vector of counts, not percentages.

hsec

Remarks

If a vector of breakpoints is specified, a final breakpoint equal to the maximum value of x will be added if the maximum breakpoint value is smaller.

If a number of categories is specified, the data will be divided into v evenly spaced categories.

Each time an element falls into one of the categories specified in b , the corresponding element of $freq$ will be incremented by one. The categories are interpreted as follows:

```
freq[1] =          x < b[1]
freq[2] = b[1]    < x < b[2]
freq[3] = b[2]    < x < b[3]
.
.
.
freq[P] = b[P-1] < x < b[P]
```

Source

phist.src

See Also

[hist](#), [histf](#), [bar](#)

hsec

Purpose

Returns the number of hundredths of a second since midnight.

Format

```
y = hsec;
```

Output

y scalar, hundredths of a second since midnight.

Remarks

The number of hundredths of a second since midnight can also be accessed as the [4,1] element of the vector returned by the **date** function.

Example

```
x = randu(1000,1000);  
tStart = hsec;  
  
y = x*x;  
tTotal = hsec-tEnd;
```

In this example, **hsec** is used to time a 1000x1000 multiplication in **GAUSS**. A 1000x1000 matrix, *x*, is created, and the current time, in hundredths of a second since midnight, is stored in the variable *tStart*. Then the multiplication is carried out. Finally, *tStart* is subtracted from **hsec** to give the time difference which is assigned to *tTotal*.

See Also

[date](#), [time](#), [timestr](#), [ethsec](#), [etstr](#)

if, else, elseif

i

if, else, elseif

Purpose

Controls program flow with conditional branching.

Format

```
if scalar_expression;  
    list of statements;  
elseif scalar_expression;  
    list of statements;  
elseif scalar_expression;  
    list of statements;  
else;  
    list of statements;  
endif;
```

Remarks

scalar_expression is any expression that returns a scalar. It is TRUE if it is not zero, and FALSE if it is zero.

A *list of statements* is any set of **GAUSS** statements.

GAUSS will test the expression after the **if** statement. If it is TRUE (nonzero), then the first list of statements is executed. If it is FALSE (zero), then **GAUSS** will move to the expression after the first **elseif** statement, if there is one, and test it. It will keep testing expressions and will execute the first list of statements that corresponds to a TRUE expression. If no expression is TRUE, then the list of statements following

the `else` statement is executed. After the appropriate list of statements is executed, the program will go to the statement following the `endif` and continue on.

`if` statements can be nested.

One `endif` is required per `if` statement. If an `else` statement is used, there may be only one per `if` statement. There may be as many `elseif`'s as are required. There need not be any `elseif`'s or any `else` statement within an `if` statement.

Note the semicolon after the `else` statement.

Example

```
if x < 0;
    y = -1;
elseif x > 0;
    y = 1;
else;
    y = 0;
endif;
```

See Also

[do](#)

imag

Purpose

Returns the imaginary part of x .

Format

```
zi = imag(x);
```

#include

Input

x	NxK matrix or N-dimensional array.
-----	------------------------------------

Output

zi	NxK matrix or N-dimensional array, the imaginary part of x .
------	--

Remarks

If x is real, zi will be an NxK matrix or N-dimensional array of zeros.

Example

```
x = { 4i 9      3,  
      2 5-6i 7i };  
y = imag(x);
```

```
y = 4      0      0  
     0     -6      7
```

See Also

[complex](#), [real](#)

#include

Purpose

Inserts code from another file into a **GAUSS** program.

Format

```
#include filename  
#include "filename"
```

Remarks

filename can be any legitimate file name.

This command makes it possible to write a section of general-purpose code, and insert it into other programs.

The code from the `#include`'d file is inserted literally as if it were merged into that place in the program with a text editor.

If a path is specified for the file, then no additional searching will be attempted if the file is not found.

If a path is not specified, the current directory will be searched first, then each directory listed in *src_path*. *src_path* is defined in *gauss.cfg*.

```
#include  
/gauss/myprog.prc  
#include myprog.prc
```

No additional search will be made if the file is not found.

The directories listed in *src_path* will be searched for *myprog.prc* if the file is not found in the current directory.

Compile time errors will return the line number and the name of the file in which they occur. For execution time errors, if a program is compiled with `#lineson`, the line number and name of the file where the error occurred will be printed. For files that have been `#include`'d this reflects the actual line number within the `#include`'d file. See `#lineson` for a more complete discussion of the use of and the validity of line numbers when debugging.

indcv

Example

```
#include
"/gauss/inc/cond.inc"
```

The command will cause the code in the program `cond.inc` to be merged into the current program at the point at which this statement appears.

See Also

[run](#), [lineson](#)

indcv

Purpose

Checks one character vector against another and returns the indices of the elements of the first vector in the second vector.

Format

```
z = indcv(what, where);
```

Input

<i>what</i>	Nx1 character vector which contains the elements to be found in vector <i>where</i> .
<i>where</i>	Mx1 character vector to be searched for matches to the elements of <i>what</i> .

Output

<i>z</i>	Nx1 vector of integers containing the indices of the
----------	--

corresponding element of *what* in *where*.

Remarks

If no matches are found for any of the elements in *what*, then the corresponding elements in the returned vector are set to the **GAUSS** missing value code.

Both arguments will be forced to uppercase before the comparison.

If there are duplicate elements in *where*, the index of the first match will be returned.

Example

```
let newVars = YEARS BONUS GENDER;
let what = AGE PAY SEX;
let where = AGE SEX JOB DATE PAY;

//Return the indices in 'where' of the items in 'what'
z = indcv(what,where);

//Replace AGE, PAY, SEX with YEARS, BONUS, GENDER
where[z] = newVars;
```

After the code above:

	YEARS	
	GENDER	1
where =	JOB	z = 5
	DATE	2
	BONUS	

See Also

[indnv](#), [indsav](#)

indexcat

indexcat

Purpose

Returns the indices of the elements of a vector which fall into a specified category

Format

$y = \text{indexcat}(x, \ v);$

Input

x	Nx1 vector.
v	scalar or 2x1 vector. If scalar, the function returns the indices of all elements of x equal to v . If 2x1, then the function returns the indices of all elements of x that fall into the range: <div>$v[1] < x \leq v[2]$</div> If v is scalar, it can contain a single missing to specify the missing value as the category.

Output

—	y	Lx1 vector, containing the indices of the elements of x which fall into the category defined by v . It will contain error code 13 if there are no elements in this category.
---	-----	--

Remarks

Use a loop to pull out indices of multiple categories.

Example

```
let x = 1.0 4.0 3.3 4.2 6.0 5.7 8.1 5.5;
let v = 4 6;
indx = indexcat(x,v);

inBds = x[indx]
```

```
      4      4.20
indx = 5  inBds = 6.00
      6      5.70
      8      5.50
```

indices

Purpose

Processes a set of variable names or indices and returns a vector of variable names and a vector of indices.

Format

```
{ name, indx } = indices(dataset, vars);
```

Input

<i>dataset</i>	string, the name of the data set.
<i>vars</i>	Nx1 vector, a character vector of names or a numeric vector of column indices.

indices

If scalar 0, all variables in the data set will be selected.

Output

<i>name</i>	Nx1 character vector, the names associated with <i>vars</i> .
<i>indx</i>	Nx1 numeric vector, the column indices associated with <i>vars</i> .

Remarks

If an error occurs, **indices** will either return a scalar error code or terminate the program with an error message, depending on the **trap** state. If the low order bit of the trap flag is 0, **indices** will terminate with an error message. If the low order bit of the trap flag is 1, **indices** will return an error code. The value of the trap flag can be tested with **trapchk**; the return from **indices** can be tested with **scalerr**. You only need to check one argument; they will both be the same. The following error codes are possible:

1	Can't open dataset.
2	Index of variable out of range, or undefined data set variables.

Source

indices.src

indices2

Purpose

Processes two sets of variable names or indices from a single file. The first is a single variable and the second is a set of variables. The first must not occur in the second set and all must be in the file.

Format

```
{ name1, indx1, name2, indx2 } = indices2(dataset, var1, var2);
```

Input

<i>dataset</i>	string, the name of the data set.
<i>var1</i>	string or scalar, variable name or index. This can be either the name of the variable, or the column index of the variable. If null or 0, the last variable in the data set will be used.
<i>var2</i>	Nx1 vector, a character vector of names or a numeric vector of column indices. If scalar 0, all variables in the data set except the one associated with <i>var1</i> will be selected.

Output

<i>name1</i>	scalar character matrix containing the name of the variable associated with <i>var1</i> .
--------------	---

indicesf

<i>indx1</i>	scalar, the column index of <i>var1</i> .
<i>name2</i>	Nx1 character vector, the names associated with <i>var2</i> .
<i>indx2</i>	Nx1 numeric vector, the column indices of <i>var2</i> .

Remarks

If an error occurs, **indices2** will either return a scalar error code or terminate the program with an error message, depending on the **trap** state. If the low order bit of the trap flag is 0, **indices2** will terminate with an error message. If the low order bit of the trap flag is 1, **indices2** will return an error code. The value of the trap flag can be tested with **trapchk**; the return from **indices2** can be tested with **scalerr**. You only need to check one argument; they will all be the same. The following error codes are possible:

- 1 Can't open dataset.
- 2 Index of variable out of range, or undefined data set variables.
- 3 First variable must be a single name or index.
- 4 First variable contained in second set.

Source

indices2.src

indicesf

Purpose

Processes a set of variable names or indices and returns a vector of variable names and a vector of indices.

Format

```
{ name, indx } = indicesf(fp, namein, indxin);
```

Input

<i>fp</i>	scalar, file handle of an open data set.
<i>namein</i>	Nx1 string array, names of selected columns in the data set. If set to a null string, columns are selected using <i>indxin</i>
<i>indxin</i>	Nx1 vector, indices of selected columns in the data set. If set to 0, columns are selected using <i>namein</i> .

Output

<i>name</i>	Nx1 string array, the names of the selected columns.
<i>indx</i>	Nx1 vector, the indices of the selected columns.

Remarks

If *namein* is a null string and *indxin* is 0, all columns of the data set will be selected.

If an error occurs, *indx* will be set to a scalar error code. The following error codes are possible:

- | | |
|---|----------------------|
| 1 | Can't open data file |
| 2 | Variable not found |

indicesfn

3 Indices outside of range of columns

Source

`indices.src`

See Also

[indicesfn](#), [indices](#)

indicesfn

Purpose

Processes a set of variable names or indices and returns a vector of variable names and a vector of indices.

Format

```
{ name, indx } = indicesfn(dataset, namein, indxin);
```

Input

<i>dataset</i>	string, name of the data set.
<i>namein</i>	Nx1 string array, names of selected columns in the data set. If set to a null string, columns are selected using <i>indxin</i>
<i>indxin</i>	Nx1 vector, indices of selected columns in the data set. If set to 0, columns are selected using <i>namein</i> .

Output

<i>name</i>	Nx1 string array, the names of the selected columns.
<i>indx</i>	Nx1 vector, the indices of the selected columns.

Remarks

If *namein* is a null string and *indxin* is 0, all columns of the data set will be selected.

If an error occurs, *indx* will be set to a scalar error code. The following error codes are possible:

- | | |
|---|-------------------------------------|
| 1 | Can't open data file |
| 2 | Variable not found |
| 3 | Indices outside of range of columns |

Source

`indices.src`

See Also

[indicesf](#), [indices](#)

indnv

Purpose

Checks one numeric vector against another and returns the indices of the elements of the first vector in the second vector.

indnv

Format

```
z = indnv(what, where);
```

Input

<i>what</i>	Nx1 numeric vector which contains the values to be found in vector <i>where</i> .
<i>where</i>	Mx1 numeric vector to be searched for matches to the values in <i>what</i> .

Output

<i>z</i>	Nx1 vector of integers, the indices of the corresponding elements of <i>what</i> in <i>where</i> .
----------	--

Remarks

If no matches are found for any of the elements in *what*, then those elements in the returned vector are set to the **GAUSS** missing value code.

If there are duplicate elements in *where*, the index of the first match will be returned.

Example

```
what = { 8, 7, 3 };  
where = { 2, 7, 8, 4, 3 };  
z = indnv(what,where);  
  
      3  
z = 2  
      5
```

indsav

Purpose

Checks one string array against another and returns the indices of the first string array in the second string array.

Format

```
indx = indsav(what, where);
```

Input

<i>what</i>	Nx1 string array which contains the values to be found in vector <i>where</i> .
<i>where</i>	Mx1 string array to be searched for the corresponding elements of <i>what</i> .

Output

<i>indx</i>	Nx1 vector of indices, the values of <i>what</i> in <i>where</i> .
-------------	--

Remarks

If no matches are found, those elements in the returned vector are set to the **GAUSS** missing value code.

If there are duplicate elements in *where*, the index of the first match will be returned.

integrate1d

integrate1d

Purpose

Integrates a user-defined function, using adaptive quadrature, over a user defined interval

Format

```
y = integrate1d(&fct, x_min, x_max);
y = integrate1d(&fct, x_min, x_max, ...);
y = integrate1d(&fct, x_min, x_max, ctl);
y = integrate1d(&fct, x_min, x_max, ..., ctl);
```

Input

<i>&fct</i>	scalar, pointer to the procedure containing the function to be integrated
<i>x_min</i>	scalar, starting point for the integration
<i>x_max</i>	scalar, ending point for the integration
<i>...</i>	Optional input, a variable number of extra arguments to pass to the user function. These arguments will be passed to the user function untouched.
<i>ctl</i>	Optional input, an instance of an integrateControl structure with members
<i>ctl.subDivisions</i>	scalar, maximum number of divisions of the region (<i>x_min</i> , <i>x_max</i>)
<i>ctl.absTol</i>	scalar, absolute accuracy requested.
<i>ctl.relTol</i>	scalar, relative accuracy requested.

Output

y scalar, the estimated integral of $\mathbf{f}(x)$ evaluated over the interval (x_min, x_max)

Examples

Example 1: Basic Example

Calculate the integral $\int_0^3 \frac{1}{x+1} dx$

```
//Define procedure to be integrated
proc (1) = fct(x);
    retp(1 ./ (x + 1));
endp;

//Calculate integral for procedure 'fct', from 0 - 3
ans = integrate1d(&fct, 0, 3);
```

will result in:

```
ans = 1.3862943611
```

Example 2: Passing extra arguments to the user function

Calculate the integral $\int_{-1000}^{1000} e^{-\frac{x^2}{2 \times a}} dx$, $a = 3$

```
//Define procedure to be integrated
proc (1) = myProc(x, var);
    retp(exp(-(x .* x) / (2 .* var)));
endp;

//Define limits of integration
x_min = -1000;
```

integrate1d

```
x_max = 1000;  
  
//Define extra argument for procedure 'myProc'  
a = 3;  
  
ans = integrate1d(&myProc, x_min, x_max, a);
```

will result in:

```
ans = 4.3416075273
```

Example 3: Bound at negative infinity

Calculate the integral $\int_{-\infty}^0 \frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{(x-\mu)^2}{2\sigma^2}} dx$

```
//Define procedure to be integrated  
proc (1) = myPdfn(x, mu, sigma);  
    retp(pdfn((x - mu) ./ sigma) ./ sigma);  
endp;  
  
//Set bounds of integration to be (-Inf, 0)  
x_min = __INFN;  
x_max = 0;  
  
//Extra inputs for user function  
mu = 0.33;  
sigma = 7;  
  
ans = integrate1d(&myPdfn, x_min, x_max, mu, sigma);
```

will result in:

```
ans = 0.481199685115
```


Example 4: Using a control structure

Calculate the integral $\int_{-\infty}^0 \frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{(x-\mu)^2}{2\sigma^2}} dx$

```
//Define procedure to be integrated
proc (1) = myPdfn(x, mu, sigma);
    retp(pdfn((x - mu) ./ sigma) ./ sigma);
endp;

//Set bounds of integration to be (0, +Inf)
x_min = 0;
x_max = __INFP;

//Extra inputs for user function
mu = 0.33;
sigma = 7;

//Declare instance of 'integrateControl' structure
//and fill with default values
struct integrateControl ctl;
ctl = integrateControlCreate();

//Lower required tolerance for faster return
ctl.absTol = 1e-2;

ans = integrate1d(&myPdfn, x_min, x_max, mu, sigma, ctl);
```

will result in:

```
ans = 0.518798668212
```

Remarks

The user-provided function must be able to accept a vector of scalar values and return

integrate1d

a vector of outputs. Make sure to use the element by element operators (`.* ./`) instead of the overloaded matrix operators (`* /`). For example, the following procedure:

```
proc (1) = myProc(x);  
    local ret;  
    ret = x / (x * x);  
    retp(ret);  
endp;
```

will work as expected for a scalar input. For example:

```
a = 2;  
b = 3;  
c = myProc(a);  
d = myProc(b);
```

will assign `c` to be equal to 0.5 and `d` to be equal to 0.334. However, if we pass in a vector like this:

```
a = { 2,  
      3 };  
c = myProc(a);
```

we will cause an the error **matrices not conformable** when we try to multiply the incoming 2x1 vector times itself inside of `myProc`. To avoid this, we simply need to change the operators `*` and `/` to the element-by-element versions by prepending the operator with a dot like this:

```
proc (1) = myProc(x);  
    local ret;  
    ret = x ./ (x .* x);  
    retp(ret);  
endp;
```

Source

`integrate.src`

See Also

[inthpControlCreate](#), [inthp2](#), [inthp3](#), [inthp4](#)

intgrat2

Purpose

Integrates the following double integral, using user-defined functions **f**, **g₁** and **g₂** and scalars *a* and *b*:

$$\int_a^b \int_{g_2(x)}^{g_1(x)} f(x, y) dy dx$$

Format

```
y = intgrat2(&f, x1, g1);
```

Input

<i>&f</i>	scalar, pointer to the procedure containing the function to be integrated.
<i>x1</i>	2x1 or 2xN matrix, the limits of <i>x</i> . These must be scalar limits.
<i>g1</i>	2x1 or 2xN matrix of function pointers, the limits of <i>y</i> . For <i>x1</i> and <i>g1</i> , the first row is the upper limit and the second row is the lower limit. N integrations are computed.

intgrat2

Global Input

<code>_intord</code>	scalar, the order of the integration. The larger <code>_intord</code> , the more precise the final result will be. <code>_intord</code> may be set to 2, 3, 4, 6, 8, 12, 16, 20, 24, 32, 40. Default = 12.
----------------------	---

Output

<code>y</code>	$N \times 1$ vector of the estimated integral(s) of $f(x, y)$, evaluated between the limits given by <code>x1</code> and <code>g1</code> .
----------------	---

Remarks

The user-defined functions specified by **f** and **g1** must either

1. Return a scalar constant

- or -
2. Return a vector of function values. **intgrat2** will pass to user-defined functions a vector or matrix for `x` and `y` and expect a vector or matrix to be returned. Use `.*` and `./` instead of `*` and `/`.

Example

```
proc (1) = f(x,y);  
    retp(cos(x) + 1) .* (sin(y) + 1);  
endp;  
  
proc (1) = g1(x);  
    retp(sqrt(1-x^2));  
endp;
```

```

proc (1) = g2(x);
  retp(0);
endp;

x1 = 1|-1;
g0 = &g1|&g2;
_intord = 40;
y = intgrat2(&f,x1,g0);

```

This will integrate the function

$$f(x,y) = (\cos(x)+1)(\sin(y)+1)$$

over the upper half of the unit circle. Note the use of the `. *` operator instead of just `*` in the

definition of $f(x,y)$. This allows f to return a vector or matrix of function values.

Source

intgrat.src

Globals

`_intord`, `_intq12`, `_intq16`, `_intq2`, `_intq20`, `_intq24`, `_intq3`, `_intq32`, `_intq4`, `_intq40`, `_intq6`, `_intq8`

See Also

[intgrat3](#), [intquad1](#), [intquad2](#), [intquad3](#), [intsimp](#)

intgrat3

Purpose

Integrates the following triple integral, using user-defined functions and scalars for bounds:

integrat3

$$\int_a^b \int_{g_2(x)}^{g_1(x)} \int_{h_2(x,y)}^{h_1(x,y)} f(x,y,z) dz dy dx$$

Format

```
y = integrat3(&f, xl, gl, hl);
```

Input

<i>&f</i>	scalar, pointer to the procedure containing the function to be integrated. <i>f</i> is a function of (<i>x</i> , <i>y</i> , <i>z</i>).
<i>xl</i>	2x1 or 2xN matrix, the limits of <i>x</i> . These must be scalar limits.
<i>gl</i>	2x1 or 2xN matrix of function pointers. These procedures are functions of <i>x</i> .
<i>hl</i>	2x1 or 2xN matrix of function pointers. These procedures are functions of <i>x</i> and <i>y</i> .

For *xl*, *gl*, and *hl*, the first row is the upper limit and the second row is the lower limit. N integrations are computed.

Global Input

<i>_intord</i>	scalar, the order of the integration. The larger <i>_intord</i> , the more precise the final result will be. <i>_intord</i> may be set to 2, 3, 4, 6, 8, 12, 16, 20, 24, 32, 40. Default = 12.
----------------	---

Output

y Nx1 vector of the estimated integral(s) of $f(x,y,z)$ evaluated between the limits given by $x1$, $g1$ and $h1$.

Remarks

User-defined functions f , and those used in $g1$ and $h1$ must either:

1. Return a scalar constant
- or -
2. Return a vector of function values. **intgrat3** will pass to user-defined functions a vector or matrix for x and y and expect a vector or matrix to be returned. Use $.*$ and $./$ operators instead of just $*$ and $/$.

Example

```
proc f(x,y,z);
  retp(2);
endp;

proc g1(x);
  retp(sqrt(25-x^2));
endp;

proc g2(x);
  retp(-g1(x));
endp;

proc h1(x,y);
  retp(sqrt(25 - x^2 - y^2));
endp;
```

inthp1

```
proc h2(x,y);  
    retp(-h1(x,y));  
endp;  
  
x1 = 5|-5;  
g0 = &g1|&g2;  
h0 = &h1|&h2;  
  
_intord = 40;  
  
y = intgrat3(&f,x1,g0,h0);
```

This will integrate the function $f(x,y,z)$ over the sphere of radius 5. The result will be approximately twice the volume of a sphere of radius 5.

Source

intgrat.src

Globals

`_intord, _intq12, _intq16, _intq2, _intq20, _intq24, _intq3, _
_intq32, _intq4, _intq40, _intq6, _intq8`

See Also

[intgrat2](#), [intquad1](#), [intquad2](#), [intquad3](#), [intsimp](#)

inthp1

Purpose

Integrates a user-defined function over an infinite interval.

Include

inthp.sdf

Format

```
y = inthp1(&f, pds, ctl);
```

Input

<i>&f</i>	scalar, pointer to the procedure containing the function to be integrated.	
<i>pds</i>	scalar, pointer to instance of a DS structure. The members of the DS are:	
	<i>pds->dataMatrix</i>	NxK matrix.
	<i>pds->dataArray</i>	NxKxL... array.
	<i>pds->vnames</i>	string array.
	<i>pds->dsname</i>	string.
	<i>pds->type</i>	scalar.
	The contents, if any, are set by the user and are passed by inthp1 to the user-provided function without modification.	
<i>ctl</i>	instance of an inthpControl structure with members	
	<i>ctl.maxEvaluations</i>	scalar, maximum number of function evaluations, default = 1e5;
	<i>ctl.p</i>	scalar, termination parameter
	0	heuristic termination, default.
	1	deterministic

inthp1

		termination with infinity norm.
	2,...	deterministic termination with p-th norm.
<i>ctl.d</i>	scalar termination parameter	
	1	if heuristic termination
	$0 < \textit{ctl.d} < \pi/2$	if deterministic termination
<i>ctl.eps</i>	scalar, relative error bound.	
	Default = 1e-6.	
A default <i>ctl</i> can be generated by calling inthpControlCreate .		

Output

<i>y</i>	scalar, the estimated integral of $\mathbf{f}(x)$ evaluated over the interval $(-\infty, +\infty)$.
----------	--

Remarks

The user-provided function must have the following format	
	<code>\mathbf{f}(struct DS *pds, x)</code>
where	

pds scalar, pointer to an instance of a **DS** structure.
x scalar, value at which integral will be evaluated.

If *ctl.d* can be specified (see Sikorski and Stenger, 1984), deterministic termination can be specified and accuracy guaranteed. if not, the heuristic method can be used and the value of *ctl.d* is disregarded.

The pointer to the instance of the data structure, *pds*, is passed untouched to the user-provided procedure computing the function to be integrated. Any information needed by that function can be put into that data structure.

Example

```
proc fct(struct DS *pds, x);
local var;
    var = pds->dataMatrix;
    retp( exp( -(x*x) / (2*var) ));
endp;

struct DS d0;
struct DS *pds;
variance = 3;
pds = &d0;
d0.dataMatrix = variance;

struct inthpControl c0;
c0 = inthpControlCreate;

r = inthp1(&fct,pds,c0);

format /ld 16,10;
print r;
print sqrt(2*pi*variance);
```

results in the following output:

inthp2

```
4.3416075273
4.3416075273
```

References

1. "Optimal Quadratures in H_p Spaces" by K. Sikorski and F. Stenger, *ACM Transactions on Mathematical Software*, 10(2):140-151, June 1984.

Source

`inthp.src`

See Also

[inthpControlCreate](#), [inthp2](#), [inthp3](#), [inthp4](#)

inthp2

Purpose

Integrates a user-defined function over the $[a, +\infty)$ interval.

Include

`inthp.sdf`

Format

```
y = inthp2(&f, pds, ctl, a);
```

Input

<code>&f</code>	scalar, pointer to the procedure containing the function to be integrated.
<code>pds</code>	scalar, pointer to instance of a DS structure. The members of the DS are:

	<i>pds->dataMatrix</i>	NxK matrix.
	<i>pds->dataArray</i>	NxKxL... array.
	<i>pds->vnames</i>	string array.
	<i>pds->dsname</i>	string.
	<i>pds->type</i>	scalar.
	The contents, if any, are set by the user and are passed by inthp1 to the user-provided function without modification.	
<i>ctl</i>	instance of an inthpControl structure with members	
	<i>ctl.maxEvaluations</i>	scalar, maximum number of function evaluations, default = 1e5;
	<i>ctl.p</i>	scalar, termination parameter
	0	heuristic termination, default.
	1	deterministic termination with infinity norm.
	2,...	deterministic termination with p-th norm.
	<i>ctl.d</i>	scalar termination parameter
	1	if heuristic termination
	$0 < \textit{ctl.d} < \pi/2$	if deterministic termination

inthp2

	<i>ctl.eps</i>	scalar, relative error bound. Default = 1e-6.
		A default <i>ctl</i> can be generated by calling inthpControlCreate .
<i>a</i>		1xN vector, lower limits of integration.

Output

<i>y</i>	Nx1 vector, the estimated integrals of f (<i>x</i>) evaluated over the interval [<i>a</i> ,+∞).
----------	---

Remarks

	The user-provided function must have the following format
	<pre>f(struct DS *<i>pds</i>, <i>x</i>)</pre>
	where
<i>pds</i>	scalar, pointer to an instance of a DS structure.
<i>x</i>	scalar, value at which integral will be evaluated.
	If <i>ctl.d</i> can be specified (see Sikorski and Stenger, 1984), deterministic termination can be specified and accuracy guaranteed. If not, the heuristic method can be used and the value of <i>ctl.d</i> is disregarded.
	The pointer to the instance of the data structure, <i>pds</i> , is passed untouched to the user-provided procedure computing the function to be integrated. Any information needed by that function can be put into that data structure.

Example

	<pre>#include inthp.sdf</pre>
--	-------------------------------

```

proc normal(struct DS *pd0, x);
    local var;
    var = pd0->dataMatrix;
    ret( (1/sqrt(2*pi*var))*exp( -(x*x) / (2*var) ));
endp;

struct DS d0;
struct DS *pd0;

pd0 = &d0;

struct inthpControl c0;
c0 = inthpControlCreate;

lim = 2;

c0.d = pi/4;
c0.p = 2;

var = 1;
d0.dataMatrix = var;

r = inthp2(&normal,pd0,c0,lim);

format /ld 16,10;
print r;
print cdfnc(2);

```

produces the following output:

```

0.0227501281
0.0227501319

```

inthp3

References

1. "Optimal Quadratures in H_p Spaces" by K. Sikorski and F. Stenger, *ACM Transactions on Mathematical Software*, 10(2):140-151, June 1984.

Source

`inthp.src`

See Also

[inthpControlCreate](#), [inthp1](#), [inthp3](#), [inthp4](#)

inthp3

Purpose

Integrates a user-defined function over the $[a, +\infty)$ interval that is oscillatory.

Include

`inthp.sdf`

Format

```
y = inthp3(&f, pds, ctl, a);
```

Input

$\&f$	scalar, pointer to the procedure containing the function to be integrated.
pds	scalar, pointer to instance of a DS structure. The members of the DS are: <div><div>$pds->dataMatrix$</div><div>NxK matrix.</div><div>$pds->dataArray$</div><div>NxKxL... array.</div></div>

<i>ctl</i>	<i>pds->vnames</i>	string array.
	<i>pds->dsname</i>	string.
	<i>pds->type</i>	scalar.
	The contents, if any, are set by the user and are passed by inthp1 to the user-provided function without modification.	
	instance of an inthpControl structure with members	
	<i>ctl.maxEvaluations</i>	scalar, maximum number of function evaluations, default = 1e5;
	<i>ctl.p</i>	scalar, termination parameter
	0	heuristic termination, default.
	1	deterministic termination with infinity norm.
	2,...	deterministic termination with p-th norm.
	<i>ctl.d</i>	scalar termination parameter
	1	if heuristic termination
	$0 < \textit{ctl.d} < \pi/2$	if deterministic termination
	<i>ctl.eps</i>	scalar, relative error bound. Default = 1e-6.

inthp3

a	A default <i>ctl</i> can be generated by calling inthpControlCreate . 1xN vector, lower limits of integration.
-----	--

Output

y	Nx1 vector, the estimated integrals of $\mathbf{f}(x)$ evaluated over the interval $[a, +\infty)$.
-----	---

Remarks

This procedure is designed especially for oscillatory functions.

The user-provided function must have the following format

```
 $\mathbf{f}(\text{struct DS } *pds, x)$ 
```

where

pds	scalar, pointer to an instance of a DS structure.
x	scalar, value at which integral will be evaluated.

If *ctl.d* can be specified (see Sikorski and Stenger, 1984), deterministic termination can be specified and accuracy guaranteed. if not, the heuristic method can be used and the value of *ctl.d* is disregarded.

The pointer to the instance of the data structure, *pds*, is passed untouched to the user-provided procedure computing the function to be integrated. Any information needed by that function can be put into that data structure.

Example

```
#include inthp.sdf
```

```

proc fct(struct DS *pd0, x);
    local m,a;
    m = pd0->dataMatrix[1];
    a = pd0->dataMatrix[2];
    retp( exp(-a*x)*cos(m*x));
endp;

struct DS d0;
struct DS *pd0;

struct inthpControl c0;
c0 = inthpControlCreate;

c0.p = 2;
c0.d = pi/3;

m = 2;
a = 1;
pd0 = &d0;
d0.dataMatrix = m | a;

lim = 0;

r = inthp3(&fct,pd0,c0,lim);

format /ld 16,10;
print r;
print a/(a*a + m*m);

```

produces the following output:

```

0.2000000000
0.2000000000

```

inthp4

References

- 1. "Optimal Quadratures in H_p Spaces" by K. Sikorski and F. Stenger, *ACM Transactions on Mathematical Software*, 10(2):140-151, June 1984.

Source

inthp.src

See Also

[inthpControlCreate](#), [inthp1](#), [inthp2](#), [inthp4](#)

inthp4

Purpose

Integrates a user-defined function over the $[a, b]$ interval.

Include

inthp.sdf

Format

$y = \text{inthp4}(\&f, pds, ctl, c);$

Input

$\&f$	scalar, pointer to the procedure containing the function to be integrated.
pds	scalar, pointer to instance of a DS structure. The members of the DS are: <div><div>$pds->dataMatrix$</div><div>$pds->dataArray$</div></div> <div><div>NxK matrix.</div><div>NxKxL... array.</div></div>

	<i>pds->vnames</i>	string array.
	<i>pds->dsname</i>	string.
	<i>pds->type</i>	scalar.
	The contents, if any, are set by the user and are passed by inthp1 to the user-provided function without modification.	
<i>ctl</i>	instance of an inthpControl structure with members	
	<i>ctl.maxEvaluations</i>	scalar, maximum number of function evaluations, default = 1e5;
	<i>ctl.p</i>	scalar, termination parameter
	0	heuristic termination, default.
	1	deterministic termination with infinity norm.
	2,...	deterministic termination with p-th norm.
	<i>ctl.d</i>	scalar termination parameter
	1	if heuristic termination
	$0 < \textit{ctl.d} < \pi/2$	if deterministic termination
	<i>ctl.eps</i>	scalar, relative error bound. Default = 1e-6.
	A default <i>ctl</i> can be generated by calling inthpControlCreate .	
<i>c</i>	2×N vector, upper and lower limits of integration, the first row contains upper limits and the second row the lower.	

Output

y	$N \times 1$ vector, the estimated integrals of $\mathbf{f}(x)$ evaluated over the interval $[a, b]$.
-----	--

Remarks

The user-provided function must have the following format

```
f(struct DS *pds, x)
```

where

pds	scalar, pointer to an instance of a DS structure.
x	scalar, value at which integral will be evaluated.

If $ctl.d$ can be specified (see Sikorski and Stenger, 1984), deterministic termination can be specified and accuracy guaranteed. if not, the heuristic method can be used and the value of $ctl.d$ is disregarded.

The pointer to the instance of the data structure, pds , is passed untouched to the user-provided procedure computing the function to be integrated. Any information needed by that function can be put into that data structure.

Example

```
#include inths.sdf

proc fct(struct DS *pd0, x);
  local a,b,c;
  a = pd0->dataMatrix[1];
  b = pd0->dataMatrix[2];
  c = pd0->dataMatrix[3];
  retp( 1/sqrt(a*x*x + b*x + c));
endp;
```

```

struct DS d0;
struct DS *pd0;

struct inthsControl c0;
c0 = inthsControlCreate;

c0.p = 2;
c0.d = pi/2;

a = -1;
b = -2;
c = 3;
pd0 = &d0;
d0.dataMatrix = a|b|c;

lim = 1 | -1;

r = inths4(&fct,pd0,c0,lim);

format /ld 16,10;
print r;
print pi/2;

```

produces the following output:

```

1.5707962283
1.5707963268

```

References

1. "Optimal Quadratures in H_p Spaces" by K. Sikorski and F. Stenger, *ACM Transactions on Mathematical Software*, 10(2):140-151, June 1984.

Source

inths.src

inthpControlCreate

See Also

[inthpControlCreate](#), [inthp1](#), [inthp2](#), [inthp3](#)

inthpControlCreate

Purpose

Creates default **inthpControl** structure.

Include

`inthp.sdf`

Format

```
c = inthpControlCreate();
```

Output

<i>c</i>	instance of inthpControl structure with members set to default values.
----------	---

Source

`inthp.src`

See Also

[inthp1](#), [inthp2](#), [inthp3](#), [inthp4](#)

intquad1

Purpose

Integrates a specified function using Gauss-Legendre quadrature. A suite of upper and lower bounds may be calculated in one procedure call.

Format

```
y = intquad1(&f, x1);  
y = intquad1(&f, x1, ...);
```

Input

$\&f$	scalar, pointer to the procedure containing the function to be integrated. This must be a function of x .
$x1$	2xN matrix, the limits of x . The first row is the upper limit and the second row is the lower limit. N integrations are computed.
\dots	Optional, a variable number of extra <i>scalar</i> arguments to pass to the user function. These arguments will be passed to the user function untouched.

Global Input

$_intord$	scalar, the order of the integration. The larger $_intord$, the more precise the final result will be. $_intord$ may be set to 2, 3, 4, 6, 8, 12, 16, 20, 24, 32, 40.
------------	--

intquad1

Default = 12.

Output

y

Nx1 vector of the estimated integral(s) of $\mathbf{f}(x)$ evaluated between the limits given by $x1$.

Remarks

The user-defined function \mathbf{f} must return a vector of function values. **intquad1** will pass to the user-defined function a vector or matrix for x and expect a vector or matrix to be returned. Use the `.*` and `./` instead of `*` and `/`.

Example

Example 1: Basic example

This will integrate the function $f(x) = x \sin(x)$ between 0 and 1. Note the use of the `.` * instead of `*`.

```
//Define function to be integrated
proc f(x);
    retp(x.*sin(x));
endp;

//Limits of integration
xlim = { 1, 0 };

//Calculate integral
y = intquad1(&f,xlim);
```

After the code above, `y` should equal:

```
0.30116868
```

Example 2: Passing in additional arguments

```
//Define function to be integrated
//with a second input
proc f(x, a);
    retp(x.*sin(x .* a));
endp;

//Create additional scalar argument 'a'
a = 3.14;

//Limits of integration
xlim = { 1, 0 };
```

intquad2

```
//Calculate integral, passing in extra input
//'a' as the final input to 'intquad1'
y = intquad1(&f, xlim, a);
```

After the code above, `y` should equal:

```
0.31863247
```

Source

`integral.src`

Globals

```
_intord, _intq12, _intq16, _intq2, _intq20, _intq24, _intq3, _
intq32, _intq4, _intq40, _intq6, _intq8
```

See Also

[intsimp](#), [intquad2](#), [intquad3](#), [intgrat2](#), [intgrat3](#)

intquad2

Purpose

Integrates a specified function using Gauss-Legendre quadrature. A suite of upper and lower bounds may be calculated in one procedure call.

Format

```
y = intquad2(&f, xl, yl);
y = intquad2(&f, xl, yl, ...);
```

Input

$\&f$	scalar, pointer to the procedure containing the function to be integrated.
$x1$	2x1 or 2xN matrix, the limits of x .
$y1$	2x1 or 2xN matrix, the limits of y .
\dots	Optional, a variable number of extra <i>scalar</i> arguments to pass to the user function. These arguments will be passed to the user function untouched.

For $x1$ and $y1$, the first row is the upper limit and the second row is the lower limit. N integrations are computed.

Global Input

$_intord$	scalar, the order of the integration. The larger $_intord$, the more precise the final result will be. $_intord$ may be set to 2, 3, 4, 6, 8, 12, 16, 20, 24, 32, 40. Default = 12.
------------	--

Output

y	Nx1 vector of the estimated integral(s) of $f(x,y)$ evaluated between the limits given by $x1$ and $y1$.
-----	---

Examples

Example 1: Basic example

```
//Define function to be integrated
proc f(x,y);
    retp(x .* sin(x + y));
endp;

//Limits of integration
xlim = { 1, 0 };
ylim = { 1, 0 };

//Calculate integral
ans = intquad2(&f, xlim, ylim);
```

After the code above, `ans` should equal:

```
0.42892501
```

Example 2: Multiple integration limits

```
//Define function to be integrated
proc (1) = myProc(x,y);
    retp(x .* sin(x + y));
endp;

//Define multiple integration limits
xlim = { 1 0.5,
        0.5 0 };

ylim = { 1 0.5,
        0.5 0.3 };
```

```
//Calculate integrals
ans = intquad2(&myProc, xlim, ylim);
```

This will integrate the function:

```
myProc(x) = x.*sin(x+y)
```

between $x=0$ and 0.5 , and between $y=0.3$ and 0.5 as well as between $x=0.5$ and 1 , and between $y=0.5$ and 1 .

The returned variable, `ans` should be equal to:

```
0.18352849
0.016593029
```

Example 3: Extra arguments to function

```
//Define function to be integrated that takes an additional
argument
proc f(x,y,a);
    retp(x .* sin(a .* x + y));
endp;

//Limits of integration
xlim = { 1, 0 };
ylim = { 1, 0 };

//Assign extra scalar argument
a = pi/2;

//Calculate integral
ans = intquad2(&f, xlim, ylim, a);
```

After the code above, `ans` should equal:

```
0.44737953
```

intquad3

Remarks

The user-defined function f must return a vector of function values. **intquad2** will pass to user-defined functions a vector or matrix for x and y and expect a vector or matrix to be returned. Use `.*` and `./` instead of `*` and `/`.

intquad2 will expand scalars to the appropriate size. This means that functions can be defined to return a scalar constant. If users write their functions incorrectly (using `*` instead of `.*`, for example), **intquad2** may not compute the expected integral, but the integral of a constant function.

To integrate over a region which is bounded by functions, rather than just scalars, use **intgrat2** or **intgrat3**.

Source

`integral.src`

Globals

`_intord`, `_intq12`, `_intq16`, `_intq2`, `_intq20`, `_intq24`, `_intq3`, `_intq32`, `_intq4`, `_intq40`, `_intq6`, `_intq8`

See Also

[intquad1](#), [intquad3](#), [intsimp](#), [intgrat2](#), [intgrat3](#)

intquad3

Purpose

Integrates a specified function using Gauss-Legendre quadrature. A suite of upper and lower bounds may be calculated in one procedure call.

Format

```
y = intquad3(&f, xl, yl, zl);  
y = intquad3(&f, xl, yl, zl, ...);
```

Input

$\&f$	scalar, pointer to the procedure containing the function to be integrated. f is a function of (x, y, z) .
xl	2x1 or 2xN matrix, the limits of x .
yl	2x1 or 2xN matrix, the limits of y .
zl	2x1 or 2xN matrix, the limits of z .
\dots	Optional, a variable number of extra <i>scalar</i> arguments to pass to the user function. These arguments will be passed to the user function untouched.

For xl , yl , and zl , the first row is the upper limit and the second row is the lower limit. N integrations are computed.

Global Input

$_intord$	scalar, the order of the integration. The larger $_intord$, the more precise the final result will be. $_intord$ may be set to 2, 3, 4, 6, 8, 12, 16, 20, 24, 32, 40. Default = 12.
------------	---

intquad3

Output

y	$N \times 1$ vector of the estimated integral(s) of $f(x,y,z)$ evaluated between the limits given by $x1$, $y1$, and $z1$.
-----	---

Example

Example 1: Basic example

```
//Define function to integrate
proc f(x,y,z);
    retp(sqrt(x.^2 + y.^2 + z.^2));
endp;

//Define limits of integration
xlim = { 1, 0 };
ylim = { 1, 0 };
zlim = { 3, 0 };

//Calculate integral
ans = intquad3(&f,xlim, ylim, zlim);
```

After the code above, *ans* should equal:

5.2994691

Example 2: Passing extra arguments

```
//Define function to integrate which takes an additional
argument
proc f(x,y,z,a);
    retp(sqrt(a .* x.^2 + y.^2 + z.^2));
endp;
```

```
//Define limits of integration
xlim = { 1, 0 };
ylim = { 1, 0 };
zlim = { 3, 0 };

//Define extra scalar argument
a = 3.14;

//Calculate integral, passing in extra scalar argument
ans = intquad3(&f,xlim, ylim, zlim, a);
```

After the code above, *ans* should equal:

5.8969356

Example 3: Multiple limits of integration

```
//Define function to integrate
proc f(x,y,z);
    retp(sqrt(x.^2 + y.^2 + z.^2));
endp;

//Define 3 sets of limits of integration
xlim = { 1, 0 };
ylim = { 1, 0 };
zlim = { 1 2 3,
         0 0 0 };

//Calculate integrals
ans = intquad3(&f,xlim, ylim, zlim);
```

This will integrate the function $f(x) = x*y*z$ over 3 sets of limits, since *zlim* is defined to be a 2x3 matrix. The value of *ans* should be:

intrleav

```
0.96059195
2.6692443
5.2994691
```

Remarks

The user-defined function f must return a vector of function values. **intquad3** will pass to the user-defined function a vector or matrix for x , y and z and expect a vector or matrix to be returned. Use `.*` and `./` instead of `*` and `/`.

intquad3 will expand scalars to the appropriate size. This means that functions can be defined to return a scalar constant. If users write their functions incorrectly (using `*` instead of `.*`, for example), **intquad3** may not compute the expected integral, but the integral of a constant function.

To integrate over a region which is bounded by functions, rather than just scalars, use **intgrat2** or **intgrat3**.

Source

`integral.src`

Globals

`_intord`, `_intq12`, `_intq16`, `_intq2`, `_intq20`, `_intq24`, `_intq3`, `_intq32`, `_intq4`, `_intq40`, `_intq6`, `_intq8`

See Also

[intquad1](#), [intquad2](#), [intsimp](#), [intgrat2](#), [intgrat3](#)

intrleav

Purpose

Interleaves the rows of two files that have been sorted on a common variable to produce a single file sorted on that variable.

Format

```
intrleav(infile1, infile2, outfile, keyvar, keytyp);
```

Input

<i>infile1</i>	string, name of input file 1.
<i>infile2</i>	string, name of input file 2.
<i>outfile</i>	string, name of output file.
<i>keyvar</i>	string, name of key variable; this is the column the files are sorted on.
<i>keytyp</i>	scalar, data type of key variable.
1	numeric key, ascending order
2	character key, ascending order
-1	numeric key, descending order
-2	character key, descending order

Remarks

The two files MUST have exactly the same variables, that is, the same number of columns AND the same variable names. They must both already be sorted on the key column. This procedure will combine them into one large file, sorted by the key variable.

If the inputs are null ("" or 0), the procedure will ask for them.

Source

sortd.src

See Also

[intrleavsa](#)

intrleavsa

intrleavsa

Purpose

Interleaves the rows of two string arrays that have been sorted on a common column.

Format

```
y = intrleavsa(sa1, sa2, ikey);
```

Input

<i>sa1</i>	NxK string array 1.
<i>sa2</i>	MxK string array 2.
<i>ikey</i>	scalar integer, index of the key column the string arrays are sorted on.

Output

<i>y</i>	LxK interleaved (combined) string array.
----------	--

Remarks

The two string arrays MUST have exactly the same number of columns AND have been already sorted on a key column.

This procedure will combine them into one large string array, sorted by the key column.

Source

sortd.src

See Also

[intrleav](#)

intrsect

Purpose

Returns the intersection of two vectors, with duplicates removed.

Format

```
y=intrsect(v1, v2);
y=intrsect(v1, v2, flag);
```

Input

v1	Nx1 vector, or string array.
v2	Mx1 vector, or string array. NOTE: v2 must be the same type as v1
flag	Optional argument, scalar, if 1, v1 and v2 are numeric; if 0, character. Default is flag equal to 1 (numeric).

Output

y	Lx1 vector containing all unique values that are in both v1 and v2, sorted in ascending order.
---	--

Remarks

1. If not matches are found, **intrsect** will return a scalar error code that can be

intrsect

tested for with **scalmiss**.

2. Place smaller vector first for fastest operation.
3. If there are a lot of duplicates within a vector, it is faster to remove them with the function **unique** before calling **intrsect**.

Examples

Example 1: Basic usage, numeric

```
//Subject ID's from study 'a'
id_a = { 3758,
         3773,
         2615,
         2511 };

//Subject ID's from study 'b'
id_b = { 3779,
         3773,
         2001,
         3758,
         1585,
         2511 };

//Find the ID's that are in both groups
id_common = intrsect(id_a, id_b);
```

After the code above, *id_common* is equal to:

```
2511
3758
3773
```

Example 2: Basic usage, string array

```
//Variable names from dataset 'a'
//Create string array with the string
```



```
//vertical concatenation operator ($|)
names_a = "oil" $| "copper" $| "silver" $| "cocoa";

//Variable names from dataset 'b'
names_b = "oil" $| "coffee" $| "cocoa" $| "tea";

//Find the variable names that are in both groups
names_common = intrsect(names_a, names_b);
```

After the code above, *names_common* is equal to:

```
cocoa
oil
```

Example 3: Character vectors

A character vector is different from a string array. A character vector is up to eight characters inside of the element of a numeric matrix.

```
//Variable names from dataset 'a'
//Create character vector array with the
//numeric vertical concatenation operator (|)
names_a = "oil" | "copper" | "silver" | "cocoa";

//Variable names from dataset 'b'
names_b = "oil" | "coffee" | "cocoa" | "tea";

//Set flag to tell 'intrsectsa' to treat input as character
data
flag = 0;

//Find the variable names that are in both groups
names_common = intrsect(names_a, names_b, flag);

//Notice the $ in front of 'names_common'
```

intrsectsa

```
//tells GAUSS to print as character data
print $names_common;
```

The code above, will print the following output:

```
cocoa
oil
```

Source

intrsect.src

See Also

[intrsectsa](#)

intrsectsa

Purpose

Returns the intersection of two string vectors, with duplicates removed. NOTE:
This function is deprecated, use **intrsect** instead.

Format

```
y = intrsectsa(sv1, sv2);
```

Input

<i>sv1</i>	Nx1 or 1xN string vector.
<i>sv2</i>	Mx1 or 1xM string vector.

Output

<code>sy</code>	Lx1 vector containing all unique strings that are in both <code>sv1</code> and <code>sv2</code> , sorted in ascending order.
-----------------	--

Remarks

1. This function is deprecated, use **intrsect** instead.
2. Place smaller vector first for fastest operation.
3. If there are a lot of duplicates it is faster to remove them with `unique` before calling **intrsectsa**.

Example

```
vars_a = "age" $| "weight" $| "bmi";
vars_b = "hdl" $| "ldl" $| "age" $| "bmi" $| "smoking";

shared_vars = intrsectsa(vars_a, vars_b);
print "Both studies reported the following variables:";
print shared_vars;
```

The code above, returns:

```
Both studies reported the following variables:
      age      bmi
```

Source

`intrsect.src`

See Also

[intrsect](#)

intsimp

intsimp

Purpose

Integrates a specified function using Simpson's method with end correction. A single integral is computed in one function call.

Format

```
y = intsimp(&f, x1, tol);
```

Input

&f	pointer to the procedure containing the function to be integrated.
x1	2x1 vector, the limits of <i>x</i> . The first element is the upper limit and the second element is the lower limit.
tol	The tolerance to be used in testing for convergence.

Output

y	The estimated integral of f(x) between <i>x1</i> [1] and <i>x1</i> [2].
----------	--

Example

```
proc f(x);  
  ret p(sin(x));  
endp;  
  
let x1 = { 1, 0 };
```

```
y = intsimp(&f,x1,1e-8);  
print y;
```

The code above, returns the following:

```
0.45969769
```

This will integrate the function between 0 and 1.

Source

`intsimp.src`

See Also

[intquad1](#), [intquad2](#), [intquad3](#), [intgrat2](#), [intgrat3](#)

inv, invpd

Purpose

inv returns the inverse of an invertible matrix. **invpd** returns the inverse of a symmetric, positive definite matrix.

Format

```
y = inv(x);  
y = invpd(x);
```

Input

x	NxN matrix or K-dimensional array where the last two dimensions are NxN.
-----	--

inv, invpd

Output

y	$N \times N$ matrix or K -dimensional array where the last two dimensions are $N \times N$, containing the inverse of x .
-----	--

Remarks

x can be any legitimate expression that returns a matrix or array that is legal for the function.

If x is an array, the result will be an array containing the inverses of each 2-dimensional array described by the two trailing dimensions of x . In other words, for a $10 \times 4 \times 4$ array, the result will be an array of the same size containing the inverses of each of the 10 4×4 arrays contained in x

For **inv**, if x is a matrix, it must be square and invertible. Otherwise, if x is an array, the 2-dimensional arrays described by the last two dimensions of x must be square and invertible.

For **invpd**, if x is a matrix, it must be symmetric and positive definite. Otherwise, if x is an array, the 2-dimensional arrays described by the last two dimensions of x must be symmetric and positive definite.

If the input matrix is not invertible by these functions, they will either terminate the program with an error message or return an error code which can be tested for with the **scalerr** function. This depends on the **trap** state as follows:

If **trap** is set to 1, they will return a scalar errorcode:

inv	invpd
50	20

If **trap** is set to 0, they will terminate with an error message:

inv	invpd
"Matrix singular"	"Matrix not positive definite"

If the input to **invpd** is not symmetric, it is possible that the function will (erroneously) appear to operate successfully.

Positive definite matrices can be inverted by **inv**. However, for symmetric, positive definite matrices (such as moment matrices), **invpd** is about twice as fast as **inv**.

Example

```
n = 4000;
x1 = rndn(n,1);
x = ones(n,1)~x1;
btrue = { 1, 0.5 };
y = x*btrue + rndn(n,1);
bols = invpd(x'x)*x'y;
```

After the code above, *bols* will be equal to:

```
1.00237215
0.48249445
```

This example simulates some data and computes the **ols** coefficient estimator using the **invpd** function. First, the number of observations is specified. Second, a vector *x1* of standard Normal random variables is generated and is concatenated with a vector of **ones**(to create a constant term). The true coefficients are specified, and the dependent variable *y* is created. Then the **ols** coefficient estimates are computed.

When computing least-squares problems with poorly conditioned matrices, the slash operator "/" and the function **olsqr** will provide greater accuracy.

invswp

Purpose

Computes a generalized sweep inverse.

iscplx

Format

```
 $y = \text{invswp}(x);$ 
```

Input

x	NxN matrix.
-----	-------------

Output

y	NxN matrix, the generalized inverse of x .
-----	--

Remarks

This will invert any general matrix. That is, even matrices which will not invert using **inv** because they are singular will invert using **invswp**.

x and y will satisfy the two conditions:

1. $xyx = x$
2. $yxy = y$

invswp returns a row and column with zeros when the pivot fails. This is good for quadratic forms since it essentially removes rows with redundant information, i.e., the statistics generated will be "correct" but with reduced degrees of freedom.

The tolerance used to determine if a pivot element is zero is taken from the **crout** singularity tolerance. The corresponding row and column are zeroed out. See **SINGULARITY TOLERANCE**, CHAPTER 1.

iscplx

Purpose

Returns whether a matrix or N-dimensional array is complex or real.

Format

```
y = iscplx(x);
```

Input

x	$N \times K$ matrix or N-dimensional array.
-----	---

Output

y	scalar, 1 if x is complex, 0 if it is real.
-----	---

Example

```
x = { 1, 2i, 3 };  
if iscplx(x);  
    //code path for complex case  
else;  
    //code path for real case  
endif;
```

See Also

[hasimag](#), [iscplxf](#)

iscplxf

Purpose

Returns whether a data set is complex or real.

Format

```
y = iscplxf(fh);
```

isden

Input

fh	scalar, file handle of an open file.
------	--------------------------------------

Output

y	scalar, 1 if the data set is complex, 0 if it is real.
-----	--

See Also

[hasimag](#), [iscplx](#)

isden

Purpose

Returns whether a scalar, matrix or N-dimensional array contains denormals.

Format

```
 $y = \text{isden}(x);$ 
```

Input

x	NxK matrix or N-dimensional array.
-----	------------------------------------

Output

y	scalar, 1 if x contains a denormal, 0 if it does not.
-----	---

Example

Sometimes denormals can unnecessarily slow down calculations and it is best to flush

them to zero. This example tests whether the vector `x` contains any denormals and then sets any values between 0 and $1e-25$ to be equal to 0.

```
tol = 1e-25;

//Create a vector that contains a denormal
x = { 1, exp(-724.5), 3 };

if isden(x);
    //Get the index of all elements between 0 and tol
    idx = indexcat(x,0|tol);
    //Set all elements between 0 and tol equal to 0
    x[idx] = 0;
endif;
```

Before the `if` block in the code above, the second element of `x` is equal to approximately $3e-57$. After the `if` block this element is set equal to 0, the other elements of `x` are unchanged.

See Also

[denToZero](#)

isinfnanmiss

Purpose

Returns true if the argument contains an infinity, NaN, or missing value.

Format

```
y = isinfnanmiss(x);
```

ismiss

Input

x	NxK matrix.
-----	-------------

Output

y	scalar, 1 if x contains any infinities, NaNs, or missing values, else 0.
-----	--

See Also

[scalinfnanmiss](#), [ismiss](#), [scalmiss](#)

ismiss

Purpose

Returns a 1 if its matrix argument contains any missing values, otherwise returns a 0.
--

Format

$y = \text{ismiss}(x);$

Input

x	NxK matrix.
-----	-------------

Output

y	scalar, 1 if x contains any missing values, otherwise 0.
-----	--

Remarks

An element of x is considered to be a missing if and only if it contains a missing value in the real part. Thus, if $x = 1 + .i$, **ismiss**(x) will return a 0.

Example

```
x = { 1, 2, 3, 4 };

//Set the second element of 'x' to be a missing value
x[2] = miss(0,0);

print "before 'if' block, x = " x;

//If there are any missing values in 'x'
if ismiss(x);
    //Remove all rows with missing values from 'x'
    x = packr(x);
endif;

print "after 'if' block, x = " x;
```

```
before 'if' block, x =
1.0000000
.
3.0000000
4.0000000
after 'if' block, x =
1.0000000
3.0000000
4.0000000
```

To reset all missing values to a specified value, replace the call to **packr** above with a call to **missrv**.

itos

See Also

[scalmiss](#), [miss](#), [missrv](#)

itos

Purpose

Converts a scalar or matrix to the string representation of an integer.

Format

```
str = itos(x);
```

Input

<i>x</i>	scalar or NxK matrix.
----------	-----------------------

Output

<i>y</i>	string or string array containing the string representation of the elements of <i>x</i> .
----------	---

Example

Example 1

```
x = 4;  
str = itos(x);  
print "x = " x;  
print "str = " str;  
  
x =
```

```
1.000000  
  
str =  
1
```

Example 2

```
for i(1, 4, 1);  
    print "iteration "$+itos(i);  
endfor;
```

See Also

[ftos](#), [stof](#)

keep (dataloop)

k

keep (dataloop)

Purpose

Specifies columns (variables) to be saved to the output data set in a data loop.

Format

```
keep variable_list;
```

Remarks

Commas are optional in *variable_list*.

Retains only the specified variables in the output data set. Any variables referenced must already exist, either as elements of the source data set, or as the result of a previous [make](#), [vector](#), or [code](#) statement.

If neither [keep](#) nor [drop](#) is used, the output data set will contain all variables from the source data set, as well as any newly defined variables. The effects of multiple [keep](#) and [drop](#) statements are cumulative.

Example

```
keep age, pay, sex;
```

See Also

[drop \(dataloop\)](#)

key

Purpose

Returns the ASCII value of the next key available in the keyboard buffer.

Format

```
y = key;
```

Output

y	scalar, ASCII value of next available key in keyboard buffer.
---	---

Remarks

If you are working in terminal mode, **key** does not "see" any keystrokes until ENTER is pressed. The value returned will be zero if no key is available in the buffer or it will equal the ASCII value of the key if one is available. The key is taken from the buffer at this time and the next call to **key** will return the next key.

Example

```
format /rds 1,0;  
kk = 0;  
do until kk == 113;  
    kk = key;  
    if kk == 0;  
        continue;  
    elseif kk == vals(" ");  
        print "space \\" kk;  
    elseif kk >= vals("0") and kk <= vals("9");  
        print "digit \\" kk chrs(kk);
```

keyav

```
    else;
        print "\\\" kk;
    endif;
endo;
```

This is an example of a loop that processes keyboard input. This loop will continue until the `q` key (ASCII 113) is pressed.

See Also

[vals](#), [chrs](#), [upper](#), [lower](#), [con](#), [cons](#)

keyav

Purpose

Check if keystroke is available.

Format

```
x = keyav;
```

Output

x	scalar, value of key or 0 if no key is available.
---	---

See Also

[keyw](#), [key](#)

keyw

Purpose

Waits for and gets a key.

Format

```
k = keyw;
```

Output

<i>k</i>	scalar, ASCII value of the key pressed.
----------	---

Remarks

If you are working in terminal mode, **GAUSS** will not see any input until you press the ENTER key. **keyw** gets the next key from the keyboard buffer. If the keyboard buffer is empty, **keyw** waits for a keystroke. For normal keys, **keyw** returns the ASCII value of the key.

See Also

[key](#)

keyword

Purpose

Begins the definition of a keyword procedure. Keywords are user-defined functions with local or global variables.

Format

```
keyword name(str);
```

keyword

Input

<i>name</i>	literal, name of the keyword. This name will be a global symbol.
<i>str</i>	string, a name to be used inside the keyword to refer to the argument that is passed to the keyword when the keyword is called. This will always be local to the keyword, and cannot be accessed from outside the keyword or from other keywords or procedures.

Remarks

A keyword definition begins with the `keyword` statement and ends with the `endp` statement. See **PROCEDURES AND KEYWORDS**, CHAPTER 1.

Keywords always have 1 string argument and 0 returns. **GAUSS** will take everything past *name*, excluding leading spaces, and pass it as a string argument to the keyword. Inside the keyword, the argument is a local string. The user is responsible to manipulate or parse the string.

An example of a keyword definition is:

```
keyword add(str);
  local tok,sum;
  sum = 0;
  do until str $== "";
    { tok, str } = token(str);
    sum = sum + stof(tok);
  endo;
  print "Sum is: " sum;
endp;
```

To use this keyword, type:

```
add 1 2 3 4 5;
```

This keyword will respond by printing:

```
Sum is: 15
```

See Also

[proc](#), [local](#), [endp](#)

lag (dataloop)

lag (dataloop)

Purpose

Lags variables a specified number of periods.

Format

```
lag nv1 = var1:p1 [[nv2 = var2:p2...]];
```

Input

<i>var</i>	name of the variable to lag.
<i>p</i>	scalar constant, number of periods to lag.

Output

<i>nv</i>	name of the new lagged variable.
-----------	----------------------------------

Remarks

You can specify any number of variables to lag. Each variable can be lagged a different number of periods. Both positive and negative lags are allowed.

Lagging is executed before any other transformations. If the new variable name is different from that of the variable to lag, the new variable is first created and appended to a temporary data set. This temporary data set becomes the input data set for the data-loop, and is then automatically deleted.

lag1

Purpose

Lags a matrix by one time period for time series analysis.

Format

```
y = lag1(x);
```

Input

x	Nx1 column vector or NxK matrix.
---	----------------------------------

Output

y	NxK matrix, x lagged 1 period.
---	--------------------------------

Remarks

lag1 lags x by one time period, so the first observations of y are missing. **lag1** assumes that each column of the input is a different time series and that each row is an observation. Therefore if a 1xK row vector is passed to **lag1**, it will return a 1xK of missing values.

Example

```
y = { 1.2,  
      3.4,  
      2.5,  
      4.1,  
      2.8 };  
y_lag = lag1(y);
```

lag

```
print y_lag;
```

will return:

```
      .  
1.200000  
3.400000  
2.500000  
4.100000
```

Source

lag.src

See Also

[lag](#), [ismiss](#), [packr](#)

lag

Purpose

Lags a matrix a specified number of time periods for time series analysis.

Format

```
y = lag(x, t);
```

Input

x	NxK matrix.
t	scalar or Px1 vector, number of time periods.

Output

y NxK matrix, x lagged t periods.

Example

```
nlags = 2;
x = { 1.4, 2.7, 3.1, 2.9, 3.2, 2.5, 2.8 };
x_lag2 = lagn(x, nlags);
```

will assign `x_lag2` to equal:

```
      .
      .
    1.4
    2.7
    3.1
    2.9
    3.2
```

Creating multiple lags

If the number of time periods to lag is a Px1 column vector, then the output matrix will be an NxP matrix where each column contains one of the lags. For example, changing the `nlags` variable from the example above to be a 3x1 column vector like this:

```
nlags = { 1, 2, 3 };
x = { 1.4, 2.7, 3.1, 2.9, 3.2, 2.5, 2.8 };
lag_mat = lagn(x, nlags);
```

will assign `lag_mat` to equal:

```
      .      .      .
    1.4      .      .
    2.7     1.4      .
    3.1     2.7     1.4
```

lapeighb

2.9	3.1	2.7
3.2	2.9	3.1
2.5	3.2	2.9

Remarks

If t is positive, **lag** lags x back t time periods, so the first t observations of y are missing. If t is negative, **lag** lags x forward t time periods, so the last t observations of y are missing.

Source

lag.src

See Also

[lagl](#)

lapeighb

Purpose

Computes eigenvalues only of a real symmetric or complex Hermitian matrix selected by bounds.

Format

```
ve = lapeighb(x, vl, vu, abstol);
```

Input

x	NxN matrix, real symmetric or complex Hermitian.
vl	scalar, lower bound of the interval to be searched for eigenvalues.

vu scalar, upper bound of the interval to be searched for eigenvalues; *vu* must be greater than *vl*.

abstol scalar, the absolute error tolerance for the eigenvalues. An approximate eigenvalue is accepted as converged when it is determined to lie in an interval $[a, b]$ of width less than or equal to:

$$abstol + EPS * \max(|a|, |b|)$$

where EPS is machine precision. If *abstol* is less than or equal to zero, then $EPS * \|T\|$ will be used in its place, where *T* is the tridiagonal matrix obtained by reducing the input matrix to tridiagonal form.

Output

ve Mx1 vector, eigenvalues, where M is the number of eigenvalues on the half open interval $[vl, vu]$. If no eigenvalues are found then *ve* is a scalar missing value.

Remarks

lapeighb computes eigenvalues only which are found on the half open interval $[vl, vu]$. To find eigenvalues within a specified range of indices see **lapeighi**. For eigenvectors see **lapeighvi**, or **lapeighvb**. **lapeighb** is based on the LAPACK drivers DSYEVX and ZHEEVX. Further documentation of these functions may be found in the LAPACK User's Guide.

Example

```
x = { 5 2 1,
```

lapeighi

```
        2 6 2,  
        1 2 9 };  
  
vl = 5;  
vu = 10;  
ve = lapeighb(x,vl,vu,1e-15);  
print ve;
```

The code above returns:

```
6.0000
```

See Also

[lapeighvi](#), [lapeighvb](#)

lapeighi

Purpose

Computes eigenvalues only of a real symmetric or complex Hermitian matrix selected by index.

Format

```
ve = lapeighi(x, il, iu, abstol);
```

Input

<i>x</i>	NxN matrix, real symmetric or complex Hermitian.
<i>il</i>	scalar, index of the smallest desired eigenvalue ranking them from smallest to largest.
<i>iu</i>	scalar, index of the largest desired eigenvalue, <i>iu</i> must be greater than <i>il</i> .

abstol

scalar, the absolute error tolerance for the eigenvalues. An approximate eigenvalue is accepted as converged when it is determined to lie in an interval $[a, b]$ of width less than or equal to $abstol + EPS * \max(|a|, |b|)$, where EPS is machine precision. If *abstol* is less than or equal to zero, then $EPS * \|T\|$ will be used in its place, where T is the tridiagonal matrix obtained by reducing the input matrix to tridiagonal form.

Output

ve

$(iu-il+1) \times 1$ vector, eigenvalues.

Remarks

lapeighi computes $iu-il+1$ eigenvalues only given a range of indices, i.e., the i th to j th eigenvalues, ranking them from smallest to largest. To find eigenvalues within a specified range see **lapeighxb**. For eigenvectors see **lapeighvi**, or **lapeighvb**. **lapeighi** is based on the LAPACK drivers DSYEVX and ZHEEVX. Further documentation of these functions may be found in the LAPACK User's Guide.

Example

```
x = { 5 2 1,
      2 6 2,
      1 2 9 };

il = 2;
iu = 3;
ve = lapeighi(x, il, iu, 0);
print ve;
```

lapeighvb

The code above calculates the second and third eigenvalues and returns:

```
6.0000
10.6056
```

To calculate the first, second and third eigenvalues, reusing the same `x` from above:

```
ve = lapeighi(x, 1, 3, 0);
print ve;
```

The output from this code is:

```
3.3944
6.0000
10.6056
```

See Also

[lapeighb](#), [lapeighvi](#), [lapeighvb](#)

lapeighvb

Purpose

Computes eigenvalues and eigenvectors of a real symmetric or complex Hermitian matrix selected by bounds.

Format

```
{ ve, va } = lapeighvb(x, vl, vu, abstol);
```

Input

<code>x</code>	NxN matrix, real symmetric or complex Hermitian.
<code>vl</code>	scalar, lower bound of the interval to be searched for

	eigenvalues.
<i>vu</i>	scalar, upper bound of the interval to be searched for eigenvalues; <i>vu</i> must be greater than <i>vl</i> .
<i>abstol</i>	scalar, the absolute error tolerance for the eigenvalues. An approximate eigenvalue is accepted as converged when it is determined to lie in an interval $[a, b]$ of width less than or equal to $abstol + EPS * \max(a , b)$, where EPS is machine precision. If <i>abstol</i> is less than or equal to zero, then $EPS * T $ will be used in its place, where <i>T</i> is the tridiagonal matrix obtained by reducing the input matrix to tridiagonal form.

Output

<i>ve</i>	Mx1 vector, eigenvalues, where M is the number of eigenvalues on the half open interval $[vl, vu]$. If no eigenvalues are found then s is a scalar missing value.
<i>va</i>	NxM matrix, eigenvectors.

Remarks

laeighvb computes eigenvalues and eigenvectors which are found on the half open interval $[vl, vu]$. **laeighvb** is based on the LAPACK drivers DSYEVX and ZHEEVX. Further documentation of these functions may be found in the LAPACK User's Guide.

Example

```
x = { 5 2 1,
      2 6 2,
      1 2 9 };
```

lapeighvi

```
v1 = 5;  
vu = 10;  
{ ve, va } = lapeighvb(x,v1,vu,0);  
  
print "Eigenvalues" ve;  
print "Eigenvectors = " va;
```

```
Eigenvalues =    6.0000  
Eigenvectors =  
-0.5774  
-0.5774  
 0.5774
```

If you increase the value of *vu* to 12.

```
{ ve, va } = lapeighvb(x,5,12,0);  
  
print "Eigenvalues" ve;  
print "Eigenvectors = " va;
```

```
Eigenvalues  
 6.0000  
10.6056  
Eigenvectors =  
-0.5774  0.3197  
-0.5774  0.4908  
 0.5774  0.8105
```

lapeighvi

Purpose

Computes selected eigenvalues and eigenvectors of a real symmetric or complex Hermitian matrix.

Format

```
{ ve, va } = lapeighvi(x, il, iu, abstol);
```

Input

<i>x</i>	NxN matrix, real symmetric or complex Hermitian.
<i>il</i>	scalar, index of the smallest desired eigenvalue ranking them from smallest to largest.
<i>iu</i>	scalar, index of the largest desired eigenvalue, <i>iu</i> must be greater than <i>il</i> .
<i>abstol</i>	scalar, the absolute error tolerance for the eigenvalues. An approximate eigenvalue is accepted as converged when it is determined to lie in an interval [<i>a</i> , <i>b</i>] of width less than or equal to <i>abstol</i> + EPS*max(<i>a</i> , <i>b</i>), where EPS is machine precision. If <i>abstol</i> is less than or equal to zero, then EPS* <i>T</i> will be used in its place, where <i>T</i> is the tridiagonal matrix obtained by reducing the input matrix to tridiagonal form.

Output

<i>ve</i>	(<i>iu</i> - <i>il</i> + 1)x1 vector, eigenvalues.
<i>va</i>	Nx(<i>iu</i> - <i>il</i> + 1) matrix, eigenvectors.

Remarks

lapeighvi computes *iu*-*il*+1 eigenvalues and eigenvectors given a range of indices, i.e., the *i*th to *j*th eigenvalues, ranking them from smallest to largest. To find eigenvalues and eigenvectors within a specified range see **lapeighvb**.
lapeighvi is based on the LAPACK drivers DSYEVX and ZHEEVX. Further

lapgeig

documentation of these functions may be found in the LAPACK User's Guide.

Example

```
x = { 5 2 1,
      2 6 2,
      1 2 9 };

il = 2;
iu = 3;
{ ve,va } = lapeighvi(x,il,iu,0);
print "ve = " ve;
print "va = " va;
```

```
ve =
6.0000
10.6056

va =
-0.5774    0.3197
-0.5774    0.4908
 0.5774    0.8105
```

See Also

[lapeighvb](#), [lapeighb](#)

lapgeig

Purpose

Computes generalized eigenvalues for a pair of real or complex general matrices.

Format

```
{ va1, va2 } = lapgeig(A, B);
```

Input

<i>A</i>	NxN matrix, real or complex general matrix.
<i>B</i>	NxN matrix, real or complex general matrix.

Output

<i>va1</i>	Nx1 vector, numerator of eigenvalues.
<i>va2</i>	Nx1 vector, denominator of eigenvalues.

Remarks

va1 and *va2* are the vectors of the numerators and denominators respectively of the eigenvalues of the solution of the generalized symmetric eigenproblem of the form $Aw = eBw$ where A and B are real or complex general matrices and $w = va1 ./ va2$. The generalized eigenvalues are not computed directly because some elements of *va2* may be zero, i.e., the eigenvalues may be infinite. This procedure calls the LAPACK routines DGGEV and ZGGEV.

See Also

[lapgeig](#), [lapgeigh](#)

lapgeigh

Purpose

Computes generalized eigenvalues for a pair of real symmetric or Hermitian matrices.

lapgeigh

Format

```
ve = lapgeigh(A, B);
```

Input

A	NxN matrix, real or complex symmetric or Hermitian matrix.
B	NxN matrix, real or complex positive definite symmetric or Hermitian matrix.

Output

ve	Nx1 vector, eigenvalues.
------	--------------------------

Remarks

ve is the vector of eigenvalues of the solution of the generalized symmetric eigenproblem of the form $Ax = \lambda Bx$.

Example

```
A = { 3 4 5,  
      2 5 2,  
      3 2 4 };  
  
B = { 4 2 2,  
      2 6 1,  
      2 1 8 };  
  
ve = lapgeigh(A,B);  
print ve;
```

The code above returns:

```

0.1219
0.6787
0.9494

```

This procedure calls the LAPACK routines DSYGV and ZHEGV.

See Also

[lapgeig](#), [lapgeighv](#)

lapgeighv

Purpose

Computes generalized eigenvalues and eigenvectors for a pair of real symmetric or Hermitian matrices.

Format

$$\{ ve, va \} = \text{lapgeighv}(A, B);$$

Input

A	NxN matrix, real or complex symmetric or Hermitian matrix.
B	NxN matrix, real or complex positive definite symmetric or Hermitian matrix.

Output

ve	Nx1 vector, eigenvalues.
va	NxN matrix, eigenvectors.

lapgeighv

Remarks

ve and va are the eigenvalues and eigenvectors of the solution of the generalized symmetric eigenproblem of the form $Ax = \lambda B$. Equivalently, va diagonalizes $U'^{-1} A^* U^{-1}$ in the following way

$$va^* U'^{-1} A^* U^{-1} va = ve$$

where $B = U'U$. This procedure calls the LAPACK routines DSYGV and ZHEGV.

Example

```
A = { 3 4 5,
      2 5 2,
      3 2 4 };

B = { 4 2 2,
      2 6 1,
      2 1 8 };

{ ve, va } = lapgeighv(A,B);
```

```
print ve;
```

```
-0.0425
0.5082
0.8694
```

```
print va;
```

```
0.3575 -0.0996 0.9286
-0.2594 0.9446 0.2012
-0.8972 -0.3128 0.3118
```

See Also

[lapgeig](#), [lapgeigh](#)

lapgeigv

Purpose

Computes generalized eigenvalues, left eigenvectors, and right eigenvectors for a pair of real or complex general matrices.

Format

$$\{ va1, va2, lve, rve \} = \mathbf{lapgeigv}(A, B);$$

Input

A	NxN matrix, real or complex general matrix.
B	NxN matrix, real or complex general matrix.

Output

$va1$	Nx1 vector, numerator of eigenvalues.
$va2$	Nx1 vector, denominator of eigenvalues.
lve	NxN left eigenvectors.
rve	NxN right eigenvectors.

Remarks

$va1$ and $va2$ are the vectors of the numerators and denominators respectively of the eigenvalues of the solution of the generalized symmetric eigenproblem of the form $Aw = \lambda Bw$ where A and B are real or complex general matrices and $w = va1 ./ va2$. The generalized eigenvalues are not computed directly because some elements of $va2$ may be zero, i.e., the eigenvalues may be infinite.

The left and right eigenvectors diagonalize $U'^{-1} * A * U^{-1}$ where $B = U' * U$, that is,

lapgsvdcst

$$\text{lve}'U'^{-1}A*U*\text{lve}' = w$$

and

$$\text{rve}'U'^{-1}*A*U^{-1}*\text{rve} = w$$

This procedure calls the LAPACK routines DGGEV and ZGGEV.

See Also

[lapgeig](#), [lapgeigh](#)

lapgsvdcst

Purpose

Compute the generalized singular value decomposition of a pair of real or complex general matrices.

Format

$$\{ C, S, R, U, V, Q \} = \text{lapgsvdcst}(A, B);$$

Input

<i>A</i>	MxN matrix.
<i>B</i>	PxN matrix.

Output

<i>C</i>	Lx1 vector, singular values for <i>A</i> .
<i>S</i>	Lx1 vector, singular values for <i>B</i> .
<i>R</i>	(K+L)x(K+L) upper triangular matrix.
<i>U</i>	MxM matrix, orthogonal transformation matrix.

V	$P \times P$ matrix, orthogonal transformation matrix.
Q	$N \times N$ matrix, orthogonal transformation matrix.

Remarks

(1) The generalized singular value decomposition of A and B is

$$U' * A * Q = D_1 * Z$$

$$V' * B * Q = D_2 * Z$$

where U , V , and Q are orthogonal matrices (see **lapgsvdcst** and **lapgsvdst**). Letting $K + L = \text{the rank of } A|B$ then R is a $(K+L) \times (K+L)$ upper triangular matrix, D_1 and D_2 are $M \times (K+L)$ and $P \times (K+L)$ matrices with entries on the diagonal, $Z = [O \ R]$, and if $M-K-L \geq 0$

$$D_1 = \begin{array}{c} \begin{array}{cc} & K \ L \\ K & [\ I \ 0 \] \\ L & [\ 0 \ C \] \\ M - K - L & [\ 0 \ 0 \] \end{array} \end{array}$$

$$D_2 = \begin{array}{c} \begin{array}{cc} & K \ L \\ P & [\ 0 \ S \] \\ P - L & [\ 0 \ 0 \] \end{array} \end{array}$$

$$[\ 0 \ R \] = \begin{array}{c} \begin{array}{ccc} & N-K-L & K \ L \\ K & [\ 0 & R_{11} \ R_{12} \] \\ L & [\ 0 & 0 \ R_{22} \] \end{array} \end{array}$$

or if $M-K-L < 0$

$$D_1 = \begin{array}{c} \begin{array}{ccc} & K & M-K & K+L-M \\ K & [\ I & 0 & 0 \] \\ M-K & [\ 0 & 0 & 0 \] \end{array} \end{array}$$

lapgsvdcst

$$\begin{matrix} & & & N-K-L & K & M-K & K+L-M \\ & & K & [& 0 & R11 & R12 & R13 &] \\ [& 0 & R &] = & M-K & [& 0 & 0 & R22 & R23 &] \\ & & K+L-M & [& 0 & 0 & 0 & R33 &] \end{matrix}$$

(2) Form the matrix

$$X = Q \begin{bmatrix} I & 0 \\ 0 & R^{-1} \end{bmatrix}$$

then

$$A = U'^{-1}E_1X$$

$$B = V'^{-1}E_2X^{-1}$$

where

$$E1 = \begin{bmatrix} 0 & D1 \end{bmatrix}$$

$$E2 = \begin{bmatrix} 0 & D2 \end{bmatrix}$$

(3) The generalized singular value decomposition of A and B implicitly produces the singular value decomposition of AB^{-1} :

$$AB^{-1} = UD_1D_2^{-1}V'$$

This procedure calls the LAPACK routines DGGSVD and ZGGSVD.

See Also

[lapgsvds](#), [lapgsvdst](#)

lapgsvds

Purpose

Compute the generalized singular value decomposition of a pair of real or complex general matrices.

Format

$$\{ C, S, R \} = \mathbf{lapgsvds}(A, B);$$

Input

A	$M \times N$ real or complex matrix.
B	$P \times N$ real or complex matrix.

Output

C	$L \times 1$ vector, singular values for A .
S	$L \times 1$ vector, singular values for B .
R	$(K+L) \times (K+L)$ upper triangular matrix.

Remarks

(1) The generalized singular value decomposition of A and B is

$$U^* A Q = D_1 Z$$

$$V^* B Q = D_2 Z$$

where U , V , and Q are orthogonal matrices (see **lapgsvdcst** and **lapgsvdst**). Letting $K+L = \text{rank of } A|B$ then R is a $(K+L) \times (K+L)$ upper triangular matrix, D_1

lapgsvds

and D_2 are $M \times (K+L)$ and $P \times (K+L)$ matrices with entries on the diagonal, $Z = [OR]$, and if $M-K-L \geq 0$

$$D1 = \begin{array}{cc} & \begin{array}{cc} K & L \end{array} \\ \begin{array}{c} K \\ L \\ M - K - L \end{array} & \begin{bmatrix} I & 0 \\ 0 & C \\ 0 & 0 \end{bmatrix} \end{array}$$

$$D2 = \begin{array}{cc} & \begin{array}{cc} K & L \end{array} \\ \begin{array}{c} P \\ P - L \end{array} & \begin{bmatrix} 0 & S \\ 0 & 0 \end{bmatrix} \end{array}$$

$$\begin{bmatrix} 0 & R \end{bmatrix} = \begin{array}{cc} & \begin{array}{cc} N-K-L & \begin{array}{cc} K & L \end{array} \end{array} \\ \begin{array}{c} K \\ L \end{array} & \begin{bmatrix} 0 & R11 & R12 \\ 0 & 0 & R22 \end{bmatrix} \end{array}$$

or if $M-K-L < 0$

$$D1 = \begin{array}{cc} & \begin{array}{ccc} K & M-K & K+L-M \end{array} \\ \begin{array}{c} K \\ M-K \end{array} & \begin{bmatrix} I & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix} \end{array}$$

$$\begin{bmatrix} 0 & R \end{bmatrix} = \begin{array}{cc} & \begin{array}{cccc} N-K-L & K & M-K & K+L-M \end{array} \\ \begin{array}{c} K \\ M-K \\ K+L-M \end{array} & \begin{bmatrix} 0 & R11 & R12 & R13 \\ 0 & 0 & R22 & R23 \\ 0 & 0 & 0 & R33 \end{bmatrix} \end{array}$$

(2) Form the matrix

$$X = Q \begin{bmatrix} I & 0 \\ 0 & R^{-1} \end{bmatrix}$$

then

$$A = U'^{-1}E_1X$$

$$B = V'^{-1}E_2X^{-1}$$

where

$$E1 = \begin{bmatrix} 0 & D1 \end{bmatrix}$$

$$E2 = \begin{bmatrix} 0 & D2 \end{bmatrix}$$

(3) The generalized singular value decomposition of A and B implicitly produces the singular value decomposition of AB^{-1} :

$$AB^{-1} = UD_1D_2^{-1}V'$$

This procedure calls the LAPACK routines DGGSD and ZGGSD.

See Also

[lapgsvdct](#), [lapgsvdst](#)

lapgsvdst

Purpose

Compute the generalized singular value decomposition of a pair of real or complex general matrices.

Format

$$\{ D1, D2, Z, U, V, Q \} = \mathbf{lapgsvdst}(A, B);$$

Input

A MxN matrix.

lapgsvdst

B	$P \times N$ matrix.
-----	----------------------

Output

$D1$	$M \times (K+L)$ matrix, with singular values for A on diagonal.
$D2$	$P \times (K+L)$ matrix, with singular values for B on diagonal.
Z	$(K+L) \times N$ matrix, partitioned matrix composed of a zero matrix and upper triangular matrix.
U	$M \times M$ matrix, orthogonal transformation matrix.
V	$P \times P$ matrix, orthogonal transformation matrix.
Q	$N \times N$ matrix, orthogonal transformation matrix.

Remarks

(1) The generalized singular value decomposition of A and B is

$$U' A Q = D_1 Z$$

$$V' B Q = D_2 Z$$

where U , V , and Q are orthogonal matrices (see **lapgsvdcst** and **lapgsvdst**). Letting $K+L$ = the rank of $A|B$ then R is a $(K+L) \times (K+L)$ upper triangular matrix, $D1$ and $D2$ are $M \times (K+L)$ and $P \times (K+L)$ matrices with entries on the diagonal, $Z = [0R]$, and if $M-K-L \geq 0$

$$D1 = \begin{matrix} & & K & L \\ & & [& I & 0 &] \\ & L & [& 0 & C &] \\ M - K - L & [& 0 & 0 &] \end{matrix}$$

$$D2 = \begin{array}{cc} & K \quad L \\ P & [\quad 0 \quad S \quad] \\ P - L & [\quad 0 \quad 0 \quad] \end{array}$$

$$\begin{array}{cc} & N-K-L & K & L \\ [\quad 0 \quad R \quad] = K & [\quad 0 & R11 & R12 \quad] \\ & L & [\quad 0 & 0 & R22 \quad] \end{array}$$

or if $M-K-L < 0$

$$D1 = \begin{array}{ccc} & K & M-K & K+L-M \\ K & [\quad I & 0 & 0 \quad] \\ M-K & [\quad 0 & 0 & 0 \quad] \end{array}$$

$$\begin{array}{cc} & N-K-L & K & M-K & K+L-M \\ & K & [\quad 0 & R11 & R12 & R13 \quad] \\ [\quad 0 \quad R \quad] = & M-K & [\quad 0 & 0 & R22 & R23 \quad] \\ & K+L-M & [\quad 0 & 0 & 0 & R33 \quad] \end{array}$$

(2) Form the matrix

$$X = Q \begin{bmatrix} I & 0 \\ 0 & R^{-1} \end{bmatrix}$$

then

$$A = U'^{-1}E_1X$$

$$B = V'^{-1}E_2X^{-1}$$

where

$$E1 = [\quad 0 \quad D1 \quad]$$

$$E2 = [\quad 0 \quad D2 \quad]$$

lapgschur

(3) The generalized singular value decomposition of A and B implicitly produces the singular value decomposition of AB^{-1} :

$$AB^{-1} = UD_1D_2^{-1}V^*$$

This procedure calls the LAPACK routines DGGSD and ZGGSD.

See Also

[lapgsvds](#), [lapgsvdst](#)

lapgschur

Purpose

Compute the generalized Schur form of a pair of real or complex general matrices.

Format

```
{ sa, sb, q, z } = lapgschur(A, B);  
{ sa, sb, q, z } = lapgschur(A, B, sort_type);
```

Input

A	NxN matrix, real or complex general matrix.
B	NxN matrix, real or complex general matrix.

sort_type

Optional input, scalar or string specifying how to sort the eigenvalues. Options include:

- | | | |
|---|-------|---|
| 1 | "udi" | Absolute value of the eigenvalue less than 1.0. (Unit disk inside). |
| 2 | "udo" | Absolute value of the eigenvalue greater than or equal to 1.0. (Unit disk outside). |
| 3 | "lhp" | Value of the real portion of the eigenvalue less than 0. (Left hand plane). |
| 4 | "rhp" | Value of the real portion of the eigenvalue greater than 0. (Right hand plane). |
| 5 | "ref" | Real eigenvalues first. (Complex portion less than <i>imagtol</i>). |
| 6 | "cef" | Complex eigenvalues first. (Complex portion greater than <i>imagtol</i>). |

Output

<i>sa</i>	NxN matrix, Schur form of A , sometimes called S .
<i>sb</i>	NxN matrix, Schur form of B , sometimes called T .
<i>q</i>	NxN matrix, left Schur vectors.
<i>z</i>	NxN matrix, right Schur vectors.

Examples

Example 1: Basic usage

```
//For repeatable random numbers
rndseed 23434;

//Matrix dimensions
order = 4;

//Create 2 square, complex matrices
A = complex(rndn(order, order), rndn(order,order));
B = complex(rndn(order, order), rndn(order,order));

//Perform 'QZ' decomposition
{ sa, sb, q, z } = lapgschur(A,B);

//Calculate generalized eigenvalues
eig_vals = diag(sa) ./ diag(sb);

print "Generalized eigenvalues = ";
print eig_vals;

print "Absolute value of the generalized eigenvalues = ";
print abs(eig_vals);
```

The above code should return the following output:

```
Generalized eigenvalues =

-0.76631163 -      1.3445924i
 0.65409426 -      0.18908938i
-0.012440975 +      0.47626474i
-0.75927986 +      1.6212326i

Absolute value of the generalized eigenvalues =
```

```

1.5476312
0.68087745
0.47642721
1.7902237

```

Example 2: Ordering eigenvalues

You can order the eigenvalues, by passing in the optional third input, *sort_type*. The code below uses the same *A* and *B* variables made in the example above.

```

//Perform 'QZ' decomposition and
//reorder generalized eigenvalues, placing
//those with absolute value less than 1
//on the upper left
{ sa, sb, q, z } = lapgschur(A, B, "udi");

//Calculate generalized eigenvalues
eig_vals = diag(sa) ./ diag(sb);

print "Generalized eigenvalues = ";
print (eig_vals);

print "Absolute value of the generalized eigenvalues = ";
print abs(eig_vals);

```

The code above should print out the sorted eigenvalues as we see below.

```

Generalized eigenvalues =

    0.65409426 -    0.18908938i
   -0.012440975 +    0.47626474i
   -0.76631163 -    1.3445924i
   -0.75927986 +    1.6212326i

```

lapgschur

Absolute value of the generalized eigenvalues =

0.68087745

0.47642721

1.5476312

1.7902237

Remarks

- The pair of matrices sa (sometimes called S) and sb (sometimes called T) are in generalized real Schur form if:
 - sb is upper triangular with non-negative diagonal.
 - sa is block upper triangular with 1x1 and 2x2 blocks. The 1x1 blocks correspond to real generalized eigenvalues and the 2x2 blocks to pairs of complex conjugate eigenvalues.
- The real generalized eigenvalues can be computed by dividing the diagonal element of sa by the corresponding diagonal element of sb .
- The complex generalized eigenvalues are computed by first constructing two complex conjugate numbers from 2x2 block where the real parts are on the diagonal of the block and the imaginary part on the off-diagonal. The eigenvalues are then computed by dividing the two complex conjugate values by their corresponding diagonal elements of sb .
- The generalized Schur vectors q and z are orthogonal matrices ($q'q = I$ and $z'z = I$) that reduce A and B to Schur form:

$$sa = q'A*z$$

$$sb = q'B*z$$

$$A = q*sa*z'$$

$$B = q*sb*z'$$

This procedure calls the LAPACK routine DGGES if both input matrices are real. If either input A or B are complex, even if the complex portion is zero, ZGGES is called.

- If only the generalized eigenvalues are needed, you can call **lapgeig**, or **lapgeigv**.

lapsevdcusv

Purpose

Computes the singular value decomposition of a real or complex rectangular matrix, returns compact U and v .

Format

$$\{ u, s, v \} = \text{lapsevdcusv}(x);$$

Input

x	MxN matrix, real or complex rectangular matrix.
-----	---

Output

u	Mxmin(M,N) matrix, left singular vectors.
s	min(M,N)xN matrix, singular values.
v	NxN matrix, right singular values.

Remarks

lapsevdcusv computes the singular value decomposition of a real or complex rectangular matrix. The SVD is

$$x = usv'$$

lapsvdcusv

where v is the matrix of right singular vectors. **lapsvdcusv** is based on the LAPACK drivers DGESVD and ZGESVD. Further documentation of these functions may be found in the LAPACK User's Guide.

Example

```
x = { 2.143 4.345 6.124,  
      1.244 5.124 3.412,  
      0.235 5.657 8.214 };
```

```
{ u,s,v } = lapsvdcusv(x);
```

```
print u;
```

```
-0.55531277  0.049048431  0.83019394  
-0.43090168  0.83684123  -0.33766923  
-0.71130266 -0.54524400  -0.44357356
```

```
print s;
```

```
13.895868  0.0000000  0.0000000  
0.0000000  2.1893939  0.0000000  
0.0000000  0.0000000  1.4344261
```

```
print v;
```

```
-0.13624432  -0.62209955  -0.77099263  
0.46497296   0.64704876  -0.60425826  
0.87477862  -0.44081748   0.20110275
```

See Also

[lapsvds](#), [lapsvdusv](#)

lapsvds

Purpose

Computes the singular values of a real or complex rectangular matrix

Format

$$s = \text{lapsvds}(x);$$

Input

x MxN matrix, real or complex rectangular matrix.

Output

s min(M,N)x1 vector, singular values.

Remarks

lapsvds computes the singular values of a real or complex rectangular matrix. The SVD is

$$x = uv^H$$

where v is the matrix of right singular vectors. For the computation of the singular vectors, see **lapsvdcusv** and **lapsvdusv**.

lapsvds is based on the LAPACK drivers DGESVD and ZGESVD. Further documentation of these functions may be found in the LAPACK User's Guide.

lapsvdusv

Example

```
x = { 2.143 4.345 6.124,  
      1.244 5.124 3.412,  
      0.235 5.657 8.214 };
```

```
va = lapsvds(x);  
print va';
```

```
13.895868 2.1893939 1.4344261
```

```
xi = { 4+1 3+1 2+2,  
       1+2 5+3 2+2,  
       1+1 2+1 6+2 };
```

```
ve = lapsvds(xi);  
print ve';
```

```
10.352877 4.0190557 2.3801546
```

Note the transpose operator (') at the end of the `print` statements. This causes the output of these column vectors to be printed as a row vector.

See Also

[lapsvdcusv](#), [lapsvdusv](#)

lapsvdusv

Purpose

Computes the singular value decomposition a real or complex rectangular matrix.

Format

```
{ u, s, v } = lapsvdusv(x);
```

Input

x	MxN matrix, real or complex rectangular matrix.
-----	---

Output

u	MxM matrix, left singular vectors.
s	MxN matrix, singular values.
v	NxN matrix, right singular values.

Remarks

lapsvdusv computes the singular value decomposition of a real or complex rectangular matrix. The SVD is

$$x = usv'$$

where v is the matrix of right singular vectors. **lapsvdusv** is based on the LAPACK drivers DGESVD and ZGESVD. Further documentation of these functions may be found in the LAPACK User's Guide.

Example

```
x = { 2.143 4.345 6.124,
      1.244 5.124 3.412,
      0.235 5.657 8.214 };

{ u,s,v } = lapsvdusv(x);

print u;
```

ldl

```
-0.5553  0.0490  0.8302
-0.4309  0.8368 -0.3377
-0.7113 -0.5452 -0.4436
```

```
print s;
```

```
13.8959 0.0000 0.0000
0.0000 2.1894 0.0000
0.0000 0.0000 1.4344
```

```
print v;
```

```
-0.1362  0.4650  0.8748
 0.6221  0.6470 -0.4408
-0.7710 -0.6043  0.2011
```

See Also

[lapsvds](#), [lapsvdcusv](#)

ldl

Purpose

Returns the L and D factors of the LDL' factorization of a real symmetric matrix.

Format

$$\{L, D\} = \mathbf{ldl}(A);$$

Input

A	$N \times N$ real symmetric matrix.
-----	-------------------------------------

Output

L	$N \times N$ scrambled, lower triangular matrix, containing the factor L .
D	$N \times N$ block diagonal matrix, containing the factor D .

Remarks

Matrix factorization is the most computationally intense part of solving a system of linear equations. The factorization can be saved and reused multiple times to prevent the need to repeat the matrix factorization step. If you only need the LDL' factorization for this purpose, the combination of **ldlp** and **ldlsol** may be a better choice.

Example

```
A = { 5   9   3   4,
      9  -6   8   1,
      3   8   2   3,
      4   1   3   9 };
```

```
//Factorize matrix 'A'
{L, D} = ldl(A);
```

```
A_new = L * D * L';
```

After the code above:

```
L =
-1.50    1.00    0.00    0.00
 1.00    0.00    0.00    0.00
```

ldlp

```
-1.33    0.81    1.00    0.00
-0.17    0.30   -0.25    1.00
```

D =

```
-6.00    0.00    0.00    0.00
 0.00   18.50    0.00    0.00
 0.00    0.00    0.50    0.00
 0.00    0.00    0.00    7.50
```

A_new =

```
5.00    9.00    3.00    4.00
9.00   -6.00    8.00    1.00
3.00    8.00    2.00    3.00
4.00    1.00    3.00    9.00
```

See Also

[ldlp](#), [ldlsol](#), [chol](#), [solpd](#)

ldlp

Purpose

Returns the Bunch-Kaufmann factorization of a real symmetric matrix along with a permutation vector.

Format

```
ldl_factor = ldlp(A);
```

Input

A NxN real symmetric matrix.

Output

`ldl_factor` (N+1)xN matrix, containing the factors L and D as well as the permutation vector P, which can be passed directly to **ldlsol** to solve a system of linear equations.

Remarks

Matrix factorization is the most computationally intense part of solving a system of linear equations. The factorization can be saved and reused multiple times to prevent the need to repeat the matrix factorization step. **ldlp** uses the LAPACK function **dsytrf** to compute the factorization.

Example

```
A = { 5    9    3    4,
      9   -6    8    1,
      3    8    2    3,
      4    1    3    9 };
b = { 1.4, 4, 0.5, 3 };

//Factorize matrix 'A'
ldl_f = ldlp(A);

//Solve system of equations
x = ldlsol(b, ldl_f);
```

The above code will solve the system of linear equations $Ax = b$, assigning x to be equal to:

```
0.5729
x = -0.1529
    -0.2829
    0.1900
```

ldlsol

See Also

[ldlsol](#), [chol](#), [solpd](#)

ldlsol

Purpose

Computes the solution to a system of linear equations given a factorized matrix returned by the function **ldlp** and one or more right hand sides.

Format

```
x = ldlsol(b, ldl_factor);
```

Input

<i>b</i>	Nx1 vector or NxK matrix, the right hand sides of the system of linear equations.
<i>ldl_factor</i>	Nx(N+1) matrix, containing the a factorization returned from the GAUSS function ldlp .

Output

<i>x</i>	Nx1 vector or NxK matrix, containing the solution to $LDL^T x = b$.
----------	--

Remarks

Matrix factorization is the most computationally intense part of solving a system of linear equations. The factorization can be saved and reused multiple times to prevent the need to repeat the matrix factorization step. **ldlsol** uses the LAPACK function **dsytrs** to solve the system of linear equations.

Example

```
A = { 5    9    3    4,
      9   -6    8    1,
      3    8    2    3,
      4    1    3    9 };
b = { 1.4, 4, 0.5, 3 };

//Factorize matrix 'A'
ldl_f = ldlp(A);

//Solve system of equations
x = ldlsol(b, ldl_f);
```

The above code will solve the system of linear equations $Ax = b$, assigning x to be equal to:

```
      0.5729
x = -0.1529
     -0.2829
      0.1900
```

See Also

[ldlp](#), [lusol](#), [solpd](#)

let

Purpose

Creates a matrix from a list of numeric or character values. The result is always of type matrix, string, or string array.

Format

```
let x = constant_list;
```

let

Remarks

Expressions and variable names are not allowed in the `let` command, expressions such as this:

```
let x[2,1] = 3*a b
```

are illegal. To define matrices by combining matrices and expressions, use an expression containing the concatenation operators: `~` and `|`.

Numbers can be entered in scientific notation. The syntax is $dE\pm n$, where d is a number and n is an integer (denoting the power of 10):

```
let x = 1e+10 1.1e-4 4.019e+2;
```

Complex numbers can be entered by joining the real and imaginary parts with a sign (+ or -); there should be no spaces between the numbers and the sign. Numbers with no real part can be entered by appending an "i" to the number:

```
let x = 1.2+23 8.56i 3-2.1i -4.2e+6i 1.2e-4-4.5e+3i;
```

If curly braces are used, the `let` is optional.

```
let x = { 1 2 3, 4 5 6, 7 8 9 };
```

```
x = { 1 2 3, 4 5 6, 7 8 9 };
```

If indices are given, a matrix of that size will be created:

```
let x[2,2] = 1 2 3 4;
```

```
x = 1 2  
    3 4
```

If indices are not given, a column vector will be created:


```
let x = 1 2 3 4;
```

```

      1
x =   2
      3
      4

```

You can create matrices with no elements, i.e., "empty matrices" . Just use a set of empty curly braces:

```
x = {};
```

Empty matrices are chiefly used as the starting point for building up a matrix, for example in a `do` loop. See **Matrices**, Section 1.0.1, for more information on empty matrices.

Character elements are allowed in a `let` statement:

```
let x = age pay sex;
```

```

      AGE
x =   PAY
      SEX

```

Lowercase elements can be created if quotation marks are used. Note that each element must be quoted.

```
let x = "age" "pay" "sex";
```

```

      age
x =   pay
      sex

```

Example

```
let x;
```

let

assigns x to be:

```
x = 0
```

```
let x = { 1 2 3, 4 5 6, 7 8 9 };
```

assigns x to be:

```
  1 2 3  
x = 3 4 5  
  6 7 8
```

```
let x[3,3] = 1 2 3 4 5 6 7 8 9;
```

assigns x to be:

```
  1 2 3  
x = 3 4 5  
  6 7 8
```

```
let x[3,3] = 1;
```

assigns x to be:

```
  1 1 1  
x = 1 1 1  
  1 1 1
```

```
let x[3,3];
```

assigns x to be:

```
  0 0 0  
x = 0 0 0  
  0 0 0
```

```
let x = dog cat;
```

assigns `x` to be:

```
x = DOG  
    CAT
```

```
let x = "dog" "cat";
```

assigns `x` to be:

```
x = dog  
    cat
```

```
let string x = { "Median Income", "Country" };
```

assigns `x` to be:

```
x = Median Income  
    Country
```

See Also

[con](#), [cons](#), [declare](#), [load](#)

lib

Purpose

Builds and updates library files.

Format

```
lib library file;  
lib library -flag;  
lib library file -flag1 -flag2;
```

Input

<i>library</i>	literal, name of library.																				
<i>file</i>	optional literal, name of source file to be updated or added.																				
<i>flags</i>	optional literal preceded by '-', controls operation of library update. To control handling of path information on source filenames: <table><tbody><tr><td><i>-addpath</i></td><td>add paths to entries without paths and expand relative paths.</td></tr><tr><td><i>-</i></td><td>reset all paths using a normal file search.</td></tr><tr><td><i>gausspath</i></td><td>(default) leave all path information untouched.</td></tr><tr><td><i>-</i></td><td></td></tr><tr><td><i>leavepath</i></td><td>drop all path information.</td></tr><tr><td><i>-nopath</i></td><td></td></tr></tbody></table> <p>To specify a library update or a complete library build:</p> <table><tbody><tr><td><i>-update</i></td><td>(default) update the symbol information for the specified file only.</td></tr><tr><td><i>-build</i></td><td>update the symbol information for every library entry by compiling the actual source file.</td></tr><tr><td><i>-delete</i></td><td>delete a file from the library.</td></tr><tr><td><i>-list</i></td><td>list files in a library.</td></tr></tbody></table>	<i>-addpath</i>	add paths to entries without paths and expand relative paths.	<i>-</i>	reset all paths using a normal file search.	<i>gausspath</i>	(default) leave all path information untouched.	<i>-</i>		<i>leavepath</i>	drop all path information.	<i>-nopath</i>		<i>-update</i>	(default) update the symbol information for the specified file only.	<i>-build</i>	update the symbol information for every library entry by compiling the actual source file.	<i>-delete</i>	delete a file from the library.	<i>-list</i>	list files in a library.
<i>-addpath</i>	add paths to entries without paths and expand relative paths.																				
<i>-</i>	reset all paths using a normal file search.																				
<i>gausspath</i>	(default) leave all path information untouched.																				
<i>-</i>																					
<i>leavepath</i>	drop all path information.																				
<i>-nopath</i>																					
<i>-update</i>	(default) update the symbol information for the specified file only.																				
<i>-build</i>	update the symbol information for every library entry by compiling the actual source file.																				
<i>-delete</i>	delete a file from the library.																				
<i>-list</i>	list files in a library.																				

To control the symbol type information placed in the library file:

- `-strong` (default) use strongly typed symbol entries.
- `-weak` save no type information. This should only be used to build a library compatible with a previous version of **GAUSS**.

To control location of temporary files for a complete library build:

- `-tmp` (default) use the directory pointed to by the `tmp_path` configuration variable. If `tmp_path` is not defined, `lib` will look for a `tmp` environment variable.
- `-disk` use the same directory listed in the `lib_path` configuration variable.

Example

Let us suppose that you have a file named `myprocs.gss` located in your GAUSS `src` directory. Let us further suppose that you would like to create a new library named `mylibrary`. You could accomplish that task like this:

```
lib mylibrary myprocs.gss;
```

Now that this library has been created, you could add other files in the same manner. To add a file named `mystats.gss` would look like this:

```
lib mylibrary mystats.gss;
```

lib

This second command will add the file `mystats.gss` to the `mylibrary` which was created in the first step above. It will not overwrite or replace the library.

You may print the list of files contained in the library by using the `-list` flag. Entering the command:

```
lib mylibrary -list;
```

at the GAUSS command line will produce the output similar to:

```
Listing library:  mylibrary.lcg
                  myprocs.gss
                  mystats.gs
```

If you add procedures to one of the files in your library, you will need to update the library to reflect these new changes. Continuing with the example from above, if you added some new procedures to the file `mystats.gss`, you could update the `mylibrary` library with the following command:

```
lib mylibrary mystats.gss -update;
```

Note that, as in the command above, the `-update` flag must be used with a file. To update, or rebuild the references for all files in the library, use the `-build` flag.

```
lib mylibrary -build;
```

Remarks

The library management functionality offered by the `lib` command can also be accomplished interactively with windows and buttons, using the Library Tool in the user interface. See **THE LIBRARY TOOL**, CHAPTER 1, for more information on using the Library Tool.

The flags can be shortened to one or two letters, as long as they remain unique—for example, `-b` to `-build` a library, `-li` to list files in a library.

If the filenames include a full path, the compilation process is faster because no unnecessary directory searching is needed during the autoloading process. The default path handling adds a path to each file listed in the library and also expands any relative paths so the system will work from any drive or subdirectory.

When a path is added to a filename containing no path information, the file is searched for on the current directory and then on each subdirectory listed in *src_path*. The first path encountered that contains the file is added to the filename in the library entry.

See Also

[library](#)

library

Purpose

Sets up the list of active libraries.

Format

```
library lib1 [[,lib2,lib3,lib4...]];
library;
```

Example

Example 1

If no arguments are given, the list of current libraries will be printed out. For example:

```
library;
```

will produce output similar to:

library

```
Library path: C:\gauss17\gaussplot\lib  
              C:\gauss17\lib
```

```
Libraries:    C:\gauss17\lib\user.lcg  
              C:\gauss17\lib\gauss.lcg
```

Example 2

Load multiple libraries by passing a comma-separated list of library names.

```
library pgraph, cmlmt;
```

After executing the code above, entering the `library` command without any arguments will produce output similar to this:

```
Library path: C:\gauss17\gaussplot\lib  
              C:\gauss17\lib  
  
Libraries:    C:\gauss17\lib\user.lcg  
              C:\gauss17\lib\pgraph.lcg  
              C:\gauss17\lib\cmlmt.lcg  
              C:\gauss17\lib\gauss.lcg
```

The output from the `library` command above is printed in the order in which GAUSS will search. For this particular example, GAUSS will first search the `user` library, then the `pgraph` library, followed by the `cmlmt` library and finally the `gauss` library.

Example 3

Loading a library or list of libraries with the `library` command will also close any open libraries other than `user` and `gauss` which are always loaded.

Continuing with the last example in which the `user`, `pgraph`, `cmlmt` and `gauss` libraries were open, executing the command:

```
library pgraph, tsmt;
```

would open `tsmt`, while closing `pgraph` and `cmlmt`.

Remarks

For more information about the library system, see **LIBRARIES**, CHAPTER 1 .

The required extension for library files is `.lcg`.

Library searching

If a list of library names is given, they will be the new set of active libraries. The two default libraries are `user.lcg` and `gauss.lcg`. Unless otherwise specified, `user.lcg` will be searched first and `gauss.lcg` will be searched last. Any other user-specified libraries will be searched after `user.lcg` in the order they were entered in the `library` statement.

If the statement:

```
y = myProc(x) ;
```

is encountered in a program, **myProc** will be searched for in the active libraries. If it is found, it will be compiled. If it cannot be found in a library, the deletion state determines how it is handled:

<code>autodelete on</code>	search for myproc.g
<code>autodelete off</code>	return Undefined symbol error message

If **myProc** calls **myRegress** and **myRegress** calls **myUtil** and they are all in separate files, they will all be found by the autoloader.

The source browser and the help facility will search for **myProc** in exactly the same sequence as the autoloader.

Library file contents

Library files are simple ASCII files that you can create with a text editor. Here is an example:

library

```
/*  
** This is a GAUSS library file.  
*/
```

```
eig.src  
    eig      : proc  
    eigsym   : proc  
    _eigerr  : matrix  
svd.src  
    cond     : proc  
    pinv     : proc  
    rank     : proc  
    svd      : proc  
    _svdtol  : matrix
```

The lines not indented are the file names. The lines that are indented are the symbols defined in that file. As you can see, a **GAUSS** library is a dictionary of files and the global symbols they contain.

Any line beginning with `/*`, `**`, or `*/` is considered a comment. Currently, `//` comments are not supported in library files. Blank lines are okay.

To make the autoloading process more efficient, you can put the full pathname for each file in the library:

```
/gauss/src/eig.src  
    eig      : proc  
    eigsym   : proc  
    _eigerr  : matrix  
/gauss/src/svd.src  
    cond     : proc  
    pinv     : proc  
    rank     : proc  
    svd      : proc  
    _svdtol  : matrix
```

Here's a debugging hint. If your program is acting strange and you suspect it is autoloading the wrong copy of a procedure, use the Library Tool on the Source Page, or the CTRL+F1 hotkey to locate the suspected function. It will use the same search path that the autoloader uses.

See Also

[declare](#), [external](#), [lib](#), [proc](#)

#lineson, #linesoff

Purpose

The `#lineson` command causes **GAUSS** to embed line number and file name records in a program for the purpose of reporting the location where an error occurs. The `#linesoff` command causes **GAUSS** to stop embedding line and file records in a program.

Format

```
#lineson  
#linesoff
```

Remarks

In the "lines on" mode, **GAUSS** keeps track of line numbers and file names and reports the location of an error when an execution time error occurs. In the "lines off" mode, **GAUSS** does not keep track of lines and files at execution time. During the compile phase, line numbers and file names will always be given when errors occur in a program stored in a disk file.

It is easier to debug a program when the locations of errors are reported, but this slows down execution. In programs with several scalar operations, the time spent tracking line numbers and file names is most significant.

linsolve

These commands have no effect on interactive programs (that is, those typed in the window and run from the command line), since there are no line numbers in such programs.

Line number tracking can be turned on and off through the user interface, but the `#lineson` and `#linesoff` commands will override that.

The line numbers and file names given at run-time will reflect the last record encountered in the code. If you have a mixture of procedures that were compiled without line and file records and procedures that were compiled with line and file records, use the `trace` command to locate exactly where the error occurs.

The Currently active call error message will always be correct. If it states that it was executing procedure `xyz` at line number `nnn` in file `ABC` and `xyz` has no line `nnn` or is not in file `ABC`, you know that it just did not encounter any line or file records in `xyz` before it crashed.

When using `#include`'d files, the line number and file name will be correct for the file the error was in within the limits stated above.

See Also

[trace](#)

linsolve

Purpose

Solves $Ax = b$ using the inverse function.

Format

```
 $x = \text{linsolve}(b, A);$ 
```

Input

b	$N \times K$ matrix.
A	$N \times N$ matrix.

Output

x	$N \times K$ matrix, the linear solution of b/A for each column in b .
-----	--

Remarks

linsolve solves for x by computing $\text{inv}(A)*b$. If A is square and b contains more than 1 column, it is much faster to use **linsolve** than the $/$ operator. However, while faster, there is some sacrifice in accuracy.

A test shows **linsolve** to be accurate to within approximately $1.2\text{e-}11$, while the slash operator $'/'$ is accurate to within approximately $4\text{e-}13$. However, the accuracy sacrifice can be much greater for poorly conditioned matrices.

Example

```
b = { 2, 3, 4 };
A = { 10 2 3, 6 14 2, 1 1 9 };
x = linsolve(b,A);
print x
```

```
0.04586330
0.13399281
0.42446043
```

See Also

[grsol](#), [qrtsol](#), [solpd](#), [cholsol](#)

listwise (dataloop)

listwise (dataloop)

Purpose

Controls listwise deletion of missing values.

Format

```
listwise [[read]] [[write]];
```

Remarks

If **read** is specified, the deletion of all rows containing missing values happens immediately after reading the input file and before any transformations. If **write** is specified, the deletion of missing values happens after any transformations and just before writing to the output file. If no **listwise** statement is present, rows with missing values are not deleted.

The default is **read**.

ln

Purpose

Computes the natural log of all elements of x .

Format

```
 $y = \ln(x);$ 
```

Input

x	$N \times K$ matrix or N -dimensional array.
-----	--

Output

y NxK matrix or N-dimensional array containing the natural log values of the elements of x .

Remarks

ln is defined for $x \neq 0$.

If x is negative, complex results are returned.

You can turn the generation of complex numbers for negative inputs on or off in the **GAUSS** configuration file, and with the **sysstate** function, case 8. If you turn it off, **ln** will generate an error for negative inputs.

If x is already complex, the complex number state doesn't matter; **ln** will compute a complex result.

x can be any expression that returns a matrix.

Example

```
y = ln(16);
```

```
y = 2.7725887
```

See Also

[log](#)

Incdfbvn

Purpose

Computes natural log of bivariate Normal cumulative distribution function.

lncdfbvn2

Format

`y = lncdfbvn(x1, x2, r);`

Input

<code>x1</code>	NxK matrix, abscissae.
<code>x2</code>	LxM matrix, abscissae.
<code>r</code>	PxQ matrix, correlations.

Output

<code>y</code>	max(N,L,P) x max(K,M,Q) matrix: <div><code>ln Pr (X < x1, X < x2 r)</code></div>
----------------	---

Remarks

`x1`, `x2`, and `r` must be ExE conformable.

Source

`lncdfn.src`

See Also

[cdfbvn](#), [lncdfmvm](#)

lncdfbvn2

Purpose

Returns natural log of standardized bivariate Normal cumulative distribution function of a bounded rectangle.

Format

$$y = \text{lncdfbvn2}(h, \text{ } dh, \text{ } k, \text{ } dk, r);$$

Input

h	Nx1 vector, upper limits of integration for variable 1.
dh	Nx1 vector, increments for variable 1.
k	Nx1 vector, upper limits of integration for variable 2.
dk	Nx1 vector, increments for variable 2.
r	Nx1 vector, correlation coefficients between the two variables.

Output

y	Nx1 vector, the log of the integral from h, k to $h+dh, k+dk$ of the standardized bivariate Normal distribution.
-----	--

Remarks

Scalar input arguments are okay; they will be expanded to Nx1 vectors.

lncdfbvn2 will abort if the computed integral is negative.

lncdfbvn2 computes an error estimate for each set of inputs-the real integral is $\exp(y) \pm err$. The size of the error depends on the input arguments. If **trap 2** is set, a warning message is displayed when $err \geq \exp(y)/100$.

For an estimate of the actual error, see **cdfBvn2e**.

lncdfmvn

Example

Example 1

```
lncdfbvn2(1,1,1,1,0.5);
```

produces:

```
-3.2180110258198771e+000
```

Example 2

```
trap 0,2;  
lncdfbvn2(1,1e-15,1,1e-15,0.5);
```

produces:

```
-7.1171016046360151e+001
```

Example 3

```
trap 2,2;  
lncdfbvn2(1,-1e-45,1,1e-45,0.5);
```

produces:

```
WARNING: Dubious accuracy from lncdfbvn2:  
0.000e+000 +/- 2.8e-060  
-INF
```

See Also

[cdfbvn2](#), [cdfbvn2e](#)

lncdfmvn

Purpose

Computes natural log of multivariate Normal cumulative distribution function.

Format

```
y = lncdfmvn(x, r);
```

Input

x	$K \times L$ matrix, abscissae.
r	$K \times K$ matrix, correlation matrix.

Output

y	$L \times 1$ vector, $\ln \Pr(X < x r)$
-----	--

Remarks

You can pass more than one set of abscissae at a time; each column of x is treated separately.

Source

lncdfn.src

See Also

[cdfmvn](#), [lncdfbvn](#)

lncdfn

Purpose

Computes natural log of Normal cumulative distribution function.

lncdfn2

Format

$y = \text{lncdfn}(x);$

Input

x NxK matrix or N-dimensional array, abscissae.

Output

y NxK matrix or N-dimensional array,

$\ln \Pr(X < x)$

Source

lncdfn.src

lncdfn2

Purpose

Computes natural log of interval of Normal cumulative distribution function.

Format

$y = \text{lncdfn2}(x, r);$

Input

x MxN matrix, abscissae.

r KxL matrix, ExE conformable with x , intervals.

Output

y

$\max(M,K) \times \max(N,L)$ matrix, the log of the integral from x to $x+dx$ of the Normal distribution, i.e.,

$$\ln \Pr(x < X < x+dx)$$

Remarks

The relative error is:

$ x < 1$	and	$dx < 1$	$\pm 1e-14$
$1 < x < 37$	and	$ dx < 1/ x $	$\pm 1e-13$
$\min(x, x+dx) > -37$	and	$y > -690$	$\pm 1e-11$ or better

A relative error of $\pm 1e-14$ implies that the answer is accurate to better than ± 1 in the 14th digit after the decimal point.

Example

```
print
lncdfN2(-10, 29);
```

```
-7.6198530241605269e-24
```

```
print
lncdfN2(0, 1);
```

```
-1.0748623268620716e+00
```

lncdfnc

```
print
lncdfN2(5,1);

-1.5068446096529453e+01
```

Source

lncdfn.src

See Also

[cdfn2](#)

lncdfnc

Purpose

Computes natural log of complement of Normal cumulative distribution function.

Format

$y = \text{lncdfnc}(x);$

Input

x NxK matrix, abscissae.

Output

y NxK matrix,
 $\ln (1 - \Pr (X < x))$

Source

lncdfn.src

Infact

Purpose

Computes the natural log of the factorial function and can be used to compute log gamma.

Format

```
y = lnfact(x);
```

Input

x	NxK matrix or N-dimensional array, all elements must be positive.
-----	---

Output

y	NxK matrix containing the natural log of the factorial of each of the elements in x .
-----	---

Remarks

For integer x , this is (approximately) $\ln(x!)$. However, the computation is done using a formula, and the function is defined for noninteger x .

In most formulae in which the factorial operator appears, it is possible to avoid computing the factorial directly, and to use **lnfact** instead. The advantage of this is that **lnfact** does not have the overflow problems that the factorial (!) operator has.

Infact

For $x > 1$, this function has at least 6 digit accuracy, for $x > 4$ it has at least 9 digit accuracy, and for $x > 10$ it has at least 12 digit accuracy. For $0 < x < 1$, accuracy is not known completely but is probably at least 6 digits.

Sometimes log gamma is required instead of log factorial. These functions are related by:

```
lngamma(x) = lnfact(x-1);
```

Example

```
let x = 100 500 1000;  
y = lnfact(x);
```

```
      363.73938  
y = 2611.3305  
      5912.1282
```

Source

lnfact.src

See Also

[gamma](#)

Technical Notes

For $x > 1$, Stirling's formula is used.

For $0 < x \leq 1$, `ln(gamma(x+1))` is used.

lngammacplx

Purpose

Returns the natural log of the Gamma function.

Format

```
 $f = \text{lngammacplx}(z);$ 
```

Input

z NxK matrix; z may be complex.

Output

f NxK matrix.

Remarks

Note that **lngammacplx**(z) may yield a result with a different imaginary part than **ln(gammacplx**(z)). This is because **lngammacplx**(z) returns the value of the logarithm of **gamma**(z) on the corresponding branch of the complex plane, while a call to **ln**(z) always returns a function value with an imaginary part within $[-\pi, \pi]$. Hence the imaginary part of the result can differ by a multiple of 2π . However, **exp(lngammacplx**(z)) = **gammacplx**(z). This routine uses a Lanczos series approximation for the complex **ln(gamma)** function.

References

1. C. Lanczos, SIAM JNA 1, 1964. pp. 86-96.
2. Y. Luke, "The Special ... approximations," 1969 pp. 29-31.
3. Y. Luke, "Algorithms ... functions," 1977.

Inpdfmvn

4. J. Spouge, SIAM JNA 31, 1994. pp. 931.
5. W. Press, "Numerical Recipes."
6. S. Chang, "Computation of special functions," 1996.
7. P. Godfrey, "A note on the computation of the convergent Lanczos complex Gamma approximation."
8. Original code by Paul Godfrey

Inpdfmvn

Purpose

Computes multivariate Normal log-probabilities.

Format

```
z = lnpdfmvn(x, s);
```

Input

x	$N \times K$ matrix, data.
s	$K \times K$ matrix, covariance matrix.

Output

z	$N \times 1$ vector, log-probabilities.
-----	---

Remarks

This computes the multivariate Normal log-probability for each row of x .

Source

lnpdfn.src

lnpdfmvt

Purpose

Computes multivariate Student's t log-probabilities.

Format

```
 $z = \text{lnpdfmvt}(x, s, nu);$ 
```

Input

x	NxK matrix, data.
s	KxK matrix, covariance matrix.
nu	scalar, degrees of freedom.

Output

z	Nx1 vector, log-probabilities.
-----	--------------------------------

Source

lnpdfn.src

See Also

[lnpdft](#)

lnpdfn

lnpdfn

Purpose

Computes standard Normal log-probabilities.

Format

```
z = lnpdfn(x);
```

Input

x NxK matrix or N-dimensional array, data.

Output

z NxK matrix or N-dimensional array, log-probabilities.

Remarks

This computes the log of the scalar Normal density function for each element of **x**. **z** could be computed by the following **GAUSS** code:

```
z = -ln(sqrt(2*pi)) - x .* x / 2;
```

For multivariate log-probabilities, see **lnpdfmvn**.

Example

```
x = { -2, -1, 0, 1, 2 };  
z = lnpdfn(x);
```

```
-2.9189385  
-1.4189385
```

```
z = -0.9189385  
    -1.4189385  
    -2.9189385
```

lnpdf_t

Purpose

Computes Student's t log-probabilities.

Format

```
z = lnpdft(x, nu);
```

Input

x	NxK matrix, data.
nu	scalar, degrees of freedom.

Output

z	NxK matrix, log-probabilities.
-----	--------------------------------

Remarks

This does not compute the log of the joint Student's t pdf. Instead, the scalar Normal density function is computed element-by-element.

For multivariate probabilities with covariance matrix see **lnpdf_{fmvt}**.

See Also

[lnpdf_{fmvt}](#)

load, loadf, loadk, loadm, loadp, loads

load, loadf, loadk, loadm, loadp, loads

Purpose

Loads from a disk file.

Format

```
load [[path=path]]x, y[ ]=filename, z = filename;
```

Remarks

All the `loadxx` commands use the same syntax—they only differ in the types of symbols you use them for:

<code>load</code> ,	matrix
<code>loadm</code>	
<code>loads</code>	string
<code>loadf</code>	function (<code>fn</code>)
<code>loadk</code>	keyword (<code>keyword</code>)
<code>loadp</code>	procedure (<code>proc</code>)

If no filename is given, as with `x` above, then the symbol name the file is to be loaded into is used as the filename, and the proper extension is added.

If more than one item is to be loaded in a single statement, the names should be separated by commas.

The filename can be either a literal or a string. If the filename is in a string variable, then the `^` (caret) operator must precede the name of the string, as in:

```
filestr = "mydata/char";  
loadm x = ^filestr;
```

If no extension is supplied, the proper extension for each type of file will be used automatically as follows:

<code>load</code>	<code>.fmt</code> - matrix file or delimited ASCII file
<code>loadm</code>	<code>.fmt</code> - matrix file or delimited ASCII file
<code>loads</code>	<code>.fst</code> - string file
<code>loadf</code>	<code>.fcg</code> - user-defined function (<code>fn</code>) file
<code>loadk</code>	<code>.fcg</code> - user-defined keyword (<code>keyword</code>) file
<code>loadp</code>	<code>.fcg</code> - user-defined procedure (<code>proc</code>) file

These commands also signal to the compiler what type of object the symbol is so that later references to it will be compiled correctly.

A dummy definition must exist in the program for each symbol that is loaded in using `loadf`, `loadk`, or `loadp`. This resolves the need to have the symbol initialized at compile time. When the load executes, the dummy definition will be replaced with the saved definition:

```
proc corrrmat;
endp;

loadp corrrmat;
y = corrrmat;

keyword regress(x); endp;
loadk regress;
regress x on y z t from data01;

fn sqrd=;
loadf sqrd;
y = sqrd(4.5);
```

To load **GAUSS** files created with the `save` command, no brackets are used with the symbol name.

load, loadf, loadk, loadm, loadp, loads

If you use `save` to save a scalar error code 65535 (i.e., `error(65535)`), it will be interpreted as an empty matrix when you `load` it again.

ASCII data files

To load ASCII data files, square brackets follow the name of the symbol.

Numbers in ASCII files must be delimited with spaces, commas, tabs, or newlines. If the size of the matrix to be loaded is not explicitly given, as in:

```
load x[] = data.asc;
```

GAUSS will load as many elements as possible from the file and create an Nx1 matrix. This is the preferred method of loading ASCII data from a file, especially when you want to verify if the load was successful. Your program can then see how many elements were actually loaded by testing the matrix with the `rows` command, and if that is correct, the Nx1 matrix can be **reshape**'d to the desired form. You could, for instance, put the number of rows and columns of the matrix right in the file as the first and second elements and **reshape** the remainder of the vector to the desired form using those values.

If the size of the matrix is explicitly given in the `load` command, then no checking will be done. If you use:

```
load x[500,6] = data.asc;
```

GAUSS will still load as many elements as possible from the file into an Nx1 matrix and then automatically reshape it using the dimensions given.

If you `load` data from a file, `data.asc`, which contains nine numbers (1 2 3 4 5 6 7 8 9), then the resulting matrix will be as follows:

```
load x[1,9] = data.asc;
```

```
x = 1 2 3 4 5 6 7 8 9
```

```
load x[3,3] = data.asc;
```



```
      1 2 3
x = 4 5 6
      7 8 9
```

```
load x[2,2] = data.asc;
```

```
x = 1 2
    3 4
```

```
load x[2,9] = data.asc;
```

```
x = 1 2 3 4 5 6 7 8 9
    1 2 3 4 5 6 7 8 9
```

```
load x[3,5] = data.asc;
```

```
      1 2 3 4 5
x = 6 7 8 9 1
    2 3 4 5 6
```

`load` accepts pathnames. The following is legal:

```
loadm k = /gauss/x;
```

This will load /gauss/x.fmt into *k*.

If the **path=** subcommand is used with `load` and `save`, the path string will be remembered until changed in a subsequent command. This path will be used whenever none is specified. There are four separate paths for:

1. `load, loadm`
2. `loadf, loadp`
3. `loads`
4. `save`

load, loadf, loadk, loadm, loadp, loads

Setting any of the four paths will not affect the others. The current path settings can be obtained (and changed) with the **sysstate** function, cases 4-7.

```
loadm path = /data;
```

This will change the **loadm** path without loading anything.

```
load path = /gauss x,y,z;
```

This will load **x.fmt**, **y.fmt**, and **z.fmt** using **/gauss** as a path. This path will be used for the next load if none is specified.

The **load** path or **save** path can be overridden in any particular **load** or **save** by putting an explicit path on the filename given to **load** from or **save** to as follows:

```
loadm path = /miscdata;  
loadm x = /data/mydata1, y, z = hisdata;
```

In the above program:

/data/mydata1.fmt would be loaded into a matrix called **x**.

/miscdata/y.fmt would be loaded into a matrix called **y**.

/miscdata/hisdata.fmt would be loaded into a matrix called **z**.

```
oldmpath = sysstate(5, "/data");  
load x, y;  
call sysstate(5, oldmpath);
```

This will get the old **loadm** path, set it to **/data**, load **x.fmt** and **y.fmt**, and reset the **loadm** path to its original setting.

See Also

[loadd](#), [dataload](#), [save](#), [let](#), [con](#), [cons](#), [sysstate](#)

loadarray

Purpose

Loads an N-dimensional array from a disk file.

Format

```
loadarray [[path=path]] x, y = filename;
```

Remarks

If no filename is given, as with *x* above, then the symbol name the file is to be loaded into is used as the filename, and the proper extension is added.

If more than one item is to be loaded in a single statement, the names should be separated by commas.

The filename can be either a literal or a string. If the filename is in a string variable, then the ^ (caret) operator must precede the name of the string, as in:

```
filestr = "mydata/adat";  
loadarray x = ^filestr;
```

If no extension is supplied, then an `.fmt` extension will be assumed.

`loadarray` accepts pathnames. The following is legal:

```
loadarray k = /gauss/a;
```

This will load `/gauss/a.fmt` into *k*.

If the **path=** subcommand is used, the path string will be remembered until changed in a subsequent command. This path will be used for all `loadarray`, `loadm`, and `load` calls whenever none is specified.

loadarray

The current path setting can be obtained (and changed) with the **sysstate** function, case 5.

```
loadarray path = /data;
```

This will change the **loadarray** path without loading anything.

```
loadarray path = /gauss a,b,c;
```

This will load **a.fmt**, **b.fmt**, and **c.fmt** using **/gauss** as a path. This path will be used for the next **loadarray**, **loadm**, or **load** call if none is specified.

The **load** path or **save** path can be overridden in any particular **load** or **save** by putting an explicit path on the filename given to **load** from or **save** to as follows:

```
loadarray path = /miscdata;  
loadarray a = /data/mydata1, b, c = hisdata;
```

In the above program:

/data/mydata1.fmt would be loaded into an array called **a**.

/miscdata/b.fmt would be loaded into an array called **b**.

/miscdata/hisdata.fmt would be loaded into an array called **c**.

```
oldarraypath = sysstate(5, "/data");  
loadarray a, b;  
call sysstate(5, oldarraypath);
```

This will get the old **loadarray** path, set it to **/data**, load **a.fmt** and **b.fmt**, and reset the **loadarray** path to its original setting.

See Also

[load](#), [loadm](#), [save](#), [let](#), [sysstate](#)

loadadd

Purpose

Loads specified variables from a GAUSS data set or matrix file.

Format

```
y = loadadd(dataset);  
y = loadadd(dataset, varnames);
```

Input

<i>dataset</i>	string, name of data set (.dat) or matrix file (.fmt).
<i>varnames</i>	Formula string , indicating which variable names to load from the data set E.g. ".", include all variables; E.g. "Income + Limit", include "Income" and "Limit"; E.g. ". - Cards", '-' means exclude "Cards".

Output

<i>y</i>	NxK matrix of data.
----------	---------------------

Examples

Example 1

Load all contents of a data set

```
//Create file name with full path
```

loadadd

```
file = getGAUSShome() $+ "examples/credit.dat";

//Load all rows from all columns of the dataset
y = loadadd(file);

//Print the first three rows of 'y'
print y[1:3,.];
```

After the above code, the following output should be printed to the program input/output window.

14.8910	3606.00	283.000	2.00000	34.0000
106.025	6645.00	483.000	3.00000	82.0000
104.593	7075.00	514.000	4.00000	71.0000

Example 2

Load specified variables from a dataset

```
//Load all variables with a formula string
dat1 = loadadd(file, "." );

//Load all observations of 'Balance' and 'Limit'
dat2 = loadadd(file, "Balance + Limit" );

//Load all variables EXCEPT for 'Cards'
dat3 = loadadd(file, ". - Cards" );

//Print first three rows of each matrix
print "All variables: " dat1[1:3,.];
print "Balance and Limit: " dat2[1:3,.];
print "All except Cards: " dat3[1:3,.];
```

After the above code,

All variables:

14.891	3606.00	283.00	2.0000	34.000	11.000
1.0000	1.0000	2.0000	3.0000	333.000	
106.03	6645.00	483.00	3.0000	82.000	15.000
2.0000	2.0000	2.0000	2.0000	903.000	
104.59	7075.00	514.00	4.0000	71.000	11.000
1.0000	1.0000	1.0000	2.0000	580.000	

Balance and Limit:

333.000	3606.00
903.000	6645.00
580.000	7075.00

All except Cards:

14.8910	3606.00	283.00	34.000	11.000	1.0000
1.0000	2.0000	3.0000	333.000		
106.025	6645.00	483.00	82.000	15.000	2.0000
2.0000	2.0000	2.0000	903.000		
104.593	7075.00	514.00	71.000	11.000	1.0000
1.0000	1.0000	2.0000	580.000		

Example 3

Load all columns of a GAUSS matrix file, `.fmt`. No variable names are stored in `.fmt` files. GAUSS allows the use of 'X1, X2, X2...XP' to reference variables in a `.fmt` file.

```
//Create a matrix
x = rndn(10, 4);

//Save to a matrix file, 'x.fmt'
save x;
```

loadadd

```
//Load all columns of 'x.fmt'
x_2 = loadadd("x.fmt");
```

Example 4

Load specified columns of a GAUSS matrix file, .fmt.

```
//Create a matrix
x = rndn(10, 4);

//Save to a matrix file, 'x.fmt'
save x;

//Load columns 2 and 4 from 'x.fmt'
x_2 = loadadd("x.fmt", "X2 + X4");
```

Remarks

- Since **loadadd** will load the entire dataset at once, the data set must be small enough to fit in memory. To read chunks of a dataset in an iterative manner, use **dataopen** and **readr**.
- If *dataset* is a null string or 0, the data set `temp.dat` will be loaded.
- To load a matrix file, use an `.fmt` extension on *dataset*.
- The supported data set types are [CSV](#), [XLS](#), [XLSX](#), [HDF5](#), [FMT](#), [DAT](#).
- For HDF5 file, the *dataset* must include [file schema](#) and both file name and data set name must be provided, e.g. **loadadd** (`"h5://C:/gauss17/examples/testdata.h5/mydata"`).

Source

saveload.src

See also

[Formula String](#), [dataopen](#), [readr](#), [saved](#)

Globals

___maxvec

loadstruct

Purpose

Loads a structure into memory from a file on the disk.

Format

```
{ instance, retcode } = loadstruct(file_name, structure_  
type);
```

Input

<i>file_name</i>	string, name of file containing structure.
<i>structure_</i> <i>type</i>	string, structure type.

Output

<i>instance</i>	instance of the structure.
<i>retcode</i>	scalar, 0 if successful, otherwise 1.

Remarks

instance can be an array of structures.

Example

```
#include ds.sdf  
struct DS p3;  
  
{ p3, retc } = loadstruct("p2", "ds");
```

loadwind

loadwind

Purpose

Load a previously saved graphic panel configuration. Note: This function is for use with the deprecated PQG graphics.

Library

pgraph

Format

```
err = loadwind(namestr);
```

Input

<i>namestr</i>	string, name of file to be loaded.
----------------	------------------------------------

Output

<i>err</i>	scalar, 0 if successful, 1 if graphic panel matrix is invalid. Note that the current graphic panel configuration will be overwritten in either case.
------------	--

Source

pwindow.src

Globals

_pwindmx

See Also

[savewind](#)

local

Purpose

Declare variables that are to exist only inside a procedure.

Format

```
local x, y, f:proc;
```

Remarks

The statement above would place the names x , y , and f in the local symbol table for the current procedure being compiled. This statement is legal only between the `proc` statement and the `endp` statement of a procedure definition.

These symbols cannot be accessed outside of the procedure.

The symbol f in the statement above will be treated as a procedure whenever it is accessed in the current procedure. What is actually passed in is a pointer to a procedure.

See **PROCEDURES AND KEYWORDS**, CHAPTER 1.

See Also

[proc](#)

locate

Purpose

Positions the cursor in the window.

loess

Format

```
locate m, n;
```

Remarks

`locate` locates the cursor in the current output window.

m and *n* denote the row and column, respectively, at which the cursor is to be located.

The origin (1,1) is the upper left corner.

m and *n* may be any expressions that return scalars. Nonintegers will be truncated to an integer.

Example

```
r = csrlin;  
c = csrcol;  
cls;  
locate r,c;
```

In this example the window is cleared without affecting the cursor position.

See Also

[csrlin](#), [csrcol](#)

loess

Purpose

Computes coefficients of locally weighted regression.

Format

```
{ yhat, ys, xs } = loess(depvar, indvars);
```

Input

<i>depvar</i>	Nx1 vector, dependent variable.
<i>indvars</i>	NxK matrix, independent variables.

Global Input

<code>__loess_Span</code>	scalar, degree of smoothing. Must be greater than 2/N. Default = .67777.
<code>__loess_NumEval</code>	scalar, number of points in <i>ys</i> and <i>xs</i> . Default = 50.
<code>__loess_Degree</code>	scalar, if 2, quadratic fit, otherwise linear. Default = 1.
<code>__loess_WgtType</code>	scalar, type of weights. If 1, robust, symmetric weights, otherwise Gaussian. Default = 1.
<code>__output</code>	scalar, if 1, iteration information and results are printed, otherwise nothing is printed.

Output

<i>yhat</i>	Nx1 vector, predicted <i>depvar</i> given <i>indvars</i> .
<i>ys</i>	<code>__loess_numEval</code> x1 vector, ordinate values given abscissae values in <i>xs</i> .
<i>xs</i>	<code>__loess_numEval</code> x1 vector, equally spaced abscissae values.

loessmt

Remarks

Based on Cleveland, William S. "Robust Locally Weighted Regression and Smoothing Scatterplots." JASA, Vol. 74, 1979, 829-836.

Source

loess.src

loessmt

Purpose

Computes coefficients of locally weighted regression.

Include

loessmt.sdf

Format

$\{ \hat{y}, y, x \} = \text{loessmt}(lc0, \text{depvar}, \text{indvars});$

Input

<i>lc0</i>	an instance of a loessmtControl structure, containing the following members:
<i>lc0.Span</i>	scalar, degree of smoothing. Must be greater than 2/N. Default = .67777.
<i>lc0.NumEval</i>	scalar, number of points in <i>ys</i> and <i>xs</i> . Default = 50.
<i>lc0.Degree</i>	scalar, if 2, quadratic fit, otherwise linear. Default = 1.
<i>lc0.WgtType</i>	scalar, type of weights. If 1, robust,

	<i>lc0.output</i>	symmetric weights, otherwise Gaussian. Default = 1. scalar, if 1, iteration information and results are printed, otherwise nothing is printed.
<i>depvar</i>	Nx1 vector, dependent variable.	
<i>indvars</i>	NxK matrix, independent variables.	

Output

<i>yhat</i>	Nx1 vector, predicted <i>depvar</i> given <i>indvars</i> .
<i>ys</i>	<i>lc0.numEval</i> x 1 vector, ordinate values given abscissae values in <i>xs</i> .
<i>xs</i>	<i>lc0.numEval</i> x 1 vector, equally spaced abscissae values.

Remarks

Based on Cleveland, William S. "Robust Locally Weighted Regression and Smoothing Scatterplots." JASA, Vol. 74, 1979, 829-836.

Source

loessmt.src

See Also

[loessmtControlCreate](#)

loessmtControlCreate

loessmtControlCreate

Purpose

Creates default **loessmtControl** structure.

Include

loessmt.sdf

Format

```
c = loessmtControlCreate();
```

Output

c

instance of a **loessmtControl** structure with members set to default values.

Example

```
struct loessmtControl lc;  
lc = loessmtControlCreate();
```

Source

loessmt.src

See Also

[loessmt](#)

log

Purpose

Computes the log of all elements of x .

Format

```
 $y = \mathbf{log}(x);$ 
```

Input

x	$N \times K$ matrix or N -dimensional array.
-----	--

Output

y	$N \times K$ matrix or N -dimensional array containing the log values of the elements of x .
-----	--

Remarks

log is defined for $x \neq 0$.

You can turn the generation of complex numbers for negative inputs on or off in the **GAUSS** configuration file, and with the **sysstate** function, case 8. If you turn it off, **log** will generate an error for negative inputs.

If x is already complex, the complex number state doesn't matter; **log** will compute a complex result.

x can be any expression that returns a matrix.

Example

```
//Create a 3x3 matrix of random uniform integers from 1
```

loglog

```
//to 11
x = round(rndu(3,3)*10+1);
y = log(x);
```

If x is equal to:

4.000	9.000	2.000
5.000	3.000	7.000
2.000	6.000	10.000

Then y will be equal to:

0.602	0.954	0.301
0.699	0.477	0.845
0.301	0.778	1.000

See Also

[ln](#)

loglog

Purpose

Graphs X vs. Y using log coordinates. Note: This function is for use with the deprecated PQG graphics. Use **plotLogLog** instead.

Library

pgraph

Format

```
loglog(x, y);
```

Input

x	Nx1 or NxM matrix. Each column contains the X values for a particular line.
y	Nx1 or NxM matrix. Each column contains the Y values for a particular line.

Source

ploglog.src

See Also

[xy](#), [logx](#), [logy](#)

logx

Purpose

Graphs X vs. Y using log coordinates for the X axis. Note: This function is for use with the deprecated PQG graphics. Use **plotLogX** instead.

Library

pgraph

Format

```
logx(x, y);
```

Input

x	Nx1 or NxM matrix. Each column contains the X values for a particular line.
y	Nx1 or NxM matrix. Each column contains the Y

logy

values for a particular line.

Source

`plogx.src`

See Also

[xy](#), [logy](#), [loglog](#)

logy

Purpose

Graphs X vs. Y using log coordinates for the Y axis. Note: This function is for use with the deprecated PQG graphics. Use **plotLogY** instead.

Library

`pgraph`

Format

```
logy(x, y);
```

Input

<i>x</i>	Nx1 or NxM matrix. Each column represents the X values for a particular line.
<i>y</i>	Nx1 or NxM matrix. Each column represents the Y values for a particular line.

Source

`plogy.src`

See Also

[xy](#), [logx](#), [loglog](#)

loopnextindex

Purpose

Increments an index vector to the next logical index and jumps to the specified label if the index did not wrap to the beginning.

Format

```
loopnextindex lab, i, o [, dim];
```

Input

<i>lab</i>	literal, label to jump to if <code>loopnextindex</code> succeeds.
<i>i</i>	Mx1 vector of indices into an array, where $M \leq N$.
<i>o</i>	Nx1 vector of orders of an N-dimensional array.
<i>dim</i>	scalar [1-M], index into the vector of indices <i>i</i> , corresponding to the dimension to walk through, positive to walk the index forward, or negative to walk backward.

Remarks

If the argument *dim* is given, `loopnextindex` will walk through only the dimension indicated by *dim* in the specified direction. Otherwise, if *dim* is not given, each call to `loopnextindex` will increment *i* to index the next element or subarray of the corresponding array.

loopnextindex

`loopnextindex` will jump to the label indicated by `lab` if the index can walk further in the specified dimension and direction, otherwise it will fall out of the loop and continue through the program.

When the index matches the vector of orders, the index will be reset to the beginning and program execution will resume at the statement following the `loopnextindex` statement.

Example

At its essence, `loopNextIndex` provides a simple way to iterate over the orders of a multi-dimensional array.

```
//The orders of the array
orders = { 2, 3, 4 };

//The starting index of the array
ind = { 1, 1, 1 };

lnilab:
print "ind = " ind;
loopNextIndex lnilab, ind, orders;
```

Running the code above, returns:

```
ind =
  1.000
  1.000
  1.000
ind =
  1.000
  1.000
  2.000
ind =
  1.000
```

```
1.000
3.000
ind =
1.000
1.000
4.000
ind =
1.000
2.000
1.000
ind =
1.000
2.000
2.000
ind =
1.000
2.000
3.000

...continuing on to end with...

ind =
2.000
3.000
4.000
```

This next example uses the variable *ind* to iterate over and make assignments to the array, *a*.

```
orders = { 2,3,4,5,6,7 };
a = arrayalloc(orders,0);
ind = { 1,1,1,1 };

loopni:
```

lower

```
setarray a, ind, rdn(6,7);  
loopnextindex loopni, ind, orders;
```

This example sets each 6x7 subarray of array *a*, by incrementing the index at each call of `loopnextindex` and then going to the label `loopni`. When *ind* cannot be incremented, the program drops out of the loop and continues.

```
ind = { 1,1,4,5 };  
  
loopni2:  
  
setarray a, ind, rdn(6,7);  
loopnextindex loopni2, ind, orders, 2;
```

Using the array and vector of orders from the example above, this example increments the second value of the index vector *ind* during each call to `loopnextindex`. This loop will set the 6x7 subarrays of *a* that begin at [1,1,4,5,1,1], [1,2,4,5,1,1], and [1,3,4,5,1,1], and then drop out of the loop.

See Also

[nextindex](#), [previousindex](#), [walkindex](#)

lower

Purpose

Converts a string or character matrix to lowercase.

Format

```
y = lower(x);
```


Input

x	string or NxK matrix of character data to be converted to lowercase.
-----	--

Output

y	string or NxK matrix which contains the lowercase equivalent of the data in x .
-----	---

Remarks

If x is a numeric matrix, y will contain garbage. No error message will be generated since **GAUSS** does not distinguish between numeric and character data in matrices.

Example

```
x = "MATH 401";  
y = lower(x);  
print y;
```

produces:

```
math 401
```

The **lower** function can be useful when performing case insensitive string comparisons. If you have a program that runs different code depending upon the variable name in a **GAUSS** dataset or spreadsheet file, you or your colleagues may want to analyze data with inconsistent use of case.

```
var1 = "Consumption";  
  
if lower(var1) == "gdp";  
    //code for gdp branch  
else if lower(var1) == "consumption";
```

lowmat, lowmat1

```
//code for consumption branch  
endif;
```

Using the **lowmat** function, the code above will operate correctly whether *var1* is Consumption, CONSUMPTION or consumption.

See Also

[upper](#)

lowmat, lowmat1

Purpose

Returns the lower portion of a matrix. **lowmat** returns the main diagonal and every element below. **lowmat1** is the same except it replaces the main diagonal with ones.

Format

```
L = lowmat(x);  
L = lowmat1(x);
```

Input

x	NxN matrix.
-----	-------------

Output

L	NxN matrix containing the lower elements of the matrix. The upper elements are replaced with zeros. lowmat returns the main diagonal intact. lowmat1 replaces the main diagonal with ones.
-----	--

Remarks

The **lowmat** function along with **upmat1** can be used to extract the LU factors from the return

Example

```
x = { 1 2 -1,
      2 3 -2,
      1 -2 4 };

L = lowmat(x);
L1 = lowmat1(x);
```

The resulting matrices are

```

      1  0  0      1  0  0
L = 2  3  0  L1 = 2  1  0
      1 -2  4      1 -2  1
```

Source

diag.src

See Also

[upmat](#), [upmat1](#), [diag](#), [diagrv](#), [crout](#), [croutp](#)

ltrisol

Purpose

Computes the solution of $Lx = b$ where L is a lower triangular matrix.

lu

Format

```
 $x = \text{ltrisol}(b, L);$ 
```

Input

b	PxK matrix.
L	PxP lower triangular matrix.

Output

x	PxK matrix, solution of $Lx = b$.
-----	------------------------------------

ltrisol applies a forward solve to $Lx = b$ to solve for x . If b has more than one column, each column will be solved for separately, i.e., **ltrisol** will apply a forward solve to $L^*x[:, i] = b[:, i]$.

lu

Purpose

Computes the LU decomposition of a square matrix with partial (row) pivoting, such that: $X = LU$.

Format

```
 $\{ L, U \} = \text{lu}(x);$ 
```

Input

x	NxN square nonsingular matrix.
-----	--------------------------------

Output

l	NxN "scrambled" lower triangular matrix. This is a lower triangular matrix that has been reordered based on the row pivoting.
u	NxN upper triangular matrix.

Example

```
//Set seed for repeatable random numbers
rndseed 13;

//Print format, display 4 digits after decimal point
format /rd 10,4;

A = rndn(3,3);
{ L, U } = lu(A);
A2 = L*U;
```

```
A =
-0.0195    0.4054   -0.0874
-1.2948    0.1734    1.9712
 0.5408   -0.1294    0.7646

L =
 0.0150    1.0000    0.0000
 1.0000    0.0000    0.0000
-0.4177   -0.1414    1.0000

U =
-1.2948    0.1734    1.9712
 0.0000    0.4028   -0.1170
 0.0000    0.0000    1.5714

L*U =
-0.0195    0.4054   -0.0874
-1.2948    0.1734    1.9712
 0.5408   -0.1294    0.7646
```

lusol

See Also

[crout](#), [croutp](#), [chol](#)

lusol

Purpose

Computes the solution of $LUx = b$ where L is a lower triangular matrix and U is an upper triangular matrix.

Format

$x = \text{lusol}(b, L, U);$

Input

b	PxK matrix.
L	PxP lower triangular matrix.
U	PxP upper triangular matrix.

Output

x	PxK matrix, solution of $LUx = b$.
-----	-------------------------------------

Remarks

If b has more than one column, each column is solved for separately, i.e., **lusol** solves $LUx[:, i] = b[:, i]$.

m

machEpsilon

Purpose

Returns the smallest number such that $1+eps > 1$.

Format

```
eps = machEpsilon;
```

Output

<i>eps</i>	scalar, machine epsilon.
------------	--------------------------

Source

`machconst.src`

make (dataloop)

Purpose

Specifies the creation of a new variable within a data loop.

Format

```
make [#] numvar = numeric_expression;  
make $charvar = character_expression;
```

makevars

Remarks

A *numeric_expression* is any valid expression returning a numeric vector. A *character_expression* is any valid expression returning a character vector. If neither '\$' nor '#' is specified, '#' is assumed.

The expression may contain explicit variable names and/or **GAUSS** commands. Any variables referenced must already exist, either as elements of the source data set, as [extern](#)'s, or as the result of a previous [make](#), [vector](#), or **code** statement. The variable name must be unique. A variable cannot be made more than once, or an error is generated.

Example

```
make sqvpt = sqrt(velocity * pressure * temp);
make $ gender = lower(gender);
```

See Also

[vector \(dataloop\)](#)

makevars

Purpose

Creates separate global vectors from the columns of a matrix.

Format

```
makevars(x, vnames, xnames);
```

Input

x	NxK matrix whose columns will be converted into individual vectors.
---	---

<i>vnames</i>	string or Mx1 character vector containing names of global vectors to create. If 0, all names in <i>xnames</i> will be used.
<i>xnames</i>	string or Kx1 character vector containing names to be associated with the columns of the matrix <i>x</i> .

Remarks

If *xnames* = 0, the prefix X will be used to create names. Therefore, if there are 9 columns in *x*, the names will be X1-X9, if there are 10, they will be X01-X10, and so on.

If *xnames* or *vnames* is a string, the individual names must be separated by spaces or commas:

```
vnames = "age pay sex";
```

Since these new vectors are created at execution time, the compiler will not know they exist until after **makevars** has executed once. This means that you cannot access them by name unless you previously **clear** them or otherwise add them to the symbol table. (See **setvars** for a quick interactive solution to this.)

This function is the opposite of **mergevar**.

Example

```
let x[3,3] = 101 35 50000
            102 29 13000
            103 37 18000;
let xnames = id age pay;
let vnames = age pay;
makevars(x, vnames, xnames);
```

Two global vectors, called *age* and *pay*, are created from the columns of *x*.

makewind

```
let x[3,3] = 101 35 50000
            102 29 13000
            103 37 18000;
xnames = "id age pay";
vnames = "age pay";
makevars (x, vnames, xnames);
```

This is the same as the example above, except that strings are used for the variable names.

Source

vars.src

Globals

___vpad

See Also

[mergevar](#), [setvars](#)

makewind

Purpose

Creates a graphic panel of specific size and position and adds it to the list of graphic panels. Note: This function is for the deprecated PQG graphics. For similar functionality, see **plotLayout** and **plotCustomLayout**.

Library

pgraph

Format

```
makewind(xsize, ysize, xshft, yshft, typ);
```

Input

<i>xsize</i>	scalar, horizontal size of the graphic panel in inches.
<i>ysize</i>	scalar, vertical size of the graphic panel in inches.
<i>xshft</i>	scalar, horizontal distance from left edge of window in inches.
<i>yshft</i>	scalar, vertical distance from bottom edge of window in inches.
<i>typ</i>	scalar, graphic panel attribute type. If this value is 1, the graphic panels will be transparent. If 0, the graphic panels will be nontransparent.

Remarks

Note that if this procedure is used when rotating the page, the passed parameters are scaled appropriately to the newly oriented page. The size and shift values will not be true inches when printed, but the graphic panel size to page size ratio will remain the same. The result of this implementation automates the rotation and eliminates the required graphic panel recalculations by the user.

See the **window** command for creating tiled graphic panels. For more information on using graphic panels, see **Tiled Graphic Panels**, Section 1.0.1.

Source

pwindow.src

See Also

[window](#), [endwind](#), [setwind](#), [getwind](#), [begwind](#), [nextwind](#)

margin

margin

Purpose

Sets the margins for the current graph's graphic panel. Note: This function is for use with the deprecated PQG graphics. For similar functionality, use `plotCustomLayout`.

Library

pgraph

Format

```
margin(l, r, t, b);
```

Input

<i>l</i>	scalar, the left margin in inches.
<i>r</i>	scalar, the right margin in inches.
<i>t</i>	scalar, the top margin in inches.
<i>b</i>	scalar, the bottom margin in inches.

Remarks

By default, the dimensions of the graph are the same as the graphic panel dimensions. With this function the graph dimensions may be decreased. The result will be a smaller plot area surrounded by the specified margin. This procedure takes into consideration the axes labels and numbers for correct placement.

All input inch values for this procedure are based on a full size window of 9x6.855 inches. If this procedure is used with a graphic panel, the values will be scaled to "window inches" automatically.

If the axes must be placed an exact distance from the edge of the page, **axmargin** should be used.

Source

pgraph.src

See Also

[axmargin](#)

matalloc

Purpose

Allocates a matrix with unspecified contents.

Format

```
y = matalloc(r, c);
```

Input

<i>r</i>	scalar, rows.
<i>c</i>	scalar, columns.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix.
----------	-----------------------------

Remarks

The contents are unspecified. This function is used to allocate a matrix that will be

matinit

written to in sections using indexing or used with the Foreign Language Interface as an output matrix for a function called with `dllcall`.

See Also

[matinit](#), [ones](#), [zeros](#), [eye](#)

matinit

Purpose

Allocates a matrix with a specified fill value.

Format

```
y=matinit(r, c, v);
```

Input

<i>r</i>	scalar, rows.
<i>c</i>	scalar, columns.
<i>v</i>	scalar, value to initialize.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix with each element equal to the value of <i>v</i> .
----------	---

Example

```
format /rd 6,2;  
print matinit(3, 4, pi);
```

```
3.14  3.14  3.14  3.14
3.14  3.14  3.14  3.14
3.14  3.14  3.14  3.14
```

See Also

[matalloc](#), [ones](#), [zeros](#), [eye](#)

mattoarray

Purpose

Converts a matrix to a type array.

Format

```
y = mattoarray(x);
```

Input

x	matrix.
---	---------

Output

y	1-or-2-dimensional array.
---	---------------------------

Remarks

If the argument `x` is a scalar, **mattoarray** will simply return the scalar, without changing it to a type array.

Example

```
x = 5*ones(2,3);
```

maxc

```
y = mattoarray(x);
```

y will be a 2x3 array of fives.

See Also

[arraytomat](#)

maxc

Purpose

Returns a column vector containing the largest element in each column of a matrix.

Format

```
y = maxc(x);
```

Input

x	$N \times K$ matrix or sparse matrix.
-----	---------------------------------------

Output

y	$K \times 1$ matrix containing the largest element in each column of x .
-----	--

Remarks

If x is complex, **maxc** uses the complex modulus (**abs**(x)) to determine the largest elements.

To find the maximum elements in each row of a matrix, transpose the matrix before applying the **maxc** function.

To find the maximum value in the whole matrix if the matrix has more than one column, nest two calls to **maxc**:

```
y = maxc (maxc (x) ) ;
```

Example

```
x = randBeta (4,2,3,1) ;  
y = maxc (x) ;
```

If **x** equals:

```
0.87174453 0.70281291  
0.90393029 0.95919009  
0.82960656 0.58022236  
0.80910492 0.61975567
```

then **y** will equal:

```
0.90393029  
0.95919009
```

See Also

[minc](#), [maxindc](#), [minindc](#)

maxindc

Purpose

Returns a column vector containing the index (i.e., row number) of the maximum element in each column of a matrix.

maxindc

Format

```
y = maxindc(x);
```

Input

x NxK matrix.

Output

y Kx1 matrix containing the index of the maximum element in each column of x .

Remarks

If x is complex, **maxindc** uses the complex modulus (**abs**(x)) to determine the largest elements.

To find the index of the maximum element in each row of a matrix, transpose the matrix before applying **maxindc**.

To find the indices of the largest element in a matrix x , use:

```
colInd = maxindc(maxc(x));  
rowInd = maxindc(x[:,colInd]);
```

If there are two or more "largest" elements in a column (i.e., two or more elements equal to each other and greater than all other elements), then **maxindc** returns the index of the first one found, which will be the smallest index.

Example

```
x = round(rndn(4,4)*5);  
mx = maxc(x);
```

```
mxInd = maxindc(x);
```

If *x* is equal to:

-2	-8	-1	-2
-1	9	0	7
9	0	4	8
-2	6	6	1

then

	9		3
mx =	9	mxInd =	2
	6		4
	8		3

See Also

[maxc](#), [minindc](#), [minc](#)

maxv

Purpose

Performs an element by element comparison of two matrices and returns the maximum value for each element.

Format

```
z = maxv(x, y);
```

Global Input

<i>x</i>	NxK matrix
<i>y</i>	NxK matrix

maxv

Output

z A NxK matrix whose values are the maximum of each element from the arguments *x* and *y*.

Remarks

maxv works for sparse matrices as well as arrays.

Example

```
//Create the sequence 1, 2, 3,...10
x = sega(1, 1, 10);

//Set 'y' equal to the reverse order of 'x'
y = rev(x);

z = maxv(x,y);
```

	1	10	10
	2	9	9
	3	8	8
	4	7	7
x = 5	y = 6	z = 6	
	6	5	6
	7	4	7
	8	3	8
	9	2	9
	10	1	10

See Also

[minv](#)

maxvec

Purpose

Returns maximum vector length allowed.

Format

```
y = maxvec ;
```

Global Input

___maxvec scalar, maximum vector length allowed.

Output

y scalar, maximum vector length.

Remarks

maxvec returns the value in the global scalar ___maxvec, which can be reset in the calling program.

maxvec is called by **Run-Time Library** functions and applications when determining how many rows can be read from a data set in one call to **readr**.

Using a value that is too large can cause excessive disk thrashing. The trick is to allow the algorithm making the disk reads to execute entirely in RAM.

Example

```
y = maxvec;  
print y;
```

maxbytes

```
20000.000
```

Source

```
system.src
```

maxbytes

Purpose

Returns maximum memory to be used.

Format

```
y = maxbytes ;
```

Global Input

_____ scalar, maximum memory to be used.
maxbytes

Output

y scalar, maximum memory to be used.

Remarks

maxbytes returns the value in the global scalar `__maxbytes`, which can be reset in the calling program.

maxbytes is called by **Run-Time Library** functions and applications when determining how many rows can be read from a data set in one call to **readr**.

maxbytes replaced the obsolete command **coreleft**. If **coreleft** returns a meaningful number for your operating system and if you wish to reference it, set `__maxbytes = 0` and then call **maxbytes**.

Example

```
y = maxbytes;  
print y;
```

```
100000000.000
```

Source

system.src

mbesseli

Purpose

Computes modified and exponentially scaled modified Bessels of the first kind of the n th order.

Format

```
y = mbesseli(x, n, alpha);  
y = mbesseli0(x);  
y = mbesseli1(x);  
y = mbesselei(x, n, alpha);  
y = mbesselei0(x);  
y = mbesselei1(x);
```

Input

x	$K \times 1$ vector, abscissae.
-----	---------------------------------

mbesseli

n	scalar, highest order.
α	scalar, $0 \leq \alpha < 1$.

Output

y	$K \times N$ matrix, evaluations of the modified Bessel or the exponentially scaled modified Bessel of the first kind of the n th order.
-----	--

Remarks

For the functions that permit you to specify the order, the returned matrix contains a sequence of modified or exponentially scaled modified Bessel values of different orders. For the i th row of y :

$$y[i, :] = I_{\alpha}(x[i]) \ I_{\alpha+1}(x[i]) \ \dots I_{\alpha+n-1}(x[i])$$

The remaining functions generate modified Bessels of only the specified order.

The exponentially scaled modified Bessels are related to the unscaled modified Bessels in the following way:

$$\text{mbesselei0}(x) = \exp(-x) * \text{mbesseli0}(x)$$

The use of the scaled versions of the modified Bessel can improve the numerical properties of some calculations by keeping the intermediate numbers small in size.

Example

This example produces estimates for the "circular" response regression model (Fisher, N.I. *Statistical Analysis of Circular Data*. NY: Cambridge University Press, 1993.), where the dependent variable varies between $-\pi$ and π in a circular manner. The model is

$$y = \mu + G(XB)$$

where **B** is a vector of regression coefficients, **x** a matrix of independent variables with a column of 1's included for a constant, and **y** a vector of "circular" dependent variables, and where $G()$ is a function mapping XB onto the $[-\pi, \pi]$ interval.

The log-likelihood for this model is from Fisher, N.I. ... 1993, 159:

$$\log L = -N \times \ln(I_0(\kappa)) + \kappa \sum_i^N \cos(y_i - \mu - G(X_i B))$$

To generate estimates it is necessary to maximize this function using an iterative method. **QNewton** is used here.

κ is required to be nonnegative and therefore in the example below, the exponential of this parameter is estimated instead. Also, the exponentially scaled modified Bessel is used to improve numerical properties of the calculations.

The **arctan** function is used in $G()$ to map XB to the $[-\pi, \pi]$ interval as suggested by Fisher, N.I. ... 1993, 158.

```
proc G(u);
    ret(2*atan(u));
endp;

proc lpr(b);
    local dev;
    /*
    ** b[1] - kappa
    ** b[2] - mu
    ** b[3] - constant
    ** b[4:rows(b)] - coefficients
    */
    dev = y - b[2] - G(b[3] + x * b[4:rows(b)]);
```

meanc

```
retp (rows (dev) * ln (mbesselei0 (exp (b[1])) -
      sumc (exp (b[1]) * (cos (dev) - 1)))));
endp;

loadm data;
y0 = data[:,1];
x0 = data[:,2:cols (data)];

b0 = 2*ones (cols (x0), 1);

{ b,fct,grd,ret } = QNewton (&lpr,b0);

cov = invpd (hessp (&lpr,b));

print "estimates standard errors";
print;
print b~sqrt (diag (cov));
```

Source

ribes1.src

meanc

Purpose

Computes the mean of every column of a matrix.

Format

$y = \text{meanc}(x);$

Input

x	$N \times K$ matrix.
-----	----------------------

Output

 y

Kx1 matrix containing the mean of every column of x .

Example

```
x = mean(randu(1e5,4));
```

After the code above, x is equal to:

```
0.5007  
0.5004  
0.4995  
0.5016
```

In this example, 4 columns of uniform random numbers are generated in a matrix, and the mean is computed for each column. Due to the use of random input data in this example, your results may differ slightly.

See Also

[stdc](#)

median

Purpose

Computes the medians of the columns of a matrix.

Format

```
m = median(x);
```

median

Input

x NxK matrix.

Output

m Kx1 vector containing the medians of the respective columns of x .

Remarks

median will return a missing value for any column that contains a missing value.

Example

```
//Set the seed for repeatable random data
rndseed 4320993;

//Create uniform random integers between 1 and 10
x = ceil(10*randu(100,3));

//Calculate the median of each column of 'x'
md = median(x);
```

After the code above, *md* is equal to:

```
5.0000
5.0000
6.0000
```

Source

median.src

mergeby

Purpose

Merges two sorted files by a common variable.

Format

```
mergeby(infile1, infile2, outfile, keytyp);
```

Input

<i>infile1</i>	string, name of input file 1.
<i>infile2</i>	string, name of input file 2.
<i>outfile</i>	string, name of output file.
<i>keytyp</i>	scalar, data type of key variable.
	1 numeric
	2 character

Remarks

This will combine the variables in the two files to create a single large file. The following assumptions hold:

1. Both files have a single (key) variable in common and it is the first variable.
2. All of the values of the key variable are unique.
3. Each file is already sorted on the key variable.

The output file will contain the key variable in its first column.

It is not necessary for the two files to have the same number of rows. For each row for which the key variables match, a row will be created in the output file. *outfile* will

mergevar

contain the columns from *infile1* followed by the columns from *infile2* minus the key column from the second file.

If the inputs are null ("" or 0), the procedure will ask for them.

Source

sortd.src

mergevar

Purpose

Accepts a list of names of global matrices, and concatenates the corresponding matrices horizontally to form a single matrix.

Format

```
x = mergevar(vnames);
```

Input

<i>vnames</i>	string or Kx1 column vector containing the names of K global matrices.
---------------	--

Output

<i>x</i>	NxM matrix that contains the concatenated matrices, where M is the sum of the columns in the K matrices specified in <i>vnames</i> .
----------	--

Remarks

The matrices specified in *vnames* must be globals and they must all have the same

number of rows.

This function is the opposite of **makevars**.

Example

```
//Random integers between 1 and 72
age = ceil(72 * rndu(100, 1));

//Random normal numbers with a mean of 70 and a standard
//deviation of 10
income = 10 * rndn(100, 1) + 70;

//Vertically concatenate the strings
vnames = "age"$|"income";

//Merge the variables into 1 matrix
agInc = mergevar(vnames);
```

The column vectors *age* and *income* will be concatenated horizontally to create *agInc*. The above call to **mergevar** is equivalent to:

```
//Combine the matrices using the horizontal concatenation
//operator
agInc = age~income;
```

Source

vars.src

See Also

[makevars](#)

minc

minc

Purpose

Returns a column vector containing the smallest element in each column of a matrix.

Format

```
y = minc(x);
```

Input

x	$N \times K$ matrix or sparse matrix.
-----	---------------------------------------

Output

y	$K \times 1$ matrix containing the smallest element in each column of x .
-----	---

Remarks

If x is complex, **minc** uses the complex modulus (**abs**(x)) to determine the smallest elements.

To find the minimum element in each row, transpose the matrix before applying the **minc** function.

To find the minimum value in the whole matrix, nest two calls to **minc**:

```
y = minc(minc(x));
```

Example

```
x = randn(4,2);
```



```
y = minc(x);
```

If x is equal to:

```
-1.9950  -1.3477  
-0.4031  -1.9137  
 0.8136  -2.3155  
-0.9947   1.4061
```

then y will equal:

```
-1.9950  
-2.3155
```

See Also

[maxc](#), [minindc](#), [maxindc](#)

minindc

Purpose

Returns a column vector containing the index (i.e., row number) of the smallest element in each column of a matrix.

Format

```
y = minindc(x);
```

Input

x NxK matrix.

minindc

Output

y

Kx1 matrix containing the index of the smallest element in each column of x .

Remarks

If x is complex, **minindc** uses the complex modulus (**abs**(x)) to determine the smallest elements.

To find the index of the smallest element in each row, transpose the matrix before applying **minindc**.

To find the index of the smallest element in a matrix x , use:

```
colInd = minindc(minc(x));  
rowInd = minindc(x[:,colInd]);
```

If there are two or more "smallest" elements in a column (i.e., two or more elements equal to each other and less than all other elements), then **minindc** returns the index of the first one found, which will be the smallest index.

Example

```
x = round(rndn(5,4)*5);  
y = minc(x);  
z = minindc(x);
```

If x is equal to:

	-5	4	-4	0
	-2	3	4	3
$x =$	-11	5	5	5
	1	2	7	4
	-2	4	-1	-5

then y and z are equal to:

$$\begin{array}{rcl} & -11 & 3 \\ y = & 2 & z = 4 \\ & -4 & 1 \\ & -5 & 5 \end{array}$$

See Also

[maxindc](#), [minc](#), [maxc](#)

minv

Purpose

Performs an element by element comparison of two matrices and returns the minimum value for each element.

Format

```
z = minv(x, y);
```

Global Input

x	NxK matrix
y	NxK matrix

Output

z	A NxK matrix whose values are the minimum of each element from the arguments x and y .
-----	--

miss, missrv

Remarks

minv works for sparse matrices as well as arrays.

Example

Σ

```
//Create the multiplicative sequence 1, 2, 4, 8
x = seqm(1,2,4);

//Reverse the order of the elements in 'x' and assign them
//to 'y'
y = rev(x);

z = minv(x,y);
```

After the code above:

1	8	1
x = 2	y = 4	z = 2
4	2	2
8	1	1

See Also

[maxv](#)

miss, missrv

Purpose

miss converts specified elements in a matrix to **GAUSS**'s missing value code.
missrv is the reverse of this, and converts missing values into specified values.

Format

```
y = miss(x, v);  
y = missrv(x, v);
```

Input

x	NxK matrix.
v	LxM matrix, ExE conformable with x.

Output

y	max(N,L) by max(K,M) matrix.
---	------------------------------

Remarks

For **miss**, elements in *x* that are equal to the corresponding elements in *v* will be replaced with the **GAUSS** missing value code.

For **missrv**, elements in *x* that are equal to the **GAUSS** missing value code will be replaced with the corresponding element of *v*.

For complex matrices, the missing value code is defined as a missing value entry in the real part of the matrix. For complex *x*, then, **miss** replaces elements with a ". + 0i" value, and **missrv** examines only the real part of *x* for missing values. If, for example, an element of *x* = 1 + .i, **missrv** will not replace it.

These functions act like element-by-element operators. If *v* is a scalar, for instance -1, then all -1's in *x* are converted to missing. If *v* is a row (column) vector with the same number of columns (rows) as *x*, then each column (row) in *x* is transformed to missings according to the corresponding element in *v*. If *v* is a matrix of the same size as *x*, then the transformation is done corresponding element by corresponding element.

miss, missrv

Missing values are given special treatment in the following functions and operators: b/A (matrix division when a is not square and neither a nor b is scalar), **counts**, **scalmiss**, **maxc**, **maxindc**, **minc**, **minindc**, **miss**, **missex**, **missrv**, **moment**, **packr**, **scalmiss**, **sortc**.

As long as you know a matrix contains no missings to begin with, **miss** and **missrv** can be used to convert one set of numbers into another. For example:

```
y = missrv(miss(x,0),1);
```

will convert 0's to 1's.

To convert a range of values, such as:

```
0.5 < x < 1.3
```

into missing values, use the **missex** function.

Example

```
//Create a 3x3 matrix with each element equal to 1
x = ones(3, 3);

//Assign the diagonal of 'x' to be equal to pi
x = diagrv(x, pi);

print "x = " x;

//Change all 1's in 'x' into missing values and assign to
//xmiss
xmiss = miss(x, 1);

print "xmiss = " xmiss;

//Change all missings in 'xmiss' into 2*pi and assign to x2
```

```
x2 = misstrv(xmiss, 2*pi);

print "x2 = " x2;
```

The code above, will return:

```
x =
    3.1415927    1.0000000    1.0000000
    1.0000000    3.1415927    1.0000000
    1.0000000    1.0000000    3.1415927

xmiss =
    3.1415927         .         .
         .    3.1415927         .
         .         .    3.1415927

x2 =
    3.1415927    6.2831853    6.2831853
    6.2831853    3.1415927    6.2831853
    6.2831853    6.2831853    3.1415927
```

See Also

[counts](#), [ismiss](#), [maxc](#), [maxindc](#), [minc](#), [minindc](#), [missex](#), [moment](#), [packr](#), [scalmiss](#), [sortc](#)

missex

Purpose

Converts numeric values to the missing value code according to the values given in a logical expression.

Format

```
y = missex(x, mask);
```

missex

Input

x	NxK matrix.
$mask$	NxK logical matrix (matrix of 0's and 1's) that serves as a "mask" for x ; the 1's in $mask$ correspond to the values in x that are to be converted into missing values.

Output

y	NxK matrix that equals x , but with those elements that correspond to the 1's in e converted to missing.
-----	--

Remarks

The matrix e will usually be created by a logical expression. For instance, to convert all numbers between 10 and 15 in x to missing, the following code could be used:

```
 $y$  = missex( $x$ , ( $x$  .> 10) .and ( $x$  .< 15));
```

Note that "dot" operators MUST be used in constructing the logical expressions.

For complex matrices, the missing value code is defined as a missing value entry in the real part of the matrix. For complex x , then, **missex** replaces elements with a ". + 0i" value.

This function is like **miss**, but is more general in that a range of values can be converted into missings.

Example

```
//Set seed for repeatable random numbers
rndseed 49728424;

 $x$  = rndu(3,2);
```



```
//Logical expression
mask =(x .> .30) .and (x .< .60);
y = missex(x,mask);
```

After the code above:

x =	0.525	0.419		1	1		.	.
	0.869	0.973	mask =	0	0	y =	0.869	0.973
	0.021	0.357		0	1		0.021	.

A 3x2 matrix of uniform random numbers is created. All values in the interval (0.30, 0.60) are converted to missing.

Source

datatran.src

See Also

[miss](#), [missrv](#)

moment

Purpose

Computes a cross-product matrix. This is the same as $x'x$.

Format

$y = \text{moment}(x, d);$

Input

x	NxK matrix or M-dimensional array where the last two dimensions are NxK.
-----	--

moment

d	scalar, controls handling of missing values.
0	missing values will not be checked for. This is the fastest option.
1	"listwise deletion" is used. Any row that contains a missing value in any of its elements is excluded from the computation of the moment matrix. If every row in x contains missing values, then moment ($x, 1$) will return a scalar zero.
2	"pairwise deletion" is used. Any element of x that is missing is excluded from the computation of the moment matrix. Note that this is seldom a satisfactory method of handling missing values, and special care must be taken in computing the relevant number of observations and degrees of freedom.

Output

y	$K \times K$ matrix or M -dimensional array where the last two dimensions are $K \times K$, the cross-product of x .
-----	---

Remarks

The fact that the moment matrix is symmetric is taken into account to cut execution time almost in half.

If x is an array, the result will be an array containing the cross-products of each 2-dimensional array described by the two trailing dimensions of x . In other words, for a

10x4x4 array x , the resulting array y will contain the cross-products of each of the 10 4x4 arrays contained in x , so $y[n,.,.] = x[n,.,.]'x[n,.,.]$ for $1 \leq n \leq 10$.

If there is no missing data then $d = 0$ should be used because it will be faster.

The `/` operator (matrix division) will automatically form a moment matrix (performing pairwise deletions if **trap 2** is set) and will compute the **ols** coefficients of a regression. However, it can only be used for data sets that are small enough to fit into a single matrix. In addition, the moment matrix and its inverse cannot be recovered if the `/` operator is used.

Example

```
xx = moment(x, 2);
ixx = invpd(xx);
b = ixx*missrv(x, 0)'y;
```

In this example, the regression of y on x is computed. The moment matrix (xx) is formed using the **moment** command (with pairwise deletion, since the second parameter is 2). Then xx is inverted using the **invpd** function. Finally, the **ols** coefficients are computed. **missrv** is used to emulate pairwise deletion by setting missing values to 0.

momentd

Purpose

Computes a moment ($x'x$) matrix from a **GAUSS** data set.

Format

```
m = momentd(dataset, vars);
```

Input

⌘

<i>dataset</i>	string, name of data set.
<i>vars</i>	Kx1 string array, names of variables
	- or -
	Kx1 numeric vector, indices of columns.
	- or -
	Formula String , e.g. "PAY + WT" or ". - 1"(include all variables besides intercept)
	These can be any size subset of the variables in the data set, and can be in any order. If a scalar 0 is passed, all columns of the data set will be used.

Global Input

<i>__con</i>	scalar, default 1. 1 a constant term will be added. 0 no constant term will be added.
<i>__miss</i>	scalar, default 0. 0 there are no missing values (fastest). 1 do listwise deletion; drop an observation if any missings occur in it. 2 do pairwise deletion; this is equivalent to setting missings to 0 when calculating <i>m</i> .
<i>__row</i>	scalar, the number of rows to read per iteration of the read loop, default 0. If 0, the number of rows will be calculated internally.

If you get an Insufficient memory error, or you want the rounding to be exactly the same between runs, you can set the number of rows to read before calling **momentd**.

Output

m

MxM matrix, where $M = K + \text{---} \text{con}$, the moment matrix constructed by calculating $X'X$ where X is the data, with or without a constant vector of ones.

Error handling is controlled by the low order bit of the trap flag.

trap 0 terminate with error message

trap 1 return scalar error code in *m*

33 too many missings

34 file not found

Examples

Example 1 Using indices of columns

```
fname = getGAUSShome() $+ "examples/freqdata.dat";

//Calculate statistics on variables in dataset: PAYand WT
//Specify the index of PAYand WT
vars = 2|4;
m = momentd(fname, vars);

print m;
```

After the above code,

momentd

400.00000	787.00000	587.98000
787.00000	1805.0000	1161.1400
587.98000	1161.1400	900.38540

Example 2 Using names of variables

```
fname = getGAUSShome() $+ "examples/freqdata.dat";  
//Calculate statistics on variables in dataset:  PAYand WT  
//Define the names string array of PAYand WT  
string vars = {"PAY", "WT"};  
m = momentd(fname, vars );  
print m;
```

After the above code,

400.00000	787.00000	587.98000
787.00000	1805.0000	1161.1400
587.98000	1161.1400	900.38540

Example 3 Using formula string

```
fname = getGAUSShome() $+ "examples/freqdata.dat";  
//Define the formula for PAY and WT, remove the intercept  
(use - 1 )  
formula_str = "-1 + PAY + WT";  
  
//Calculate statistics on variables in dataset:  PAYand WT  
m = momentd(fname, formula_str);  
print m;
```

After the above code,

1805.0000	1161.1400
1161.1400	900.38540

Remarks

- The supported data set types are [CSV](#), [XLS](#), [XLSX](#), [HDF5](#), [FMT](#), [DAT](#).

For HDF5 file, the *dataset* must include [file schema](#) and both file name and data set name must be provided, e.g. **momentd** ("h5://C:/gauss17/examples/testdata.h5/mydata", *vars*).

- Character vector is supported for backward comparability, but it have been deprecated.

See also

[Formula String](#)

Source

momentd.src

movingave

Purpose

Computes moving average of a series.

Format

```
y = movingave(x, d);
```

Input

<i>x</i>	NxK matrix.
<i>d</i>	scalar, order of moving average.

movingaveExpwgt

Output

y	$N \times K$ matrix, filtered series. The first $d-1$ rows of x are set to missing values.
-----	--

Remarks

movingave is essentially a smoothing time series filter. The moving average is performed by column and thus it treats the $N \times K$ matrix as K time series of length N .

See Also

[movingaveWgt](#), [movingaveExpwgt](#)

movingaveExpwgt

Purpose

Computes exponentially weighted moving average of a series.

Format

$y = \text{movingaveExpwgt}(x, d, p);$

Input

x	$N \times K$ matrix.
d	scalar, order of moving average.
p	scalar, smoothing coefficient where $0 > p > 1$.

Output

y	$N \times K$ matrix, filtered series. The first $d-1$ rows of x are
-----	---

set to missing values.

Remarks

movingaveExpwgt is smoothing time series filter using exponential weights. The moving average as performed by column and thus it treats the NxK matrix as K time series of length N.

See Also

[movingaveWgt](#), [movingave](#)

movingaveWgt

Purpose

Computes weighted moving average of a series

Format

$y = \text{movingaveWgt}(x, d, w);$

Input

x	NxK matrix.
d	scalar, order of moving average.
w	$d \times 1$ vector, weights.

Output

y	NxK matrix, filtered series. The first $d-1$ rows of x are set to missing values.
-----	---

msym

Remarks

movingaveWgt is essentially a smoothing time series filter with weights. The moving average as performed by column and thus it treats the NxK matrix as K time series of length N.

See Also

[movingave](#), [movingaveExpwgt](#)

msym

Purpose

Allows the user to set the symbol that **GAUSS** uses when missing values are converted to ASCII and vice versa.

Format

```
msym str;
```

Input

str

literal or ^string (up to 8 letters) which, if not surrounded by quotes, is forced to uppercase. This is the string to be printed for missing values. The default is '.'.

Remarks

The entire string will be printed out when converting to ASCII in [print](#) and **printfm** statements.

When converting ASCII to binary in `loadm` and `let` statements, only the first character is significant. In other words,

```
msym HAT;
```

will cause 'H' to be converted to missing on input.

This does not affect **writer**, which outputs data in binary format.

Note that `msym` is a keyword and not a variable being assigned to, so there is no equals sign between `msym` and the string that is being passed to it.

Example

In the example below, you first create simulated data. The data represents the scores that a group of students received on a particular test and also the time that they took. For your calculations, you only want to consider data from students that completed the test in less than 80 minutes.

The code below replaces the scores from students that took more than 80 minutes with missing values. It uses the `msym` keyword to change the visual representation used for missing values from a '.' to a 'T'. Though, note that the underlying elements are still missing values, not character or string elements.

```
//Set seed for repeatable random numbers
rndseed 543124;

//Random integers with a mean of 70 and range of 20 to
//represent time taken for test
testTime = ceil(30 * rndu(10, 1)) + 60;

//Random integers with a mean of 1000 and a standard
//deviation of 10
score = ceil(10 * rndn(10, 1)) + 1000;

//Maximum allowed time for test
```

msym

```
maxTime = 80;

//Create a mask for times greater than maxTime
mask = testTime .> maxTime;

//Set scores to be missing values if testTime is greater
//than maxTime
mScores = missex(score, mask);

//Set missing values to print as 'T' to represent that the
//score was invalid because the student took too much time
msym "T";

format /rd 4,0;
print mScores;
```

The code above will return:

```
T
1010
997
1002
985
997
1007
995
T
T
```

See Also

[print](#), [printfm](#)

n

new

Purpose

Erases everything in memory including the symbol table; closes all open files as well as the auxiliary output and turns the window on if it was off; also allows the size of the new symbol table and the main program space to be specified.

Format

```
new;  
new nos;
```

Input

<i>nos</i>	scalar, optional input which indicates the maximum number of global symbols allowed.
------------	--

Remarks

Procedures, user-defined functions, and global matrices, strings, and string arrays are all global symbols.

If you would like your user-defined procedures to not be cleared after a `new` statement, you can either add them to a **GAUSS Library** or create a file in your GAUSSHOME directory with the same name as your procedure and a `.g` file extension. This file `.g` file should only contain your procedure.

nextindex

This command can be used with arguments as the first statement in a program to clear the symbol table and to allocate only as much space for program code as your program actually needs. When used in this manner, the auxiliary output will not be closed. This will allow you to open the auxiliary output from the command level and run a program without having to remove the `new` at the beginning of the program. If this command is not the first statement in your program, it will cause the program to terminate.

Example

```
new; /* clear global symbols. */

new 300; /* clear global symbols, set maximum
        ** number of global symbols to 300,
        ** and leave program space unchanged.
        */
```

See Also

[clear](#), [delete](#), [output](#)

nextindex

Purpose

Returns the index of the next element or subarray in an array.

Format

```
ni = nextindex(i, o);
```

Input

<i>i</i>	Mx1 vector of indices into an array, where $M \leq N$.
<i>o</i>	Nx1 vector of orders of an N-dimensional array.

Output

ni

Mx1 vector of indices, the index of the next element or subarray in the array corresponding to *o*.

Remarks

nextindex will return a scalar error code if the index cannot be incremented.

Example

```
//Dimensions of an array
orders = { 3, 4, 5, 6, 7 };

//Starting index
ind = { 2, 3, 5 };

//Return the index for the next element
ind = nextindex(ind,orders);
```

After the code above, *ind* will be equal to:

```
2
4
1
```

In this example, **nextindex** incremented *ind* to index the next 6x7 subarray in array *a*.

Using the same data from above, a subsequent call to **nextindex**:

```
ind = nextindex(ind,orders);
```

will assign *ind* to be equal to:

nextn, nextnevn

```
2
4
2
```

See Also

[previousindex](#), [loopnextindex](#), [walkindex](#)

nextn, nextnevn

Purpose

Returns allowable matrix dimensions for computing FFT's.

Format

```
n = nextn(n0);
n = nextnevn(n0);
```

Input

$n0$	scalar, the length of a vector or the number of rows or columns in a matrix.
------	--

Output

n	scalar, the next allowable size for the given dimension for computing an FFT or RFFT. $n > n0$.
-----	--

Remarks

nextn and **nextnevn** determine allowable matrix dimensions for computing FFT's.

The Temperton FFT routines (see table below) can handle any matrix whose dimensions can be expressed as:

$$2^p \times 3^q \times 5^r \times 7^s$$

where p , q and r are nonnegative integers and s is equal to 0 or 1.

with one restriction: the vector length or matrix column size must be even (p must be positive) when computing RFFT's.

fftn, etc., automatically pad matrices (with zeros) to the next allowable dimensions; **nextn** and **nextnevn** are provided in case you want to check or fix matrix sizes yourself.

Use the following table to determine what to call for a given function and matrix:

FFT	Vector	Matrix	Matrix
Function	Length	Rows	Columns
fftn	nextn	nextn	nextn
rfftn	nextnevn	nextn	nextnevn
rfftnp	nextnevn	nextn	nextnevn

Example

```
n = nextn(456);
```

The code above will assign n to be equal to 480.

Source

optim.src

See Also

[fftn](#), [optn](#), [optnevn](#), [rfftn](#), [rfftnp](#)

nextwind

nextwind

Purpose

Set the current graphic panel to the next available graphic panel. Note: This function is for use with the deprecated PQG graphics. For similar functionality use `plotLayout` instead.

Library

pgraph

Format

```
nextwind;
```

Remarks

This function selects the next available graphic panel to be the current graphic panel. This is the graphic panel in which the next graph will be drawn.

See the discussion on using graphic panels in **Tiled Graphic Panels**, Section 1.0.1.

Source

pwindow.src

See Also

[endwind](#), [begwind](#), [setwind](#), [getwind](#), [makewind](#), [window](#)

ntos

Purpose

Converts a floating point number to a string or string array with optionally specified precision.

Format

```
str = ntos(num);
str = ntos(num, prec);
```

Input

<i>num</i>	scalar or NxK matrix; the numbers to be converted to a string.
<i>prec</i>	Scalar, optional argument; the number of digits to display. If the precision input is not specified, the default value is 6. Valid input values are: $1 \leq prec \leq 15$.

Output

<i>str</i>	String or NxK string array containing the string representation of the input.
------------	---

Examples

Example 1

```
//Set 'pi_num' equal to the constant 'pi'
pi_num = pi;

//Create a string containing the first 6 digits of pi
pi_str = ntos(pi_num);
print pi_str;
```

returns:

```
3.14159
```

Example 2

ntos

```
roi = 6.725301;

//Convert to string with 3 digits
roi_str = ntos(roi, 3);

//Combine strings
out = "The project had an ROI of " $+ roi_str $+ "%";
print out;
```

returns:

```
The project had an ROI of 6.73%
```

Example 3

```
parm = { 1982 2.75000,
         1983 2.20272,
         1984 2.55102 };

//Convert to string array with max of 5 digits per element
parms = ntos(parm, 5);

print parms;
```

returns:

1982	2.75
1983	2.2027
1984	2.551

Remarks

This function will convert numbers to either decimal representation or scientific notation, depending upon which is most compact. The behavior is equivalent to the '%' format specifier to the 'C' language function **printf**. The precision of an individual number will be the smaller of the *prec* input and the maximum number of significant digits.

See Also

[ftos](#), [stof](#)

null

Purpose

Computes an orthonormal basis for the (right) null space of a matrix.

Format

`b = null(x);`

Input

`x` NxM matrix.

Output

`b` MxK matrix, where K is the nullity of `x`, such that:

`x * b = 0 //NxK matrix of 0's`

and

`b'b = I //MxM identity matrix`

The error returns are returned in `b`:

error code	reason
1	there is no null space
2	<code>b</code> is too large to return in a single matrix

null

Use **scalerr** to test for error returns.

Remarks

The orthogonal complement of the column space of x' is computed using the QR decomposition. This provides an orthonormal basis for the null space of x .

Example

```
let x[2,4] = 2 1 3 -1
            3 5 1  2;
```

```
b = null(x);
z = x*b;
i = b'b;
```

After the code above:

```
      -0.804  0.142
b =  0.331 -0.473  z = 0  0  i = 1  0
      0.473  0.331      0  0      0  1
      0.142  0.804
```

Source

null.src

Globals

_qrdc, _grsl

null1

Purpose

Computes an orthonormal basis for the (right) null space of a matrix and writes it to a **GAUSS** dataset.

Format

```
nu = null1(x, dataset);
```

Input

<i>x</i>	NxM matrix.
<i>dataset</i>	string, the name of a data set null1 will write.

Output

<i>nu</i>	scalar, the nullity of <i>x</i> .
-----------	-----------------------------------

Remarks

null1 computes an MxK matrix *b*, where K is the nullity of *x*, such that:

```
x * b = 0 //NxK matrix of 0's
```

and

```
b'b = I //MxM identity matrix
```

The transpose of *b* is written to the data set named by *dataset*, unless the nullity of *x* is zero. If *nu* is zero, the data set is not written.

numCombinations

Source

null.src

Globals

_qrdc, _qrs1

numCombinations

Purpose

Computes number of combinations of n things taken k at a time.

Format

$y = \text{numCombinations}(n, k);$

Input

n	scalar.
k	scalar.

Output

y	scalar, number of combinations of n things take k at a time.
-----	--

Remarks

To calculate all of the combinations, use the function **combine**.

Example

```
y = numCombinations(25,5);  
  
print y;
```

The code above, returns:

```
53130.0000
```

See Also

[combine](#), [combined](#)

o

ols

Purpose

Computes a least squares regression.

Format

```
{ vnam, m, b, stb, vc, stderr, sigma, cx, rsq, resid, dwstat } = ols  
(dataset, depvar, indvars);  
{ vnam, m, b, stb, vc, stderr, sigma, cx, rsq, resid, dwstat } = ols  
(dataset, formula);
```

Input

<i>dataset</i>	string, name of data set or null string. If <i>dataset</i> is a null string, the procedure assumes that the actual data has been passed in the next two arguments.
<i>depvar</i>	If <i>dataset</i> contains a string: string, name of dependent variable - or - scalar, index of dependent variable. If scalar 0, the last column of the data set will be used. If <i>dataset</i> is a null string or 0: Nx1 vector, the dependent variable.
<i>indvars</i>	If <i>dataset</i> contains a string:

Kx1 character vector, names of independent variables

- or -

Kx1 numeric vector, indices of independent variables.

These can be any size subset of the variables in the data set and can be in any order. If a scalar 0 is passed, all columns of the data set will be used except for the one used for the dependent variable.

If *dataset* is a null string or 0:

NxK matrix, the independent variables.

formula

String, [formula string](#) of the model.

E.g. "*y* ~ *X1* + *X2*", '*y*' is the name of dependent variable, '*X1*' and '*X2*' are names of independent variables;

E.g. "*y* ~ ." , '.' means including all variables except dependent variable '*y*';

E.g. "*y* ~ -1 + *X1* + *X2*", '-1' means no intercept model.

Global Input

Defaults are provided for the following global input variables, so they can be ignored unless you need control over the other options provided by this procedure.

`__altnam`

character vector, default 0.

This can be a (K+1)x1 or (K+2)x1 character vector of alternate variable names for the output. If `__con` is 1, this must be (K+2)x1. The name of the dependent variable is the last element.

___ <i>con</i>	scalar, default 1.
	1 a constant term will be added, $D = K+1$.
	0 no constant term will be added, $D = K$.
	A constant term will always be used in constructing the moment matrix m .
___ <i>miss</i>	scalar, default 0.
	0 there are no missing values (fastest).
	1 listwise deletion, drop any cases in which missings occur.
	2 pairwise deletion, this is equivalent to setting missings to 0 when calculating m . The number of cases computed is equal to the total number of cases in the data set.
___ <i>olsalg</i>	string, default "cholup." Selects the algorithm used for computing the parameter estimates. The default Cholesky update method is more computationally efficient; however, accuracy can suffer for poorly conditioned data. For higher accuracy, set ___ <i>olsalg</i> to either <i>qr</i> or <i>svd</i> .
	<i>qr</i> Solves for the parameter estimates using a <i>qr</i> decomposition.
	<i>svd</i> Solves for the parameter estimates using a singular value decomposition.
___ <i>output</i>	scalar, default 1.
	1 print the statistics.
	0 do not print statistics.
___ <i>row</i>	scalar, the number of rows to read per iteration of the read loop. Default 0.

If 0, the number of rows will be calculated internally. If you get an Insufficient memory error while executing `ols`, you can supply a value for `__row` that works on your system.

The answers may vary slightly due to rounding error differences when a different number of rows is read per iteration. You can use `__row` to control this if you want to get exactly the same rounding effects between several runs.

<code>__olsres</code>	scalar, default 0.
1	compute residuals (<i>resid</i>) and Durbin-Watson statistic (<i>dwstat</i>).
0	<i>resid</i> = 0, <i>dwstat</i> = 0.

Output

<code>vnam</code>	(K+2)x1 or (K+1)x1 character vector, the variable names used in the regression. If a constant term is used, this vector will be (K+2)x1, and the first name will be "CONSTANT". The last name will be the name of the dependent variable.		
<code>m</code>	MxM matrix, where M = K+2, the moment matrix constructed by calculating $x'x$ where x is a matrix containing all useable observations and having columns in the order:		
	1.0 (constant)	indvars (independent variables)	depvar (dependent variable)
	A constant term is always used in computing <code>m</code> .		
<code>b</code>	Dx1 vector, the least squares estimates of parameters Error handling is controlled by the low order bit of the trap flag.		

trap 0	terminate with error message
trap 1	return scalar error code in <i>b</i>
30	system singular
31	system underdetermined
32	same number of columns as rows
33	too many missings
34	file not found
35	no variance in an independent variable

The system can become underdetermined if you use listwise deletion and have missing values. In that case, it is possible to skip so many cases that there are fewer useable rows than columns in the data set.

<i>stb</i>	Kx1 vector, the standardized coefficients.
<i>vc</i>	DxD matrix, the variance-covariance matrix of estimates.
<i>stderr</i>	Dx1 vector, the standard errors of the estimated parameters.
<i>sigma</i>	scalar, standard deviation of residual.
<i>cx</i>	(K+1)x(K+1) matrix, correlation matrix of variables with the dependent variable as the last column.
<i>rsq</i>	scalar, R square, coefficient of determination.
<i>resid</i>	residuals, $resid = y - x * b$.

If *_olsres* = 1, the residuals will be computed.

If the data is taken from a data set, a new data set will be created for the residuals, using the name in the global string

variable `_olsrnam`. The residuals will be saved in this data set as an Nx1 column. The `resid` return value will be a string containing the name of the new data set containing the residuals.

If the data is passed in as a matrix, the `resid` return value will be the Nx1 vector of residuals.

`dwstat`

scalar, Durbin-Watson statistic.

Remarks

- For poorly conditioned data the default setting for `__olsalg`, using the Cholesky update, may produce only four or five digits of accuracy for the parameter estimates and standard error. For greater accuracy, use either the `qr` or singular value decomposition algorithm by setting `__olsalg` to `qr` or `svd`. If you are unsure of the condition of your data, set `__olsalg` to `qr`.
- No output file is modified, opened, or closed by this procedure. If you want output to be placed in a file, you need to open an output file before calling `ols`.
- The supported data set types are [CSV](#), [XLS](#), [XLSX](#), [HDF5](#), [FMT](#), [DAT](#).
- For HDF5 file, the `dataset` must include [file schema](#) and both file name and data set name must be provided, e.g. `ols ("h5://C:/gauss17/examples/testdata.h5/mydata", formula)`.

Examples

Example 1

```
y = { 2,
      3,
      1,
      7,
      5 };

x = { 1 3 2,
```

```
2 3 1,
7 1 7,
5 3 1,
3 5 5 };
```

```
output file = ols.out reset;
call ols(0,y,x);
output off;
```

In this example, the output from **ols** is put into a file called `ols.out` as well as being printed to the window. This example will compute a least squares regression of *y* on *x*. The return values are discarded by using a `call` statement.

```
data = "olsdat";
depvar = { score };
indvars = { region, age, marstat };
_olsres = 1;
output file = lpt1 on;
{ nam, m, b, stb, vc, std, sig, cx, rsq, resid, dbw } = ols
(data, depvar, indvars);
output off;
```

In this example, the data set `olsdat.dat` is used to compute a regression. The dependent variable is *score*. The independent variables are: *region*, *age*, and *marstat*. The residuals and Durbin-Watson statistic will be computed. The output will be sent to the printer as well as the window and the returned values are assigned to variables.

Example 2 Pass in a data set name and variable names

```
fname = getGAUSShome() $+ "examples/credit.dat";
// Specify the formula, Limit is dependent variable and Bal-
ance, Income and Age are independent variables
dep = "Limit";
string indep = {"Balance", "Income", "Age"};
```



```
call ols(fname, dep, indep);
```

After the above code,

Valid cases:	400	Dependent variable:
Limit		
Missing cases:	0	Deletion method:
None		
Total SS:	2125784986.000	Degrees of freedom:
396		
R-squared:	0.939	Rbar-squared:
0.939		
Residual SS:	129727134.947	Std error of est:
572.358		
F(3,396):	2031.029	Probability of F:
0.000		

Standard Variable Estimate	Estimate Dep Var	Prob Error	Standardized t-value	Cor with > t
-----	-----	-----	-----	-----
CONSTANT	1521.904666	102.228802	14.887240	0.000
---	---			
Balance	3.168467	0.070635	44.856923	0.000
0.631111	0.861697			
Income	32.566995	0.935925	34.796581	0.000
0.497271	0.792088			
Age	1.677855	1.694288	0.990301	0.323
0.012539	0.100888			

Example 3 Pass in a data set name and a [formula string](#)

Command Reference

```
fname = getGAUSShome() $+ "examples/credit.dat";

// Specify the formula, 'Limit' is dependent variable and
// 'Balance', 'Income' and 'Age' are independent variables, '-
// 1' means remove the intercept in the model
formula = "Limit ~ - 1 + Balance + Income + Age ";

call ols(fname, formula);
```

After the above code,

Valid cases:	400	Dependent variable:
Limit		
Missing cases:	0	Deletion method:
None		
Total SS:	11096147930.000	Degrees of freedom:
397		
R-squared:	0.982	Rbar-squared:
0.982		
Residual SS:	202331711.222	Std error of est:
713.899		
F(3,397):	7125.008	Probability of F:
0.000		

Standard Variable Estimate	Estimate Dep Var	Prob Error	Standardized t-value	Cor with > t
Balance	3.429796	0.085339	40.190438	0.000
0.451757	0.923618			
Income	33.447531	1.165041	28.709327	0.000
0.363912	0.922459			
Age	23.718127	1.027629	23.080436	0.000
0.262414	0.871984			

Source

`ols.src`

See Also

[olsqr](#), [Formula String](#)

olsmt

Purpose

Computes a least squares regression.

Format

```
oout = olsmt(oc0, dataset, depvar, indvars);  
oout = olsmt(oc0, dataset, formula);
```

Input

<i>oc0</i>	instance of an olsmtControl structure containing the following members:
<i>oc0.altnam</i>	character vector, default 0. This can be a (K+1)x1 or (K+2)x1 character vector of alternate variable names for the output. If <i>oc0.con</i> is 1, this must be (K+2)x1. The name of the dependent variable is the last element.
<i>oc0.con</i>	scalar, default 1. 1 a constant term will be added, $D = K+1$. 0 no constant term will be

oc0.miss

added, $D = K$.

A constant term will always be used in constructing the moment matrix m .

scalar, default 0.

- 0 there are no missing values (fastest).
- 1 listwise deletion, drop any cases in which missings occur.
- 2 pairwise deletion, this is equivalent to setting missings to 0 when calculating m . The number of cases computed is equal to the total number of cases in the data set.

oc0.row

scalar, the number of rows to read per iteration of the read loop. Default 0.

If 0, the number of rows will be calculated internally. If you get an Insufficient memory error message while executing **olsmt**, you can supply a value for *oc0.row* that works on your system.

The answers may vary slightly due to rounding error differences when a different number of rows is read per iteration. You can use *oc0.row* to control this if you want to get exactly the same rounding effects between several runs.

<i>oc0.vpad</i>	<p>scalar, default 1.</p> <p>If 0, internally created variable names are not padded to the same length (e.g. "X1, X2,..., X10").</p> <p>If 1, they are padded with zeros to the same length (e.g., "X01, X02,..., X10").</p>
<i>oc0.output</i>	<p>scalar, default 1.</p> <p>1 print the statistics.</p> <p>0 do not print statistics.</p>
<i>oc0.res</i>	<p>scalar, default 0.</p> <p>1 compute residuals (<i>resid</i>) and Durbin-Watson statistic (<i>dwstat</i>.)</p> <p>0 <i>oout.resid</i> = 0, <i>oout.dwstat</i> = 0.</p>
<i>oc0.rnam</i>	<p>string, default "_olsmtres".</p> <p>If the data is taken from a data set, a new data set will be created for the residuals, using the name in <i>oc0.rnam</i>.</p>
<i>oc0.maxvec</i>	<p>scalar, default 20000.</p> <p>The largest number of elements allowed in any one matrix.</p>
<i>oc0.fcmtol</i>	<p>scalar, default 1e-12.</p> <p>Tolerance used to fuzz the comparison operations to allow for round off error.</p>
<i>oc0.alg</i>	<p>string, default "cholup".</p>

	<p>Selects the algorithm used for computing the parameter estimates. The default Cholesky update method is more computationally efficient. However, accuracy can suffer for poorly conditioned data. For higher accuracy set <i>oc0.alg</i> to either <i>qr</i> or <i>svd</i>.</p> <p><i>qr</i> Solves for the parameter estimates using a <i>qr</i> decomposition.</p> <p><i>svd</i> Solves for the parameter estimates using a singular value decomposition.</p>
<i>dataset</i>	string, name of data set or null string.
<i>depvar</i>	<p>If <i>dataset</i> is a null string, the procedure assumes that the actual data has been passed in the next two arguments.</p> <p>If <i>dataset</i> contains a string:</p> <p>string, name of dependent variable</p> <p>- or -</p> <p>scalar, index of dependent variable. If scalar 0, the last column of the data set will be used.</p> <p>If <i>dataset</i> is a null string or 0:</p> <p>Nx1 vector, the dependent variable.</p>
<i>indvars</i>	<p>If <i>dataset</i> contains a string:</p> <p>Kx1 character vector, names of independent variables</p> <p>- or -</p> <p>Kx1 numeric vector, indices of independent variables.</p> <p>These can be any size subset of the variables in the data set and can be in any order. If a scalar 0 is passed, all columns of the data set will be used except for the one</p>

used for the dependent variable.

If `dataset` is a null string or 0:
 NxK matrix, the independent variables.

formula String, [formula string](#) of the model.

E.g. "**y** ~ **X1** + **X2**", 'y' is the name of dependent variable, 'X1' and 'X2' are names of independent variables;

E.g. "**y** ~ .", '.' means including all variables except dependent variable 'y';

E.g. "**y** ~ -1 + **X1** + **X2**", '-1' means no intercept model.

Output

oolt instance of an **olsmtOut** structure containing the following members:

oolt.vnam (K+2)x1 or (K+1)x1 character vector, the variable names used in the regression. If a constant term is used, this vector will be (K+2)x1, and the first name will be "CONSTANT". The last name will be the name of the dependent variable.

oolt.m MxM matrix, where M = K+2, the moment matrix constructed by calculating $X'X$ where X is a matrix containing all useable observations and having columns in the order:

1.0	indvars	depvar	
	(constant)	(independent variables)	(dependent variable)

A constant term is always used in computing *m*.

oolt.b Dx1 vector, the least squares estimates of

parameters

Error handling is controlled by the low order bit of the trap flag.

trap 0 terminate with error message

trap 1 return scalar error code in *b*

30 system singular

31 system underdetermined

32 same number of columns as rows

33 too many missings

34 file not found

35 no variance in an independent variable

The system can become underdetermined if you use listwise deletion and have missing values. In that case, it is possible to skip so many cases that there are fewer useable rows than columns in the data set.

oout.stb Kx1 vector, the standardized coefficients.

oout.vc DxD matrix, the variance-covariance matrix of estimates.

oout.stder Dx1 vector, the standard errors of the estimated parameters.

oout.sigma scalar, standard deviation of residual.

oout.cx (K+1)x(K+1) matrix, correlation matrix of

	variables with the dependent variable as the last column.
<code>oout.rsq</code>	scalar, R square, coefficient of determination.
<code>oout.resid</code>	residuals, $oout.resid = y - x * oout.b$. If <code>oc0.olsres = 1</code> , the residuals will be computed. If the data is taken from a data set, a new data set will be created for the residuals, using the name in <code>oc0.rnam</code> . The residuals will be saved in this data set as an Nx1 column. The <code>oout.resid</code> return value will be a string containing the name of the new data set containing the residuals. If the data is passed in as a matrix, the <code>oout.resid</code> return value will be the Nx1 vector of residuals.
<code>oout.dwstat</code>	scalar, Durbin-Watson statistic.

Remarks

- For poorly conditioned data the default setting for `oc0.alg`, using the Cholesky update, may produce only four or five digits of accuracy for the parameter estimates and standard error. For greater accuracy, use either the `qr` or singular value decomposition algorithm by setting `oc0.alg` to `qr` or `svd`. If you are unsure of the condition of your data, set `oc0.alg` to `qr`.
- No output file is modified, opened, or closed by this procedure. If you want output to be placed in a file, you need to open an output file before calling `olsmt`.
- The supported data set types are [CSV](#), [XLS](#), [XLSX](#), [HDF5](#), [FMT](#), [DAT](#).
- For HDF5 file, the `dataset` must include [file schema](#) and both file name and data set name must be provided, e.g. `olsmt("h5://testdata.h5/mydata", formula)`.

Examples

Example 1 Use data matrices

```
struct olsmtControl oc0;
struct olsmtOut oOut;
oc0 = olsmtControlCreate;

y = { 2,
      3,
      1,
      7,
      5 };

x = { 1 3 2,
      2 3 1,
      7 1 7,
      5 3 1,
      3 5 5 };

output file = olsmt.out reset;
oOut = olsmt(oc0,0,y,x);
output off;
```

In this example, the output from **olsmt** is put into a file called `olsmt.out` as well as being printed to the window. This example will compute a least squares regression of y on x .

Example 2 Use a data set and variable names

```
struct olsmtControl oc0;
struct olsmtOut oOut;
oc0 = olsmtControlCreate;

data = "credit.dat";
depvar = "Limit";
string indvars = { "Balance", "Income", "Age" };
```

```

oc0.res = 1;
output file = lpt1 on;
oOut = olsmt(oc0, data, depvar, indepvars);
output off;

```

In this example, the data set "**credit.dat**" is used to compute a regression. The dependent variable is *Limit*. The independent variables are: *Balance*, *Income*, and *Age*. The residuals and Durbin-Watson statistic will be computed. The output will be sent to the printer as well as the window and the returned values are assigned to variables.

Example 3 Use a data set and variable indices

```

struct olsmtControl oc0;
struct olsmtOut oOut;
oc0 = olsmtControlCreate;

data = "credit.dat";
depvar = 2;

indepvar = {11,1, 5};

oc0.res = 1;

oOut = olsmt(oc0, data, depvar, indepvar);

```

In this example, the data set "**credit.dat**" is used to compute a regression. The dependent variable is *Limit* which is the second column in the data set. The independent variables are: *Balance*, *Income*, and *Age*. They are the 11, 1 and 5 columns in the data set. The residuals and Durbin-Watson statistic will be computed. The output is :

Valid cases:	400	Dependent variable:
Limit		
Missing cases:	0	Deletion method:
None		
Total SS:	2125784986.000	Degrees of freedom:

```

              396
R-squared:          0.939      Rbar-squared:
              0.939
Residual SS:      129727134.947  Std error of est:
              572.358
F(3,396):          2031.029      Probability of F:
              0.000
Durbin-Watson:          1.953
```

Standard Variable Estimate	Estimate Dep Var	Prob Error	Standardized t-value	Cor with > t
-----	-----	-----	-----	-----
CONSTANT	1521.904666	102.228802	14.887240	0.000
---	---	---	---	---
Balance	3.168467	0.070635	44.856923	0.000
0.631111	0.861697			
Income	32.566995	0.935925	34.796581	0.000
0.497271	0.792088			
Age	1.677855	1.694288	0.990301	0.323
0.012539	0.100888			

Example 4 Use a data set and a [formula string](#)

```
struct olsmtControl oc0;
struct olsmtOut oOut;
oc0 = olsmtControlCreate;

data = "credit.dat";
formula = "Limit ~ Balance + Income + Age" };
oc0.res = 1;

oOut = olsmt(oc0, data, formula);
```

In this example, the data set "**credit.dat**" is used to compute a regression. The dependent variable is *Limit*. The independent variables are: *Balance*, *Income*, and *Age*. The residuals and Durbin-Watson statistic will be computed. The output is :

```
Valid cases:                400      Dependent variable:
                               Limit
Missing cases:              0      Deletion method:
                               None
Total SS:                   2125784986.000  Degrees of freedom:
                               396
R-squared:                  0.939      Rbar-squared:
                               0.939
Residual SS:                129727134.947  Std error of est:
                               572.358
F(3,396):                   2031.029      Probability of F:
                               0.000
Durbin-Watson:              1.953
```

Standard Variable Estimate	Estimate Dep Var	Prob Error	Standardized t-value	Cor with > t
-----	-----	-----	-----	-----
CONSTANT	1521.904666	102.228802	14.887240	0.000
---	---	---	---	---
Balance	3.168467	0.070635	44.856923	0.000
0.631111	0.861697			
Income	32.566995	0.935925	34.796581	0.000
0.497271	0.792088			
Age	1.677855	1.694288	0.990301	0.323
0.012539	0.100888			

Source

olsmt.src

See Also

[olsmtControlCreate](#), [olsqrmt](#), [Formula String](#)

olsmtControlCreate

Purpose

Creates default **olsmtControl** structure.

Include

`olsmt.sdf`

Format

```
c = olsmtControlCreate();
```

Output

c	instance of an olsmtControl structure with members set to default values.
-----	--

Example

Since structures are strongly typed in **GAUSS**, each structure must be declared before it can be used.

```
// declare 'ctl' as an olsmtControl structure
struct olsmtControl ctl;

// initialize structure 'ctl'
ctl = olsmtControlCreate;
```

The members of the **olsmtControl** structure and their default values are described in the manual entry for **olsmt**.

Source

`olsmt.src`

See Also

[olsmt](#)

olsqr

Purpose

Computes OLS coefficients using QR decomposition.

Format

```
b = olsqr(y, x);
```

Input

<i>y</i>	Nx1 vector containing dependent variable.
<i>x</i>	NxP matrix containing independent variables.

Output

<i>b</i>	Px1 vector of least squares estimates of regression of <i>y</i> on <i>x</i> . If <i>x</i> does not have full rank, then the coefficients that cannot be estimated will be zero.
----------	---

Remarks

This provides an alternative to y/x for computing least squares coefficients.

This procedure is slower than the $/$ operator. However, for near singular matrices it may produce better results.

olsqr handles matrices that do not have full rank by returning zeros for the coefficients that cannot be estimated.

Example

```
A = rndn(4,4);
b = rndn(4,1);
x = olsqr(b,A);
```

See Also

[ols](#), [olsqr2](#), [orth](#), [qqr](#)

olsqr2

Purpose

Computes OLS coefficients, residuals, and predicted values using the QR decomposition.

Format

```
{ b, r, p } = olsqr2(y, x);
```

Input

y	Nx1 vector containing dependent variable.
x	NxP matrix containing independent variables.

Output

b	Px1 vector of least squares estimates of regression of y on x . If x does not have full rank, then the coefficients that cannot be estimated will be zero.
r	Px1 vector of residuals. ($r = y - x * b$)
p	Px1 vector of predicted values. ($p = x * b$)

Remarks

This provides an alternative to y/x for computing least squares coefficients.

This procedure is slower than the $/$ operator. However, for near singular matrices, it may produce better results.

olsqr2 handles matrices that do not have full rank by returning zeros for the coefficients that cannot be estimated.

See Also

[olsqr](#), [orth](#), [qqr](#)

olsqrmt

Purpose

Computes OLS coefficients using QR decomposition.

Format

```
 $b = \text{olsqrmt}(y, x, \text{tol});$ 
```

Input

y	Nx1 vector containing dependent variable.
x	NxP matrix containing independent variables.
tol	scalar, the tolerance for testing if diagonal elements are approaching zero. The default value is 10^{-14} .

Output

b	Px1 vector of least squares estimates of regression of y on x . If x does not have full rank, then the coefficients that cannot be estimated will be zero.
-----	--

Remarks

This provides an alternative to y/x for computing least squares coefficients.

This procedure is slower than the $/$ operator. However, for near singular matrices it may produce better results.

olsqrmt handles matrices that do not have full rank by returning zeros for the coefficients that cannot be estimated.

Source

`olsmt.src`

See Also

[olsmt](#), [olsqr2](#)

ones

Purpose

Creates a matrix of ones.

Format

```
y = ones(r, c);
```

Input

<i>r</i>	scalar, number of rows.
<i>c</i>	scalar, number of columns.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of ones.
----------	-------------------------------------

Remarks

Noninteger arguments will be truncated to an integer.

Example

```
x = ones(3,2);
```

The code above assigns *x* to be equal to:

1.0000000	1.0000000
1.0000000	1.0000000
1.0000000	1.0000000

See Also

[zeros](#), [eye](#)

open

Purpose

Opens an existing GAUSS data file.

Format

```
open fh = filename;  
open fh = filename for mode;  
open fh = filename for mode varindxi offs;
```

Input

Command Reference	<i>filename</i>	literal or ^string. <i>filename</i> is the name of the file on the disk. The name can include a path if the directory to be used is not the current directory. This filename will automatically be given the extension .dat. If an extension is specified, the .dat will be overridden. If the file is an .fmt matrix file, the extension must be explicitly given. If the name of the file is to be taken from a string variable, the name of the string must be preceded by the ^ (caret) operator.
	<i>mode</i>	literal, the modes supported with the optional <code>for</code> subcommand are: read This is the default file opening mode and will be the one used if none is specified. Files opened in this mode cannot be written to. The pointer is set

	to the beginning of the file and the writer function is disabled for files opened in this way. This is the only mode available for matrix files (<code>.fmt</code>), which are always written in one piece with the <code>save</code> command.
append	Files opened in this mode cannot be read. The pointer will be set to the end of the file so that a subsequent write to the file with the writer function will add data to the end of the file without overwriting any of the existing data in the file. The readr function is disabled for files opened in this way. This mode is used to add additional rows to the end of a file.
update	Files opened in this mode can be read from and written to. The pointer will be set to the beginning of the file. This mode is used to make changes in a file.

offs scalar, offset added to "index variables."

The optional **varindx** subcommand tells **GAUSS** to create a set of global scalars that contain the index (column position) of the variables in a **GAUSS** data file. These "index variables" will have the same names as the corresponding variables in the data file but with "i" added as a prefix. They can be used inside index brackets, and with functions like **submat** to access specific columns of a matrix without having to remember the column position.

The optional *offs* argument is an offset that will be added to the index variables. This is useful if data from multiple files are concatenated horizontally in one matrix. It can be any scalar expression. The default is 0.

The index variables are useful for creating submatrices of specific variables without requiring that the positions of the variables be known. For instance, if there are two variables, *xvar* and *yvar* in the data set, the index variables will have the names *ixvar*, *iyvar*. If *xvar* is the first column in the data file, and *yvar* is the second, and if no offset, *offs*, has been specified, then *ixvar* and *iyvar* will equal 1 and 2 respectively. If an offset of 3 had been specified, then these variables would be assigned the values 4 and 5 respectively.

The **varindx** option cannot be used with *.fmt* matrix files because no column names are stored with them.

If **varindx** is used, **GAUSS** will ignore the Undefined symbol error for global symbols that start with "i". This makes it much more convenient to use index variables because they don't have to be cleared before they are accessed in the program. Clearing is otherwise necessary because the index variables do not exist until execution time when the data file is actually opened and the names are read in from the header of the file. At compile time a statement like: *y=x[.,ixvar];* will be illegal if the compiler has never heard of *ixvar*. If **varindx** is used, this error will be ignored for symbols beginning with "i". Any symbols that are accessed before they have been initialized with a real value will be trapped at execution time with a Variable not initialized error.

Output

<i>fh</i>	scalar, file handle. <i>fh</i> is the file handle which will be used by most commands to refer to the file within GAUSS . This file handle is actually a scalar containing an integer value that uniquely identifies each file. This value is assigned by GAUSS when the <code>open</code> command is executed. If the file was not successfully opened, the file handle will be set to -1.
-----------	--

Remarks

The file must exist before it can be opened with the `open` command. To create a new file, see `create` or `save`.

A file can be opened simultaneously under more than one handle. See the second example following.

If the value that is in the file handle when the `open` command begins to execute matches that of an already open file, the process will be aborted and a File already open message will be given. This gives you some protection against opening a second file with the same handle as a currently open file. If this happens, you would no longer be able to access the first file.

It is important to set unused file handles to zero because both `open` and `create` check the value that is in a file handle to see if it matches that of an open file before they proceed with the process of opening a file. This should be done with `close` or `closeall`.

Example

```
fname = "/data/rawdat";  
open dt = ^fname for append;
```

```

if dt == -1;
    print "File not found";
end;
endif;
y = writer(dt,x);
if y /= rows(x);
    print "Disk Full";
end;
endif;

dt = close(dt);

```

In the example above, the existing data set `/data/rawdat.dat` is opened for appending new data. The name of the file is in the string variable `fname`. In this example the file handle is tested to see if the file was opened successfully. The matrix `x` is written to this data set. The number of columns in `x` must be the same as the number of columns in the existing data set. The first row in `x` will be placed after the last row in the existing data set. The **writer** function will return the number of rows actually written. If this does not equal the number of rows that were attempted, then the disk is probably full.

```

open fin = mydata for read;
open fout = mydata for update;

do until eof(fin);
    x = readr(fin,100);
    x[.,1 3] = ln(x[.,1 3]);
    call writer(fout,x);
end;

closeall fin,fout;

```

In the above example, the same file, `mydata.dat`, is opened twice with two different file handles. It is opened for read with the handle `fin`, and it is opened for

update with the handle *fout*. This will allow the file to be transformed in place without taking up the extra space necessary for a separate output file. Notice that *fin* is used as the input handle and *fout* is used as the output handle. The loop will terminate as soon as the input handle has reached the end of the file. Inside the loop the file is read into a matrix called *x* using the input handle, the data are transformed (columns 1 and 3 are replaced with their natural logs), and the transformed data is written back out using the output handle. This type of operation works fine as long as the total number of rows and columns does not change.

The following example assumes a data file named `dat1.dat` that has the variables: *visc*, *temp*, *lub*, and *rpm*:

```
open f1 = dat1 varindxi;
dtx = readr(f1,100);
x = dtx[.,irpm ilub ivisc];
y = dtx[.,itemp];
call seekr(f1,1);
```

In this example, the data set `dat1.dat` is opened for reading (the `.dat` and the **for read** are implicit). **varindxi** is specified with no constant. Thus, index variables are created that give the positions of the variables in the data set. The first 100 rows of the data set are read into the matrix *dtx*. Then, specified variables in a specified order are assigned to the matrices *x* and *y* using the index variables. The last line uses the **seekr** function to reset the pointer to the beginning of the file.

```
open q1 = dat1 varindx;
open q2 = dat2 varindx colsf(q1);
nr = 100;
y = readr(q1,nr)~readr(q2,nr);
closeall q1,q2;
```

In this example, two data sets are opened for reading and index variables are created for each. A constant is added to the indices for the second data set (*q2*), equal to the number of variables (columns) in the first data set (*q1*). Thus, if there are three variables *x1*, *x2*, *x3* in *q1*, and three variables *y1*, *y2*, *y3* in *q2*, the index variables

that were created when the files were opened would be *ix1*, *ix2*, *ix3*, *iy1*, *iy2*, *iy3*. The values of these index variables would be 1, 2, 3, 4, 5, 6, respectively. The first 100 rows of the two data sets are read in and concatenated to produce the matrix *y*. The index variables will thus give the correct positions of the variables in *y*.

```
open fx = x.fmt;
rf = rowsf(fx);
sampsiz = round(rf*0.1);
rndsmpx = zeros(sampsiz,colsf(fx));

for(1, sampsiz, 1);
    r = ceil(rndu(1,1)*rf);
    call seekr(fx,r);
    rndsmpx[i,.] = readr(fx,1);
endfor;

fx = close(fx);
```

In this example, a 10% random sample of rows is drawn from the matrix file *x.fmt* and put into the matrix ***rndsmpx***. Note that the extension *.fmt* must be specified explicitly in the ***open*** statement. The ***rowsf*** command is used to obtain the number of rows in *x.fmt*. This number is multiplied by 0.10 and the result is rounded to the nearest integer; this yields the desired sample size. Then random integers (*r*) in the range 1 to *rf* are generated. ***seekr*** is used to locate to the appropriate row in the matrix, and the row is read with ***readr*** and placed in the matrix ***rndsmpx***. This is continued until the complete sample has been obtained.

See Also

[dataopen](#), [create](#), [close](#), [closeall](#), [readr](#), [writer](#), [seekr](#), [eof](#)

optn, optnevn

Purpose

Returns optimal matrix dimensions for computing FFT's.

Format

```
 $n = \text{optn}(n0);$ 
 $n = \text{optnevn}(n0);$ 
```

Input

$n0$	scalar, the length of a vector or the number of rows or columns in a matrix.
------	--

Output

n	scalar, the next optimal size for the given dimension for computing an FFT or RFFT. $n > n0$.
-----	--

Remarks

optn and **optnevn** determine optimal matrix dimensions for computing FFT's. The Temperton FFT routines (see table following) can handle any matrix whose dimensions can be expressed as:

$$2^p \times 3^q \times 5^r \times 7^s$$

where p , q and r are nonnegative integers and s is equal to 0 or 1.

with one restriction: the vector length or matrix column size must be even (p must be positive) when computing RFFT's.

fftn, etc., pad matrices to the next allowable dimensions; however, they generally run faster for matrices whose dimensions are highly composite numbers, that is, products of several factors (to various powers), rather than powers of a single factor. For example, even though it is bigger, a 33600x1 vector can compute as much as 20% faster than a 32768x1 vector, because 33600 is a highly composite number, $2^6 * 3 * 5^2 * 7$, whereas 32768 is a simple power of 2, 2^{15} . **optn** and **optnevn** are provided so

you can take advantage of this fact by hand-sizing matrices to optimal dimensions before computing the FFT.

Use the following table to determine what to call for a given function and matrix:

FFT	Vector	Matrix	Matrix
Function	Length	Rows	Columns
fftn	optn	optn	optn
rfftn	optnevn	optn	optnevn
rfftnp	optnevn	optn	optnevn

Example

```
n = optn(231);
```

The above code assigns *n* to be equal to 240.

See Also

[fftn](#), [nextn](#), [nextnevn](#), [rfftn](#), [rfftnp](#)

orth

Purpose

Computes an orthonormal basis for the column space of a matrix.

Format

```
y = orth(x);
```

Input

<i>x</i>	NxK matrix.
----------	-------------

Global Input

`orthtol` scalar, the tolerance for testing if diagonal elements are approaching zero. The default is 1.0e-14.

Output

`y` NxL matrix such that $y'y = \mathbf{eye}(L)$ and whose columns span the same space as the columns of `x`; L is the rank of `x`.

Example

```
x = { 6 5 4,
      2 7 5 };

y = orth(x);
```

After the code above:

```
y = -0.58123819    -0.81373347    y'y = 1    0
      -0.81373347     0.58123819         0    1
```

Source

`qqr.src`

See Also

[qqr](#), [olsqr](#)

output

Purpose

This command makes it possible to direct the output of `print` statements to two different places simultaneously. One output device is always the window or standard output. The other can be selected by the user to be any disk file or other suitable output device such as a printer.

Format

```
output file=filename
output file=filename [on|off|reset];
```

Input

<i>filename</i>	literal or ^string. The <i>file=filename</i> subcommand selects the file or device to which output is to be sent. If the name of the file is to be taken from a string variable, the name of the string must be preceded by the ^ (caret) operator. The default file name is <code>output.out</code> .
<i>on, off, reset</i>	literal, mode flag: <i>on</i> opens the auxiliary output file or device and causes the results of all <code>print</code> statements to be sent to that file or device. If the file already exists, it will be opened for appending. If the file does not already exist, it will be created.

<i>off</i>	closes the auxiliary output file and turns off the auxiliary output.
<i>reset</i>	similar to the <i>on</i> subcommand, except that it always creates a new file. If the file already exists, it will be destroyed and a new file by that name will be created. If it does not exist, it will be created.

Remarks

After you have written to an output file you have to close the file before you can print it or edit it with the **GAUSS** editor. Use

```
output off;
```

The selection of the auxiliary output file or device remains in effect until a new selection is made, or until you get out of **GAUSS**. Thus, if a file is named as the output device in one program, it will remain the output device in subsequent programs until a new *file=filename* subcommand is encountered.

The command

```
output file=filename;
```

will select the file or device but will not open it. A subsequent **output on** or **output reset** will open it and turn on the auxiliary output.

The command **output off** will close the file and turn off the auxiliary output. The filename will remain the same. A subsequent **output on** will cause the file to be opened again for appending. A subsequent **output reset** will cause the existing file to be destroyed and then recreated and will turn on the auxiliary output.

The command **output** by itself will cause the name and status (i.e., open or closed) of the current auxiliary output file to be printed to the window.

The output to the console can be turned off and on using the **screen off** and **screen on** commands. Output to the auxiliary file or device can be turned off or on using the **output off** or **output on** command. The defaults are **screen on** and **output off**.

The auxiliary file or device can be closed by an explicit **output off** statement, by an **end** statement, or by an interactive **new** statement. However, a **new** statement at the beginning of a program will not close the file. This allows programs with **new** statements in them to be run without reopening the auxiliary output file.

If a program sends data to a disk file, it will execute much faster if the window is off.

The **outwidth** command will set the line width of the output file. The default is 80.

Example

```
output file = out1.out on;
```

This statement will open the file `out1.out` and will cause the results of all subsequent **print** statements to be sent to that file. If `out1.out` already exists, the new output will be appended.

```
output file = out2.out;  
output on;
```

This is equivalent to the previous example.

```
output reset;
```

This statement will create a new output file using the current filename. If the file already exists, any data in it will be lost.

```
output file = mydata.asc reset;  
screen off;  
format /m1/rz 1,8;  
open fp = mydata;
```



```
do until eof(fp);  
    print readr(fp,200);;  
endo;  
  
fp = close(fp);  
end;
```

The program above will write the contents of the **GAUSS** file `mydata.dat` into an ASCII file called `mydata.asc`. If there had been an existing file by the name of `mydata.asc`, it would have been overwritten.

The `/m1` parameter in the `format` statement in combination with the `;;` at the end of the `print` statement will cause one carriage return/line feed pair to be written at the beginning of each row of the output file. There will not be an extra line feed added at the end of each 200 row block.

The `end` statement above will automatically perform **output off** and **screen on**.

See Also

[outwidth](#), [screen](#), [end](#), [new](#)

outtyp (dataloop)

Purpose

Specifies the precision of the output data set.

Format

```
outtyp num_constant;
```

Input

<code>num_ constant</code>	scalar, precision of output data set.
--------------------------------	---------------------------------------

Remarks

`num_constant` must be 2, 4, or 8, to specify integer, single precision, or double precision, respectively.

If `outtyp` is not specified, the precision of the output data set will be that of the input data set. If character data is present in the data set, the precision will be forced to double.

Example

```
outtyp 8;
```

outwidth

Purpose

Specifies the width of the auxiliary output.

Format

```
outwidth n;
```

Input

<code>n</code>	scalar, width of auxiliary output.
----------------	------------------------------------

Remarks

n specifies the width of the auxiliary output in columns (characters). After printing *n* characters on a line, **GAUSS** will output a line feed.

If a matrix is being printed, the line feed sequence will always be inserted between separate elements of the matrix rather than being inserted between digits of a single element.

n may be any scalar-valued expressions in the range of 2-256. Nonintegers will be truncated to an integer. If 256 is used, no additional lines will be inserted.

The default is setting is 256.

Example

```
outwidth 132;
```

This statement will change the auxiliary output width to 132 columns.

See Also

[output](#), [print](#)

pacf

p

pacf

Purpose

Computes sample partial autocorrelations.

Format

$rk = \text{pacf}(y, \quad k, \quad d);$

Input

y	Nx1 vector, data.
k	scalar, maximum number of partial autocorrelations to compute.
d	scalar, order of differencing.

Output

rk	Kx1 vector, sample partial autocorrelations.
------	--

Example

Example 1

A sample partial autocorrelation function example.

```
//Create short time-series column vector
```

```
x = { 12.92,
      14.28,
      13.31,
      13.34,
      12.71,
      13.08,
      11.86,
      9.000,
      8.190,
      7.970,
      8.350,
      8.200,
      8.120,
      8.390,
      8.660 };

//Maximum number of lags
k = 4;

//Order of differencing
d = 1;

//Calculate and print result of partial autocorrelation
function
rk = pacf(x,k,d);

print rk;
```

The code above produces the following output:

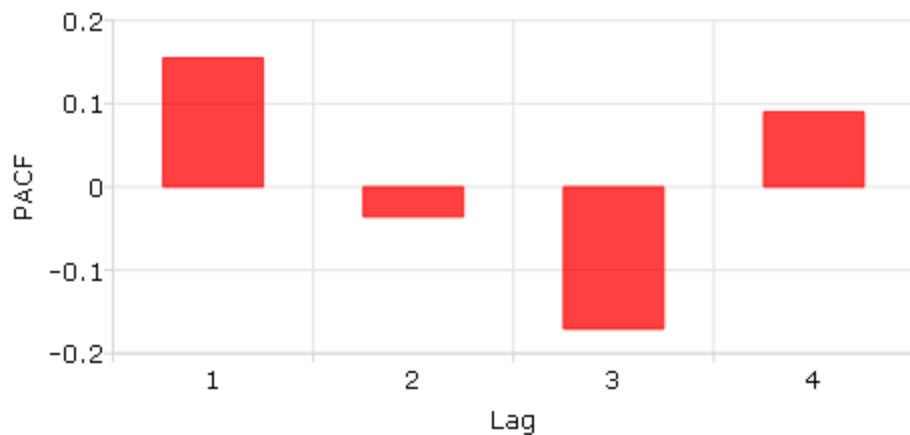
```
0.15488076
-0.035928234
-0.17063786
0.089875096
```

pacf

Plot the PACF results with **plotBar**. Passing in 0 as the first input tells **GAUSS** to create a sequential series from 1 to the number of elements in *rk* as the x-tic labels.

```
plotBar(0, rk);
```

You can add labels for x-axis and y-axis interactively on the Graphics Page by selecting "View->Graph Settings" from the main menu. The plot is shown below:



Example 2

Calculate the partial autocorrelation function (PACF) and plot the results for "beef_prices" data.

```
//Clear out variables in GAUSS workspace
```

```
new;

//Create file name with full path
file = getGAUSSHome() $+ "examples/beef_prices.csv";

//Import data set starting with row 2 and column 2
beef = csvReadM(file,2,2);

//Max lags
k = 10;

//Order of differencing
d = 0;

//Call pacf function
beef_pacf = pacf(beef, k, d);
```

Creat a time series plot and sample partial autocorrelation (PACF) plot based on the *beef* and *beef_pacf* variables created above:

```
//Time series plot
//Declare a plotControl structure
struct plotControl ctl;
ctl = plotGetDefaults("xy");

//Make a 1 by 2 plot with the time series
//plot in the [1,1] location
plotLayout(1,2,1);

//Labels and format settings for 'beef' matrix plot
plotSetYLabel(&ctl, "cents/lb");
plotSetXLabel(&ctl, "Year");
plotSetXTicLabel(&ctl, "YYYY");
plotSetXTicInterval(&ctl, 120, 199501);
```

pacf

```
//Time plot with plotTS function
plotTS(ctl, 1992, 12, beef);

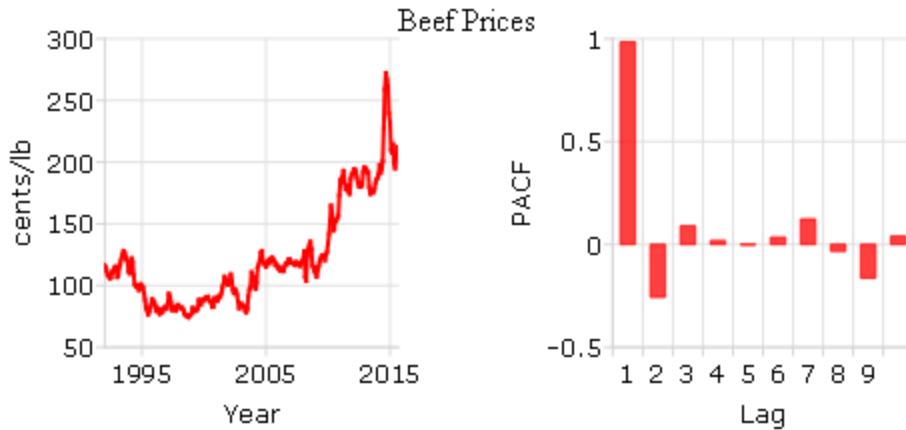
//Making a 1 by 2 plot, the second plot is the PACF plot
plotLayout(1,2,2);

//ACF plot
//Fill 'ctl' structure with defaults settings for bar plots
ctl = plotGetDefaults("bar");

//Setting labels and format based on 'beef_acf' matrix
plotSetYLabel(&ctl, "PACF");
plotSetXLabel(&ctl, "Lag");
plotSetXTicInterval(&ctl, 1, 5);

//PACF plot with plotBar function
plotBar(ctl, seqa(1,1,k), beef_pacf);
```

You can use 'Add Text' to type 'Beef Prices' as the title in the graphics window. The plot is:



Example 3

Compare ACF and PACF for "cow" data.

```
new;
cls;

file = getGAUSSHome() $+ "examples/cows.fmt";

//Import '.fmt' data
load data = ^file;

//Max lags
k = 10;

//Order of differencing
d = 0;

//call pacf function
```

pacf

```
data_pacf = pacf(data, k, d);  
  
//call acf function  
data_acf = acf(data, k, d);
```

In this example, we compute the ACF and PACF for cow's temperature and save them in *data_acf* and *data_pacf*.

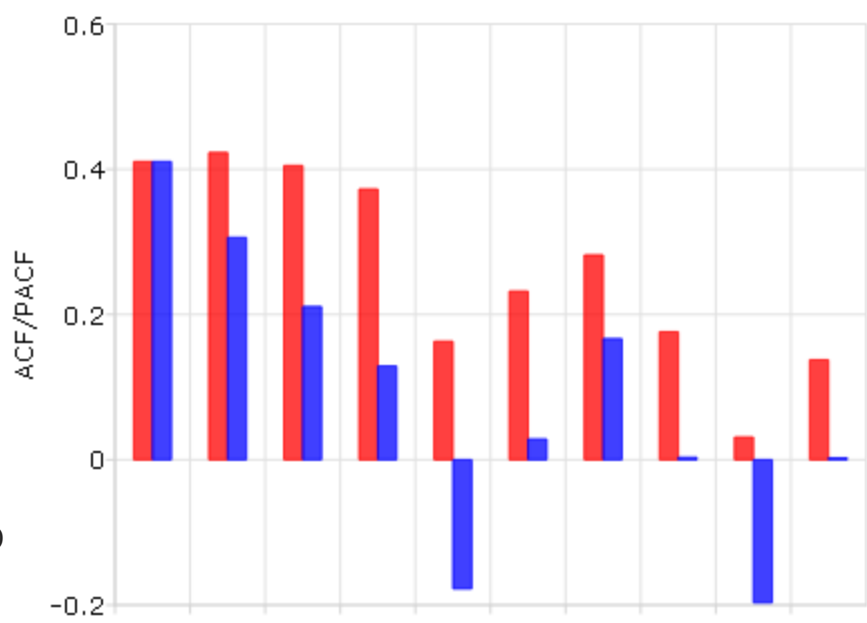
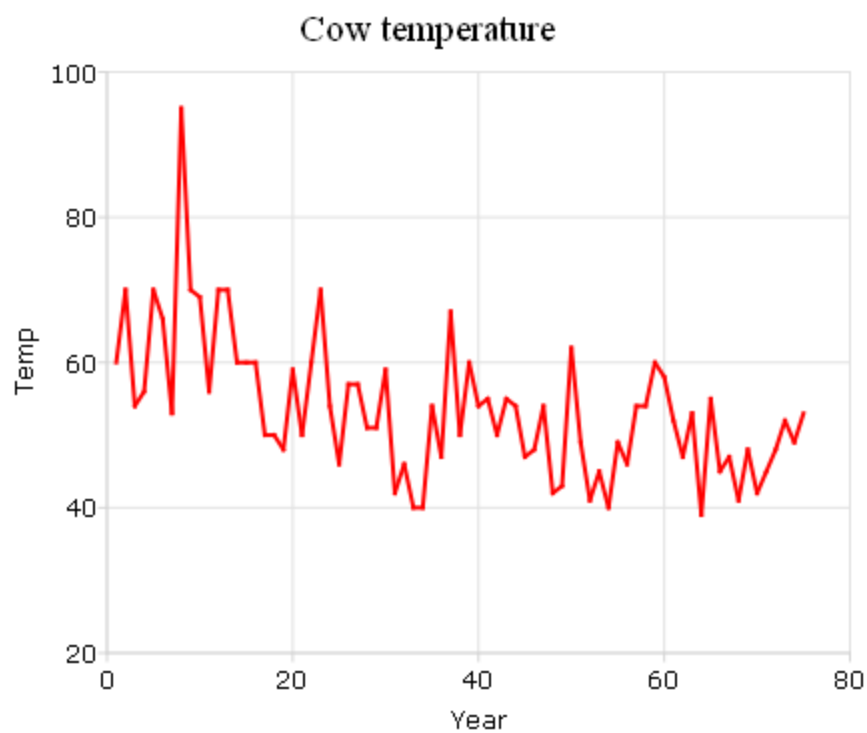
The following code plot autpccorrelation (ACF) and sample partial autocorrelation (PACF) :

```
//Compare ACF and PACF for cow's temperature data  
//Create sequential numbers  
years = seqa(1, 1, rows(data));  
  
//Declare a plotControl structure  
struct plotControl cow_ctl;  
cow_ctl = plotGetDefaults("xy");  
  
//Set plot title for top graph  
plotSetTitle(&cow_ctl, "Cow Temperature");  
  
//Labels and format setting based on 'data_acf' matrix  
plotSetYLabel(&cow_ctl, "Temp");  
plotSetXLabel(&cow_ctl, "Year");  
  
//Making a 2 by 1 plot, the first plot is the time plot  
plotLayout(2,1,1);  
  
//Time plot  
plotXY(cow_ctl,years, data );  
  
//Change type of plotControl struct  
cow_ctl = plotgetdefaults("bar");  
  
//Setting labels and format based on 'data_pacf' matrix
```

```
plotSetYLabel (&cow_ctl, "ACF/PACF");  
plotSetXLabel (&cow_ctl, "Lag");  
  
//Place the 2nd plot in the second cell of a 2 by 1 grid  
plotLayout (2,1,2);  
  
//ACF plot  
plotBar (cow_ctl, seqa (1,1,k), data_acf);  
  
//PACF plot  
plotAddBar (seqa (1,1,k), data_pacf);  
  
//Clear 2 by 1 plot layout for next plots  
plotClearLayout ();
```

The plot produced by the code above should look like this:

pacf



p

35-1170

Source

tsutil.src

See Also

[acf](#)

packedToSp

Purpose

Creates a sparse matrix from a packed matrix of non-zero values and row and column indices.

Format

$y = \text{packedToSp}(r, c, p);$

Input

r	scalar, rows of output matrix.
c	scalar, columns of output matrix.
p	$N \times 3$ or $N \times 4$ matrix, containing non-zero values and row and column indices.

Output

y	$r \times c$ sparse matrix.
-----	-----------------------------

Remarks

If p is $N \times 3$, y will be a real sparse matrix. Otherwise, if p is $N \times 4$, y will be

packedToSp

complex.

The format for `p` is as follows:

If `p` is Nx3:

Column 1	Column 2	Column 3
non-zero values	row indices	column indices

If `p` is Nx4:

Column 1	Column 2	Column 3	Column 4
real non-zero values	imaginary non-zero values	row indices	column indices

Note that **spCreate** may be faster.

Since sparse matrices are strongly typed in **GAUSS**, `y` must be defined as a sparse matrix before the call to **packedToSp**.

Example

p

```
//Declare 'y' to be a sparse matrix
sparse matrix y;

//Create a 15x10 matrix 'y' in which:
//y[2,4] = 1.1; y[5,1] = 2.3; y[8,9] = 3.4;
//y[13,5] = 4.2
//all other values in 'y' will be zeros
p = { 1.1 2 4, 2.3 5 1, 3.4 8 9, 4.2 13 5 };
y = packedToSp(15,10,p);
```

After the code above, `y` is a sparse matrix, containing the following non-zero values:

Non-zero value	Index
1.1	(2, 4)
2.3	(5, 1)
3.4	(8, 9)
4.2	(13, 5)

See Also

[spCreate](#), [denseToSp](#)

packr

Purpose

Deletes the rows of a matrix that contain any missing values.

Format

$y = \text{packr}(x);$

Input

x	$N \times K$ matrix.
-----	----------------------

Output

y	$L \times K$ submatrix of x containing only those rows that do not have missing values in any of their elements.
-----	--

Remarks

This function is useful for handling missing values by "listwise deletion," particularly

packr

prior to using the / operator to compute least squares coefficients.

If all rows of a matrix contain missing values, **packr** returns a scalar missing value. This can be tested for quickly with the **scalmiss** function.

Example

Example 1

```
//Set the rng seed for repeatable random numbers
rndseed 7342692;

//Create a 3x3 matrix of random integers between 1 and 10
x = ceil(rndu(3, 3) * 10);

//Turn all elements with a value of 8 into missing values
x2 = miss(ceil(rndu(3,3)*10),8);

//Remove all rows that contain missing values
y = packr(x2);
```

After the code above:

	6	10	3		6	10	3	
x =	8	7	8	x2 =	.	7	.	y = 6 10 3
	8	6	7		.	6	7	

Example 2

```
//Open a GAUSS data file for reading
open fp = mydata;
obs = 0;
sum = 0;

//Continue looping until the end of the file has been
//reached
```



```

do until eof(fp);
  //Read in 100 lines of the data file and remove any rows
  //with missing values
  x = packr(readr(fp,100));
  //Check to see if 'packr' returned a missing value; if
  //not, update 'obs' and 'sum'
  if not scalmiss(x);
    obs = obs + rows(x);
    sum = sum + sumc(x);
  endif;
enddo;
mean = sum/obs;

```

In this example the sums of each column in a data file are computed as well as a count of the rows that do not contain any missing values. **packr** is used to delete rows that contain missings and **scalmiss** is used to skip the two sum steps if all the rows are deleted for a particular iteration of the read loop. Then the sums are divided by the number of observations to obtain the means.

See Also

[scalmiss](#), [miss](#), [missrv](#)

parse

Purpose

Parses a string, returning a character vector of tokens.

Format

```
tok = parse(str, delim);
```

parse

Input

<i>str</i>	string consisting of a series of tokens and/or delimiters.
<i>delim</i>	NxK character matrix of delimiters that might be found in <i>str</i> .

Output

<i>tok</i>	Mx1 character vector consisting of the tokens contained in <i>str</i> . All tokens are returned; any delimiters found in <i>str</i> are ignored.
------------	--

Remarks

The tokens in *str* must be 8 characters or less in size. This is because they are returned in a character vector in which each element is represented as a double precision value. If they are longer, the contents of *tok* is unpredictable. Use string arrays to create arrays of text with elements longer than 8 characters.

Example

Example 1

```
names = "GDP;GNP;M1;M2";
namesVec = parse(names, ";");

//The '$' is used when printing character vectors
print $namesVec;
```

The code above will return:

```
GDP
GNP
```

M1

M2

Example 2

```

obs = 1000;
names = "Age,Weight,Height";

//Create uniform random integers between 1 and 77
data1 = ceil(77 * rndu(obs,1));

//Create normal random integers centered at 100 with a
//standard deviation of 9
data2 = ceil(100 + 9*rndn(obs,1));

//Create uniform random numbers between 0 and 60
data3 = ceil(60 * rndu(obs,1));

//Horizontally concatenate data into 'obs'x3 matrix
data = data1~data2~data3;

//Print the data using the procedure below
printStats(names, data);

//Create procedure to take our data, calculate some basic
//stats and print them
proc (0) = printStats( names, data);
    local title, vars, sepVars;

    //Set to print with 6 spaces between numbers and 0
    //digits after the decimal
    format /rd 6,0;

    //Create the titles to print for each column
    title = parse("var,mean,max,min", ",");

```

p

parse

```
//Extract the substrings from 'names' into a character
//array using the comma as a separator between tokens
sepVars = parse(names, ",");
print "-----";

//The '$' tells GAUSS to print as character data
print $title';
print "-----"
//Loop through as many times as there are rows in
//'sepVars'
for i( 1, rows(sepVars), 1);
    //Two semi-colons at the end of a print statement
    //prevents a new-line after the print
    print $sepVars[i];;
    print meanc(data[:,i]);;
    print maxc(data[:,i]);;
    print minc(data[:,i]);
endfor;
print "-----";
endp;
```

The code above will produce output like this:

var	mean	max	min

Age	38	77	1
Weight	101	135	75
Height	31	60	1

See Also

[token](#)

pause

Purpose

Pauses for a specified number of seconds.

Format

```
pause(sec);
```

Input

<i>sec</i>	scalar, seconds to pause.
------------	---------------------------

Remarks

This function can be used to delay a program, allowing users time to view graphics and/or data printed to the program output window.

Source

`pause.src`

See Also

[wait](#)

pdfBinomial

Purpose

Computes the binomial probability density function.

Format

```
p = pdfBinomial(successes, trials, prob);
```

pdfBinomial

Input

<i>successes</i>	NxK matrix, Nx1 vector or scalar. <i>successes</i> must be a positive number and < trials
<i>trials</i>	ExE conformable with <i>successes</i> . <i>trials</i> must be > <i>successes</i> .
<i>prob</i>	The probability of success on any given trial. ExE conformable with <i>successes</i> . $0 < \text{prob} < 1$.

Output

<i>p</i>	The probability of the specified number of successes, NxK matrix, Nx1 vector or scalar.
----------	---

Example

A polling company randomly selects 1,024 prospective voters in a region where 55% support their candidate. What is the probability that exactly 600 of those selected support their candidate?

```
p = pdfBinomial(600, 1024, 0.55);
```

After running the code above, *p* is equal to:

```
0.0017226334
```

Continuing with the example above, what would be the probability of selecting the same number of voters that support their candidate if their candidate's support in the region was 50% or 60%?

```
p_support = { 0.5, 0.6 };  
p = pdfBinomial(600, 1024, p_support);
```

After running the code above, *p* is equal to:

```
6.3351627e-09
0.016621105
```

Remarks

The probability density function for the binomial distribution is defined as:

$$P(x = k|n, p) = \binom{n}{k} p^k (1 - p)^{n-k}$$

where k is the number of successes, n is the number of trials and p is the probability of success on each trial.

For invalid inputs, **pdfBinomial** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[cdfBinomial](#), [cdfBinomialInv](#)

pdfCauchy

Purpose

Computes the probability density function for the Cauchy distribution.

Format

```
y = pdfCauchy(x, mu, sigma);
```

p

pdfexp

Input

x	NxK matrix, an Nx1 vector or scalar.
μ	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x .
σ	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . σ must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

The probability density function for the Cauchy distribution is defined as

$$f(x) = \left(\pi \sigma \left(1 + \left(\frac{x - \mu}{\sigma} \right)^2 \right) \right)^{-1}$$

See Also

[cdfCauchy](#)

p

pdfexp

Purpose

Computes the probability density function for the exponential distribution.

Format

$y = \text{pdfexp}(x, a, b);$

Input

x	NxK matrix, Nx1 vector or scalar. x must be greater than a .
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x .
b	Scalar, the scale parameter sometimes called beta. b must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

pdfExp calculates the probability density function for the two-parameter exponential distribution, which is defined as

$$f(x) = \frac{1}{b} \exp\left(-\frac{x-a}{b}\right)$$

See Also

[cdfexp](#)

pdfGenPareto

Purpose

Computes the probability density function for the Generalized Pareto distribution.

Format

```
 $y = \text{pdfGenPareto}(x, a, \text{sigma}, k);$ 
```

Input

x	NxK matrix, an Nx1 vector or scalar.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x .
$sigma$	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . $sigma$ must be greater than 0.
k	Shape parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x .

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

The probability density function for the Generalized Pareto distribution is defined as

$$f(x) = \begin{cases} \frac{1}{\sigma} \left(1 + k \frac{(x-\mu)}{\sigma}\right)^{-1-1/k} & k \neq 0 \\ \frac{1}{\sigma} \exp\left(-\frac{(x-\mu)}{\sigma}\right) & k = 0 \end{cases}$$

See Also

[cdfGenPareto](#)

pdfHyperGeo

Purpose

Computes the probability mass function for the hypergeometric distribution.

Format

$p = \text{pdfHyperGeo}(x, m, k, n);$

Input

x	NxK matrix, Nx1 vector or scalar. x must be a positive number and $< m$
m	The size of the population from which draws will be made. ExE conformable with x . m must be $> x$, k and n .
k	The number of marked items. ExE conformable with x .
n	The number of items drawn from the population. ExE conformable with x . $0 < k < m$.

Output

p	The probability of drawing x marked items. NxK matrix, Nx1 vector or scalar.
-----	--

Example

You are given 50 hard drives, 4 of which are known to be bad. What is the probability of drawing exactly 1 bad hard drive if you randomly select 6 drives?

p

pdfHyperGeo

```
p = pdfHyperGeo(1, 50, 4, 6);
```

After running the code above, p is equal to:

```
0.34504559
```

Continuing with the example above, what are the probabilities of drawing exactly 2 or exactly 4 bad hard drives?

```
x = { 2, 4 };  
p = pdfHyperGeo(x, 50, 4, 6);
```

After running the code above, p is equal to:

```
0.061615284  
6.5132436e-05
```

Remarks

The probability density function for the hypergeometric distribution is defined as:

$$P(x|m, k, n) = \frac{\binom{k}{x} \binom{m-k}{n-x}}{\binom{m}{n}}$$

For invalid inputs, **pdfHyperGeo** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

See Also

[cdfHyperGeo](#), [rndHyperGeo](#), [pdfBinomial](#)

pdfLaplace

Purpose

Computes the probability density function for the Laplace distribution.

Format

```
y = pdfLaplace(x, a, b);
```

Input

x	NxK matrix, Nx1 vector or scalar.
a	Scalar, location parameter.
b	Scalar, scale parameter. b must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

The probability density function for the Laplace distribution is defined as

$$f(x) = \frac{1}{2b} \exp\left(-\frac{|x-a|}{b}\right)$$

See Also

[cdfCauchy](#), [pdfCauchy](#)

pdflogistic

pdflogistic

Purpose

Computes the probability density function for the logistic distribution.

Format

$y = \text{pdflogistic}(x, a, b);$

Input

x	NxK matrix, an Nx1 vector or scalar.
a	Location parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x .
b	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with x . b must be greater than 0.

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

The probability density function for the logistic distribution is defined as

$$f(x) = \frac{\exp(z)}{b(1 + \exp(z))^2}$$
$$z = -\left(\frac{x-a}{b}\right)$$

See Also

[cdflogistic](#)

pdfn

Purpose

Computes the standard Normal (scalar) probability density function.

Format

```
y = pdfn(x);
```

Input

x	$N \times K$ matrix.
-----	----------------------

Output

y	$N \times K$ matrix containing the standard Normal probability density function of x .
-----	--

Remarks

This does not compute the joint Normal density function. Instead, the scalar Normal density function is computed element-by-element. y could be computed by the following **GAUSS** code:

```
y = (1/sqrt(2*pi)) * exp(-(x.*x)/2);
```

Example

```
x = { -3, -2, 0, 2, 3 };  
y = pdfn(x);
```

After the code above:

pdfPoisson

```
0.0044318484
0.053990967
y = 0.39894228
0.053990967
0.0044318484
```

pdfPoisson

Purpose

Computes the Poisson probability mass function.

Format

```
p = pdfPoisson(x, lambda);
```

Input

<i>x</i>	NxK matrix, Nx1 vector or scalar. <i>x</i> must be a positive whole number.
<i>lambda</i>	ExE conformable with <i>x</i> . The mean parameter.

Output

<i>p</i>	NxK matrix, Nx1 vector or scalar.
----------	-----------------------------------

Remarks

For invalid inputs, **pdfPoisson** will return a scalar error code which, when its value is assessed by function **scalerr**, corresponds to the invalid input. If the first input is out of range, **scalerr** will return a 1; if the second is out of range, **scalerr** will return a 2; etc.

p

Example

Example 1: Basic example

```
p = pdfPoisson(190,200);
```

After the code above, p is equal to:

```
0.02243
```

Example 2: Vector input

```
events = { 170,  
           180,  
           190,  
           200 };  
p = pdfPoisson(events,200);
```

After the code above, p is equal to:

```
0.00285  
0.01056  
0.02243  
0.02820
```

Example 3: Vector Inputs

```
events = { 170,  
           180,  
           190,  
           200 };  
lambda = { 180,  
           190,  
           200,  
           210 };
```

p

pdfRayleigh

```
p = pdfPoisson(events,lambda);
```

After the code above, *p* is equal to:

```
0.02304
0.02274
0.02243
0.02214
```

See Also

[cdfPoisson](#), [cdfPoissonInv](#), [rndPoisson](#)

pdfRayleigh

Purpose

Computes the probability density function of the Rayleigh distribution.

Format

```
y = pdfRayleigh(x,b);
```

Input

p

<i>x</i>	NxK matrix, an Nx1 vector or scalar. <i>x</i> must be greater than 0.
<i>b</i>	Scale parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with <i>x</i> . <i>b</i> must be greater than 0.

Output

<i>y</i>	NxK matrix, Nx1 vector or scalar.
----------	-----------------------------------

Remarks

The probability density function of the Rayleigh distribution is defined as

$$f(x) = \frac{x}{b^2} \exp\left(\frac{-x^2}{2b^2}\right)$$

See Also

[cdfRayleighinv](#)

pdfWeibull

Purpose

Computes the probability density function of a Weibull random variable.

Format

```
y = pdfWeibull(x, k, lambda);
```

Input

<i>x</i>	NxK matrix, Nx1 vector or scalar. <i>x</i> must be greater than 0.
<i>k</i>	Shape parameter; NxK matrix, Nx1 vector or scalar, ExE conformable with <i>x</i> . <i>k</i> must be greater than 0.
<i>lambda</i>	Scale parameter; may be matrix, Nx1 vector or scalar, ExE conformable with <i>x</i> . <i>lambda</i> must be greater than 0.

pdfWishartInv

Output

y	NxK matrix, Nx1 vector or scalar.
-----	-----------------------------------

Remarks

The probability density function of a Weibull random variable is defined as

$$f(x, \lambda, k) = \begin{cases} \frac{k}{\lambda} \left(\frac{x}{\lambda}\right)^{k-1} e^{-(x/\lambda)^k} & x \geq 0 \\ 0 & x < 0 \end{cases}$$

See Also

[cdfWeibull](#), [cdfWeibullInv](#)

pdfWishartInv

Purpose

Computes the probability density function of the inverse Wishart distribution.

Format

$y = \text{pdfWishartInv}(IW, S, df);$

Input

p

IW	p x p positive definite matrix. T
S	p x p positive definite scale matrix. Ψ
df	Scalar, degree of freedom. ν

Output

y

Scalar, probability density function.

Remarks

pdfWishartInv calculates the probability density function for the inverse Wishart distribution, which is defined as

$$f(T) = \frac{|\Psi|^{\nu/2}}{|T|^{\frac{\nu+p+1}{2}} 2^{\frac{\nu p}{2}} \Gamma_p\left(\frac{\nu}{2}\right)} \exp\left(-\frac{1}{2} \text{tr}(\Gamma T^{-1})\right)$$

Example

```
new ;
cls ;
rndseed 2223;

x = {9.2517907  7.4283670,
      7.4283670 10.503325 };

S = {1 .5, .5 1};

df = 3;

// pdf of inverse Wishart distribution
y = pdfWishartInv(x, S, df);

print y;
```

After above code,

6.0267322e-007

pi

See also

[rndWishart](#), [rndWishartInv](#)

pi

Purpose

Returns the mathematical constant π .

Format

$y = \text{pi};$

Output

y scalar, the value of π .

Example

```
//Print 14 digits and allow 16 digits worth of space for  
//each printed number  
format /rdn 16,14;  
print pi;
```

will return:

3.14159265358979

pinv

Purpose

Computes the Moore-Penrose pseudo-inverse of a matrix, using the singular value decomposition. This pseudo-inverse is one particular type of generalized inverse.

Format

$y = \text{pinv}(x);$

Input

x	NxM matrix.
-----	-------------

Global Input

$_svdtol$	scalar, any singular values less than $_svdtol$ are treated as zero in determining the rank of the input matrix. The default value for $_svdtol$ is 1.0e-13.
------------	--

Output

y	MxN matrix that satisfies the 4 Moore-Penrose conditions: $xyx = x$ $yxy = y$ xy is symmetric yx is symmetric
-----	---

pinv

Global Output

`_svderr` scalar, if not all of the singular values can be computed `_svderr` will be nonzero.

Example

pinv can be used to solve an undertermined least squares problem.

```
//Create an underdetermined system of equations 'A'
A = randn(4, 5);

//Create a right hand side
b = randn(4,1);

if rank(A) < cols(A);
    print "A does not have full rank, using pinv to solve";
    Api = pinv(A);
    x = Api*b;
else;
    print "A has full rank, solve with '/' operator";
    x = b/A;
endif;
```

Least squares problems with full rank can also be solved with the **GAUSS** functions: **ols**, **olsqr** and **olsqr2**.

Source

svd.src

pinvmt

Purpose

Computes the Moore-Penrose pseudo-inverse of a matrix, using the singular value decomposition. This pseudo-inverse is one particular type of generalized inverse.

Format

```
{ y, err } = pinvmt(x, tol);
```

Input

<i>x</i>	NxM matrix.
<i>tol</i>	scalar, any singular values less than <i>tol</i> are treated as zero in determining the rank of the input matrix.

Output

<i>y</i>	MxN matrix that satisfies the 4 Moore-Penrose conditions: $xyx = x$ $yxy = y$ xy is symmetric yx is symmetric
<i>err</i>	scalar, if not all of the singular values can be computed <i>err</i> will be nonzero.

pinvmt can be used to solve an undertermined least squares problem.

```
tol = 1e-13;

//Create an underdetermined system of equations 'A'
```

plotAddArea

```
A = rndn(4, 5);

//Create a right hand side
b = rndn(4,1);

if rank(A) < cols(A);
    print "A does not have full rank, using pinvmt to
solve";
    Api = pinvmt(A, tol);
    x = Api*b;
else;
    print "A has full rank, solve with '/' operator";
    x = b/A;
endif;
```

Least squares problems with full rank can also be solved with the **GAUSS** functions: **ols**, **olsqr** and **olsqr2**.

Source

svdmt.src

plotAddArea

Purpose

Adds a cumulative area plot to an existing graph.

Format

```
plotAddArea(myPlot, x, y);
plotAddArea(x, y);
plotAddArea(myPlot, x, y, base);
plotAddArea(x, y, base);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Nx1 vector containing the X coordinates.
<i>y</i>	Nx1 or NxM matrix. Each column contains the height for the corresponding section of the graph. If <i>y</i> contains more than one column, each column will be stacked on top of the previous column.
<i>base</i>	Scalar optional argument. The height for the base of the added area plot. The default value is zero. plotAddArea does not yet support a vector input for <i>base</i> .

Remarks

plotAddArea may only add curves to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

Example

Creating confidence intervals with **plotAddArea**

```
//Create the 'x' and 'y' for the normal
//probability density function
x = sega(-3.5, 0.1, 71);
y = pdfn(x);

//Create the 'x' and 'y' points
//for the left tail
edge = cdfni(0.05);
x_ci = selif(x, (x .< edge));
y_ci = y[1:rows(x_ci)];
```

plotAddArea

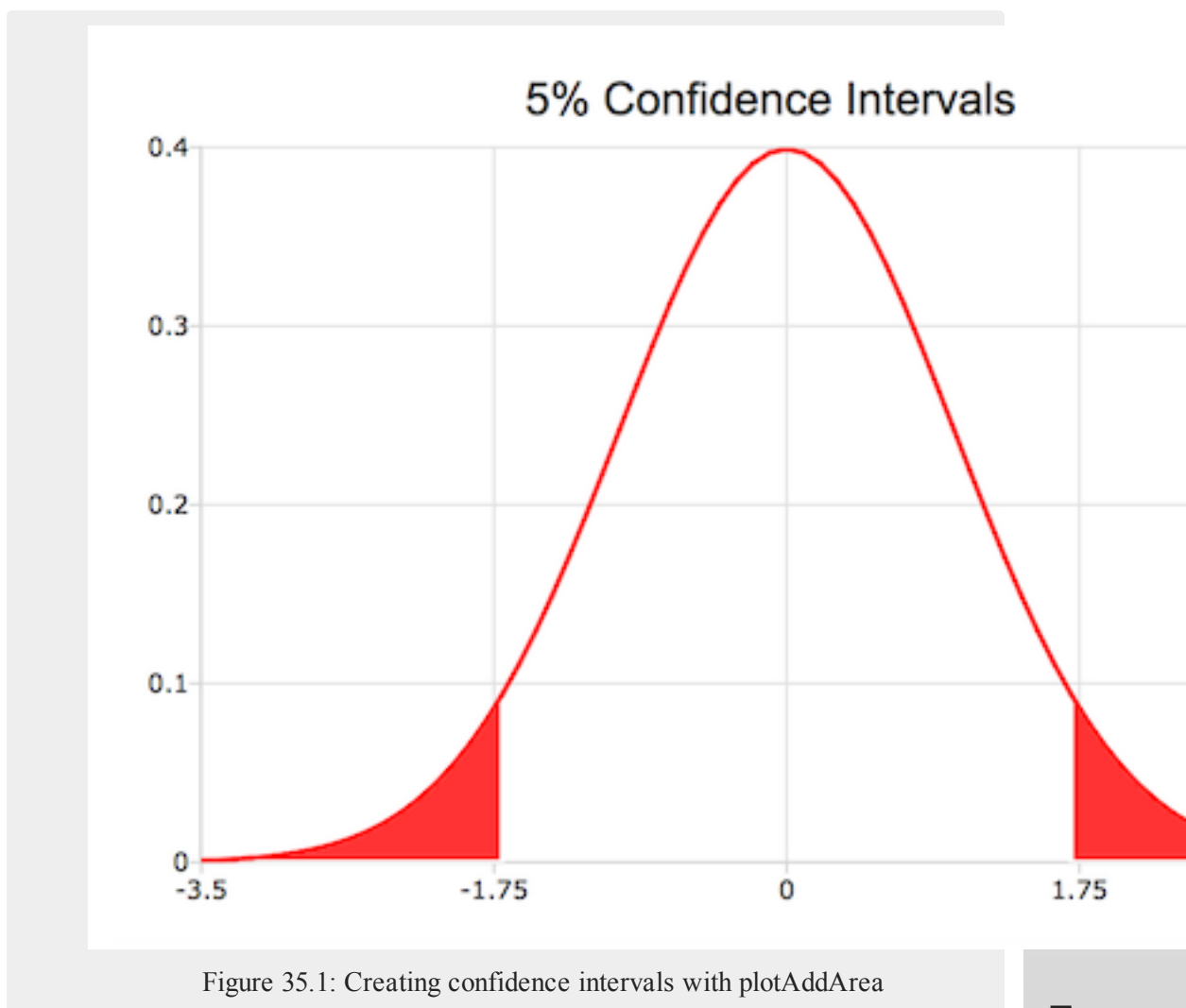
```
//Draw filled in left tail
plotArea(x_ci, y_ci);

//Create the 'x' and 'y' points
//for the right tail
edge = cdfni(0.95);
x_ci = selif(x, (x .> edge));
y_ci = y[rows(y)-rows(x_ci)+1:rows(y)];

//Add right tail to graph
plotAddArea(x_ci, y_ci);

//Add pdfn line
plotAddXY(x, y);
```

The code to create the graph below can be found in the file `plotaddci.e` in your GAUSS examples directory.

**See Also**

[plotAddBar](#), [plotAddHist](#), [plotAddHistF](#), [plotAddHistP](#), [plotAddPolar](#)

plotAddArrow

plotAddArrow

Purpose

Adds an arrow to an existing graph.

Format

```
plotAddArrow(myAnnotation, x_start, y_start, x_end, y_end, head_size);
plotAddArrow( x_start, y_start, x_end, y_end, head_size);
```

Input

<i>myAnnotation</i>	Optional argument, an instance of a plotAnnotation structure.
<i>x_start</i>	Scalar or Nx1 vector, the X coordinate for the start of each respective arrow.
<i>y_start</i>	Scalar or Nx1 vector, the Y coordinate for the start of each respective arrow.
<i>x_end</i>	Scalar or Nx1 vector, the X coordinate for the end of each respective arrow.
<i>y_end</i>	Scalar or Nx1 vector, the Y coordinate for the end of each respective arrow.
<i>head_size</i>	2x1 vector, the size of the arrowhead(s) in pixels. The first element of <i>head_size</i> is the size for head at the end of the arrow. The second element is the size of the head at the start of the arrow.

Remarks

Please note that **plotAddArrow** will add arrows to existing graphs, it will not create

a new graph if one does not exist. **plotAddArrow** is not yet supported for surface plots.

Examples

Example 1: Basic usage

```
x_start = 0.2;
y_start = 0.25;
x_end = 0.4;
y_end = 0.5;

//Set arrowhead at the end to 15 px
//No arrowhead at the beginning of the arrow
head_size = { 15, 0 };

//Add an arrow to graph
plotAddArrow(x_start, y_start, x_end, y_end, head_size);
```

Example 2: Add an arrow between points

```
//Draw random scatter plot
x = rndu(10,1);
y = rndu(10, 1);
plotScatter(x, y);

//Add arrow from the first point to the ninth point
plotAddArrow(x[1], y[1], x[9], y[9], 12);
```

See Also

[plotAddTextbox](#), [annotationGetDefaults](#), [annotationSetLineColor](#)

plotAddBar

plotAddBar

Purpose

Adds a bar or a set of bars to an existing graph.

Format

```
plotAddBar(myPlot, val, ht);  
plotAddBar(val, ht);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>val</i>	Nx1 numeric vector, bar labels. If scalar 0, a sequence from 1 to rows (<i>ht</i>) will be created.
<i>ht</i>	NxK numeric vector, bar heights. K overlapping or side-by-side sets of N bars will be graphed.

Remarks

plotAddBar may only add bars to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

See Also

[plotAddHist](#), [plotAddHistF](#), [plotAddHistP](#), [plotAddPolar](#), [plotAddXY](#)

plotAddBox

Purpose

Adds a box graph to an existing graph.

Format

```
plotAddBox(myPlot, grp, y);  
plotAddBox(grp, y);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>grp</i>	1xM vector. This contains the group numbers corresponding to each column of <i>y</i> data. If scalar 0, a sequence from 1 to cols (<i>y</i>) will be generated automatically for the X axis.
<i>y</i>	NxM matrix. Each column represents the set of <i>y</i> values for an individual percentiles box symbol.

Remarks

plotAddBox may only add a box graph to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

See Also

[plotAddHist](#), [plotAddHistF](#), [plotAddHistP](#), [plotAddPolar](#), [plotAddXY](#)

plotAddErrorBar

plotAddErrorBar

Purpose

Adds an error bar or a set of bars to an existing 2-D graph.

Format

```
plotAddErrorBar(myPlot, x, y, err);
plotAddErrorBar(myPlot, x, y, lwr, uppr);
plotAddErrorBar(x, y, err);
plotAddErrorBar(x, y, lwr, uppr);
```

Input

<i>myPlot</i>	Optional input, a plotControl structure.
<i>x</i>	Scalar, Nx1 or NxM matrix. Each column contains the X values for a particular line.
<i>y</i>	Scalar, Nx1 or NxM matrix. Each column contains the Y values for a particular line.
<i>err</i>	Scalar, Nx1 or NxM matrix. Each column contains the error values for each Y column.
<i>lwr</i>	Nx1 or NxM matrix. Each column contains the lower bar heights for assymetrical

uppr

error bars.

Nx1 or NxM matrix. Each column contains the upper bar heights for assymetrical error bars.

Examples

Example 1: Basic addition of error bars to scatter plot

```
new;

//Sequence 1, 2, 3...12
x = seqa(1, 1, 12);

//Create some random normal data, y ~ N(0, 1)
y = rndn(12,1);

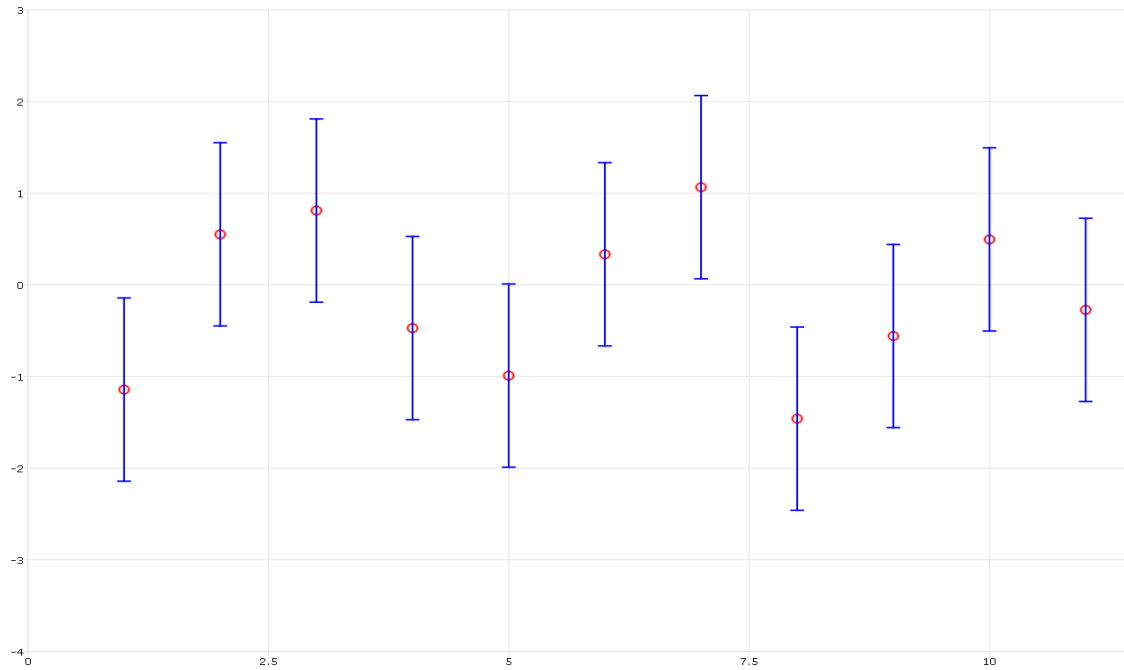
//12x1 vector of all 1's
sd = ones(12,1);

//Draw basic scatter plot, using default settings
plotScatter(x, y);

//Add error bars with height of 1
plotAddErrorBar(x, y, sd);
```

The plot is

plotAddErrorBar



Example 2: Assymetrical error bars to plot median and range

```
new;  
  
//Create the sequence 1, 2, 3...9  
x = seqa(1, 1, 9);  
  
//Create 9 gamma distributed column vectors  
y = rndGamma(100, 9, 2, 4);  
  
//Calculate medians of each column  
med = median(y);  
  
//Draw scatter plot of each median
```

```

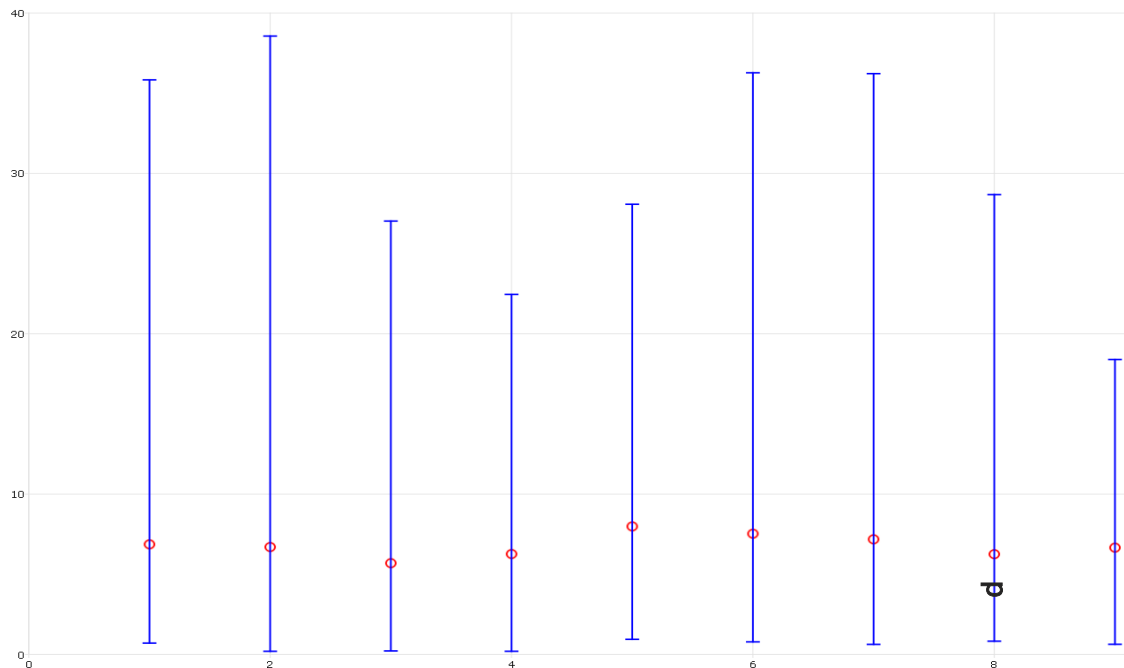
plotScatter(x, med);

//Calculate the distance between the median
//and the min and max for each column
err_high = maxc(y) - med;
err_low = (med - minc(y));

//Add assymetrical error bars to span
//the range of each column
plotAddErrorBar(x, med, err_low, err_high);

```

The plot is



Example 3: Add error bars to XY plot, using plotControl structure

plotAddErrorBar

```
new;

//Create x values
x = seqa(0,1,11);

//Create y values
y = 10*randn(11,1);

//Define plotControl Structure
struct plotControl myPlot;
myPlot = plotGetDefaults("xy");

//Set color for main line
plotSetLineColor(&myPlot, "steel blue");

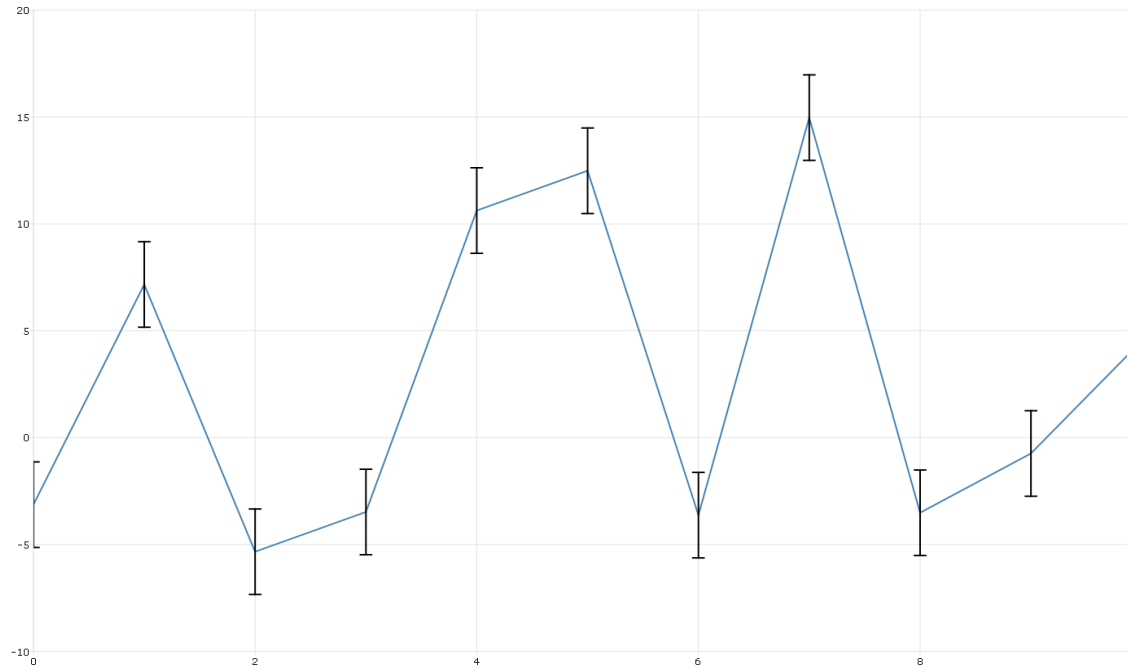
//Plot line
plotXY(myPlot, x, y);

//Set error bar values
err = ones(11,1)*2;

//Set error bar color to be different than XY line
plotSetLineColor(&myPlot, "black");

//Add error bars
plotAddErrorBar(myPlot, x,y,err);
```

The plot is



Example 4: Add error bars to bar plot

```
new;  
  
//Load 'Age' and 'Limit' variables  
//into a 2 column matrix  
file = getGAUSSHome() $+ "/examples/credit.dat";  
data = loadadd(file, "Age + Limit");  
  
age = data[:,1];  
limit = data[:,2];  
  
num_ranges = 4;
```

plotAddErrorBar

```
age_ranges = { 20 40,
               40 60,
               60 80,
               80 100 };

//Pre-allocate vector to hold means
//and standard deviation of the samples
mu = zeros(num_ranges,1);
s = zeros(num_ranges,1);

for i(1, num_ranges, 1);
    // Get the index for different age level
    level_idx = indexcat(age, age_ranges[i,.]');

    // mean of limit for different age level
    mu[i] = mean(limit[level_idx]);

    // Calculate the standard deviation of the sample
    s[i] = std(limit[level_idx]);
endfor;

// Note: < is '&lt;'; in html
// '$|' is string concatenation
labels = "20 &lt; age &le; 40" $|
         "40 &lt; age &le; 60" $|
         "60 &lt; age &le; 80" $|
         "80 &lt; age &le; 100";

// Declare 'myPlot' to be a plotControl structure
// and fill with default settings for bar plots
struct plotControl myPlot;
myPlot = plotGetDefaults("bar");

//Set bar fill to be: solid, 100% opaque and steel blue
plotSetFill(&myPlot, 1, 1, "steel blue");
```



```
//Set title and axes labels
plotSetTitle(&myPlot, "Credit Limits and Age", "arial",
20);
plotSetYLabel(&myPlot, "Credit Limits", "arial", 18);
plotSetXLabel(&myPlot, "Age", "arial", 18);

//Draw bar plot
plotBar(myPlot, labels, mu);

//File 'myPlot' plotControl structure with
//default settings for 'xy' plots
myPlot = plotGetDefaults("xy");

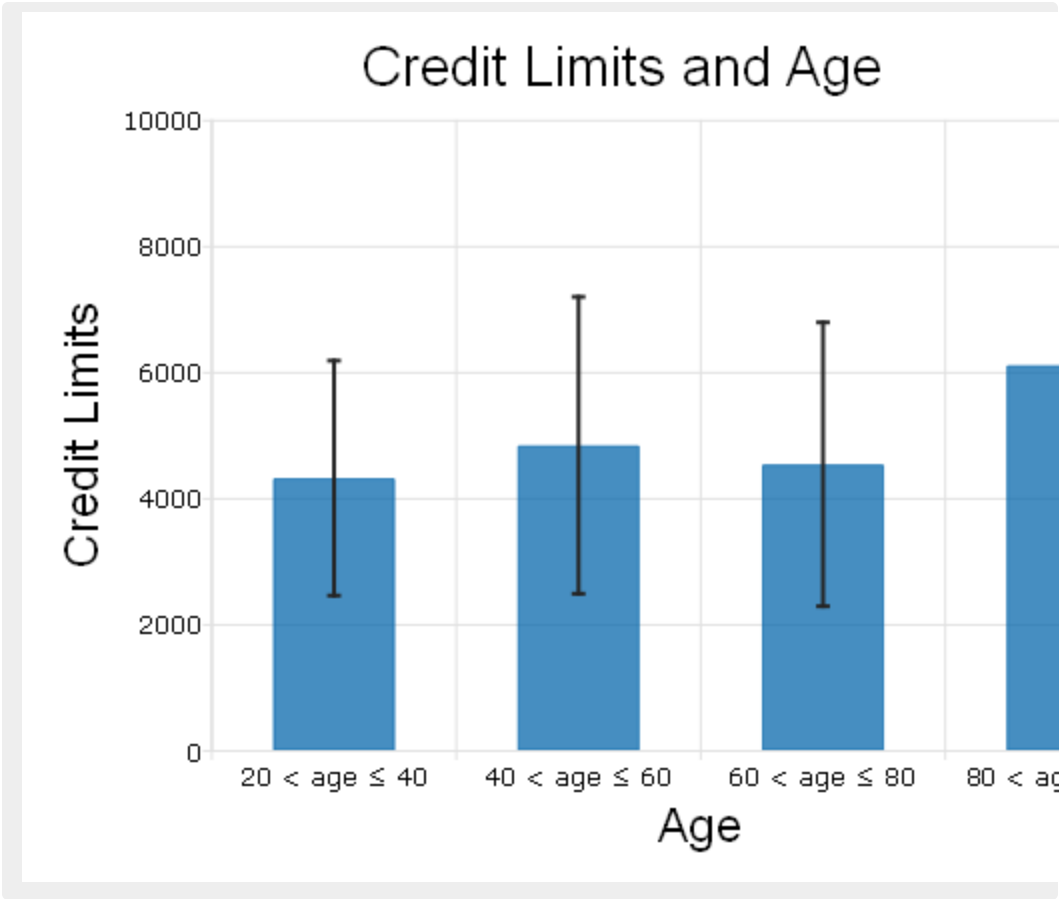
plotSetLineColor(&myPlot, "black");

//'x' location of error bars
//1 is first bar, 2 is second bar, etc
x = seqa(1, 1, num_ranges);

//Draw error bars on bar plot
plotAddErrorBar(myPlot, x, mu, s);
```

The plot is

plotAddHist



See Also

[plotBar](#)

plotAddHist

Purpose

Adds a histogram to an existing graph.

Format

```
plotAddHist(myPlot, x, v);  
plotAddHist(x, v);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Mx1 vector of data.
<i>v</i>	Nx1 vector, the breakpoints to be used to compute the frequencies - or - scalar, the number of categories.

Remarks

plotAddHist may only add a histogram to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

See Also

[plotAddBar](#), [plotAddHistF](#), [plotAddHistP](#), [plotAddPolar](#), [plotAddXY](#)

plotAddHistF

Purpose

Adds a frequency histogram to an existing graph.

plotAddHistP

Format

```
plotAddHistF(myPlot, f, c);  
plotAddHistF(f, c);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>f</i>	Nx1 vector, frequencies to be graphed.
<i>c</i>	Nx1 vector, numeric labels for categories. If this is a scalar 0, a sequence from 1 to rows (<i>f</i>) will be created.

Remarks

plotAddHistF may only add a histogram to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

See Also

[plotAddBar](#), [plotAddHist](#), [plotAddHistP](#), [plotAddPolar](#), [plotAddXY](#)

plotAddHistP

Purpose

Adds a percent histogram to an existing graph.

Format

```
plotAddHistP(myPlot, x, v);  
plotAddHistP(x, v);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Mx1 vector of data.
<i>v</i>	Nx1 vector, the breakpoints to be used to compute the frequencies - or - scalar, the number of categories.

Remarks

plotAddHistP may only add a histogram to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

See Also

[plotAddBar](#), [plotAddHist](#), [plotAddHistF](#), [plotAddPolar](#), [plotAddXY](#)

plotAddPolar

Purpose

Adds a graph using polar coordinates to an existing polar graph.

Format

```
plotAddPolar(myPlot, radius, theta);  
plotAddPolar(radius, theta);
```

Input

<i>myPlot</i>	A plotControl structure.
---------------	---------------------------------

plotAddScatter

<i>radius</i>	Nx1 or NxM matrix. Each column contains the magnitude for a particular line.
<i>theta</i>	Nx1 or NxM matrix. Each column represents the angle values for a particular line.

Remarks

plotAddPolar may only add curves to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

See Also

[plotAddBar](#), [plotAddHist](#), [plotAddHistF](#), [plotAddHistP](#), [plotAddXY](#)

plotAddScatter

Purpose

Adds a 2-dimensional scatter plot to an existing graph.

Format

```
plotAddScatter(myPlot, x, y);  
plotAddScatter(x, y);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Nx1 or NxM matrix. Each column contains the X values for a particular data point.
<i>y</i>	Nx1 or NxM matrix. Each column contains the Y

values for a particular data point.

Remarks

plotAddScatter may only add a scatter plot to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

See Also

[plotAddBar](#), [plotAddHist](#), [plotAddHistF](#), [plotAddHistP](#), [plotAddScatter](#), [plotAddXY](#)

plotAddShape

Purpose

Adds an arrow, line, ellipse or rectangle to an existing graph.

Format

```
plotAddShape(myAnnotation, which_shape, x_start, y_start, x_end, y_end);
plotAddShape(which_shape, x_start, y_start, x_end, y_end);
```

Input

<i>myAnnotation</i>	Optional argument, an instance of a plotAnnotation structure.
<i>which_shape</i>	String, indicating which shape to create, options include: <ul style="list-style-type: none">• ellipse

2

plotAddShape

	<ul style="list-style-type: none">• line (to which you may add an arrow head)• rectangle
<i>x_start</i>	Scalar or Nx1 vector, the X coordinate for the start of the bounding box for each respective shape.
<i>y_start</i>	Scalar or Nx1 vector, the Y coordinate for the start of the bounding box for each respective shape.
<i>x_end</i>	Scalar or Nx1 vector, the X coordinate for the end of the bounding box for each respective shape.
<i>y_end</i>	Scalar or Nx1 vector, the Y coordinate for the end of the bounding box for each respective shape.

Remarks

plotAddShape will add shapes to existing graphs. It will not create a new graph, however, if one does not already exist. Please also note that the top left corner of the bounding box for the shape will be placed at the coordinates that you specify. The bounding box is rectangular and will therefore not touch the edge of an ellipse at that point.

plotAddShape is not yet supported for surface plots.

Examples

Example: Add a rectangle

```
//Draw simple graph
x = rndu(10, 1);
y = rndu(10, 1);
plotScatter(x, y);

//The rectangle will be drawn between
//third and sixth points on the plot
x_start = x[3];
```



```

y_start = y[3];
x_end = x[6];
y_end = y[6];

//Shape type will be rectangle
annotation_type = "rectangle";

//Add rectangle to graph
plotAddShape(annotation_type, x_start, y_start, x_end, y_end);

```

See Also

[plotAddTextbox](#), [annotationGetDefaults](#)

plotAddSurface

Purpose

Add a surface plot to an existing plot.

Format

```

plotAddSurface(myPlot, x, y, z);
plotAddSurface(x, y, z);

```

Input

<i>myPlot</i>	Optional input, a plotControl structure.
<i>x</i>	1xK vector, the X axis data.
<i>y</i>	Nx1 vector, the Y axis data.
<i>z</i>	NxK matrix, the matrix of height data to be plotted.

plotAddSurface

Example

Example 1: Basic case with default settings

```
new;
cls;

// n: Number of observation
n = 20;

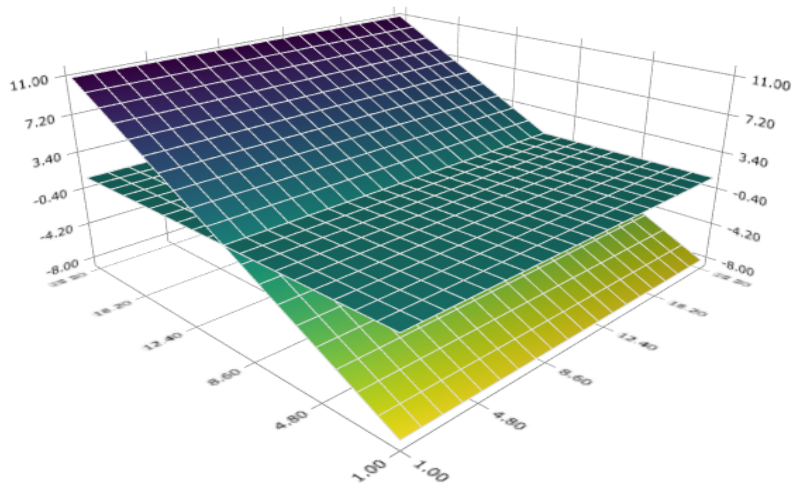
// First Surface
x = seqa(1, 1, n)';
z = ones(n,n);

// Plot surface
plotSurface(x, x', z);

// Second Surface
z2 = reshape(seqa(-8,1,n),n,n);

// Plot the second surface
plotAddSurface(x, x', z2');
```

The plot is



Example 2: With a plotControl structure and color map

```
new;  
cls;  
  
// n: Number of observation  
n = 50;  
  
// x: n * 1 vector from -3 to +3  
x = sega(-3, 6/(n-1), n);  
  
// y: 1 * n row vector  
y = x';
```

plotAddSurface

```
// z: Joint probability of 'x' and 'y'
z = pdfn(x) .* pdfn(y);

// Define plotControl structure
struct plotcontrol myPlot;
myPlot = plotGetDefaults("surface");

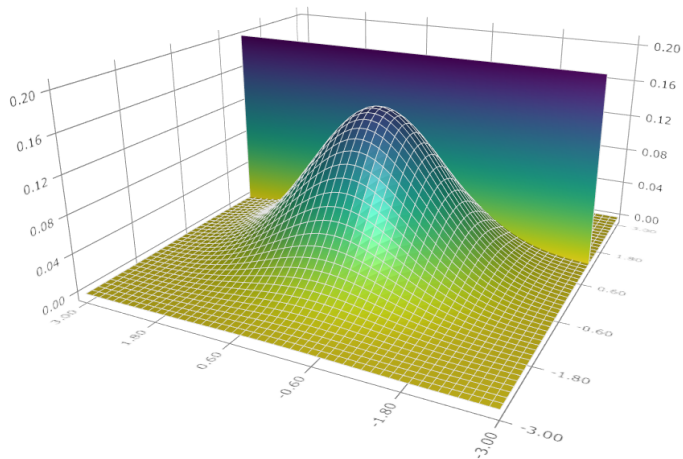
// Set a color map
plotSetColormap(&myPlot, "viridis");

// Plot bivariate normal distribution
plotSurface(myPlot, y, x, z);

// Add a plane with the following 4 corners: (x,y,z)
// (1, -3, 0), (1, 3, 0)
// (1, -3, 0.2), (1, 3, 0.2)
x_plane = 1 ~ 1;
y_plane = -3 | 3;
z_plane = { 0 0.2,
            0 0.2 };

// Add plane
plotAddSurface(x_plane, y_plane, z_plane);
```

The plot is



Remarks

If a `plotControl` structure is passed in to **plotAddSurface**, the only settings that will be checked are:

- The extent of the x, y and z axes (which may be expanded by the `plotControl` structure).
- The instructions for turning the wireframe on or off for the surface being added.

See Also

[plotSurface](#)

plotAddTextbox

plotAddTextbox

Purpose

Adds a textbox to an existing graph.

Format

```
plotAddTextbox(text, x_start, y_start);  
plotAddTextbox(myAnnotation, text, x_start, y_start);
```

Input

<code>myAnnotation</code>	Optional input, a plotAnnotation structure.
<code>text</code>	String, the text to place in the textbox.
<code>x_start</code>	Scalar or Nx1 vector, the X coordinate for the start of the bounding box for each respective text box.
<code>y_start</code>	Scalar or Nx1 vector, the Y coordinate for the start of the bounding box for each respective text box.

Remarks

plotAddTextbox will only add a textbox to an existing graph. It will not create a new graph if one does not already exist. You should also note that the top left corner of the bounding box will be located at the point on the graph that you specify. If the border is turned off, the text will not touch the exact coordinate that you input. In many cases this makes it simpler to label a point that is already part of a scatter or line series without covering it.

Unlike the functions that add data to a plot, if a textbox created by **plotAddTextbox** lies outside of the current bounds of the X and Y axes, the axes will not extend further to provide room for the textbox. This gives you the ability to add text to any part of the scene, or between different subplots.

plotAddTextbox does not currently support surface plots.

Examples

Example 1: Basic textbox

```
//Create text for textbox
box_text = "Periods of recession are highlighted";

x_start = 4;
y_start = 3;

//Add textbox to the (4,3) location on the last draw graph
plotAddTextbox(box_text, x_start, y_start);
```

Example 2: Customized textbox

```
//Simulate and plot simple linear model
b_0 = 2;
b_1 = 1.7;
x = rndn(100, 1);
y = b_0 + b_1 .* x + rndn(100, 1);
plotScatter(x, y);

//Declare instance of plotAnnotation structure
//and fill in with default values
struct plotAnnotation myTextbox;
myTextbox = annotationGetDefaults();

//Set textbox background to 'light gray' with 20% opacity
annotationSetBkd(&myTextbox, "light gray", 0.2);

//Turn off line surrounding textbox by setting thickness to
0px
annotationSetLineThickness(&myTextbox, 0);

//Create text for textbox, using HTML
```

plotAddTextbox

```
box_text = "&alpha; = 2; &beta;<sub>1</sub> = 1.7";

//The top-left corner of the text box
//will be located at the coordinates (0, -1)
x_start = 0;
y_start = -1;

//Add textbox to last draw graph
plotAddTextbox(myTextbox, box_text, x_start, y_start);
```

Example 3: Using a procedure to apply your settings

If you use textboxes often and usually want the same styling, instead of going through the steps above every time you would like to add a text box, you should create a simple procedure to do the set up for you. Here is an example of a procedure that will return a customized **plotAnnotation** structure. You can pass this function in to **plotAddTextbox**.

```
//Add the procedure below to your user library
//and you will only need one line for all the settings
plotAddTextbox(grayTextSettings(), "My customized text
box", 0.15, 0.2);

proc (1) = grayTextSettings();
    struct plotAnnotation mytextbox;

    mytextbox = annotationGetDefaults();
    annotationSetBkd(&mytextbox, "#DDDDDD", 0.3);
    annotationSetFont(&mytextbox, "times", 18, "#555555");
    annotationSetLineThickness(&mytextbox, 2);
    annotationSetLineColor(&mytextbox, "#555555");
    retp(mytextbox);
endp;
```

See Also

[plotAddShape](#), [annotationGetDefaults](#)

plotAddTS

Purpose

Adds a curve of time series data to an existing time series plot.

Format

```
plotAddTS(myPlot, dtstart, frequency, y);  
plotAddTS(dtstart, frequency, y);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>dtstart</i>	Scalar, starting date in DT scalar format.
<i>frequency</i>	Scalar, frequency of the data per year. Valid options include: 1 Yearly 4 Quarterly 12 Monthly
<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular line.

Examples

Example 1

```
//Create some data to plot  
y = rndn(100, 1);  
  
//The first input starts the series in January of 1982  
//The second input specifies the data to be monthly  
plotTS(1982, 12, y);
```

plotAddXY

```
y2 = rndu(28, 1);

//Add the data from 'y2' as quarterly data
//starting in Q2 of 1980
plotAddTS(198004, 4, y2);
```

Remarks

You may only add time series graphs to other time series graphs. For more information on time series graphs, see **Time Series Plots in GAUSS**, Section 1.1.

By default missing values in the *y* variable will be represented as gaps in the line.

See Also

[plotSetXTicLabel](#), [plotSetXTicInterval](#), [plotTS](#)

plotAddXY

Purpose

Adds an XY graph to an existing graph.

Format

```
plotAddXY(myPlot, x, y);
plotAddXY(x, y);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Nx1 or NxM matrix. Each column contains the X values for a particular line.

<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular line.
----------	---

Remarks

plotAddXY may only add curves to 2-D graphs.

This function will not change any of the current graph's settings other than to resize the view as necessary to display the new curve.

See Also

[plotAddBar](#), [plotAddHist](#), [plotAddHistF](#), [plotAddHistP](#), [plotAddPolar](#)

plotArea

Purpose

Creates a stacked area plot.

Format

```
plotArea(myPlot, x, y);
plotArea(x, y);
plotArea(myPlot, x, y, base);
plotArea(x, y, base);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Nx1 matrix. The X values for a particular line.
<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular line. If <i>y</i> contains more than one column, each column will be stacked on top of the

d

plotBar

<i>base</i>	previous column. Scalar optional argument. The height for the base of the area plot. The default value is zero. plotArea does not yet support a vector input for <i>base</i> .
-------------	--

Example

```
x = { 1, 2, 3, 4 };
y = { 1 1.5 0.9,
      0.8 1.2 1.8,
      1 0.7 2,
      1.2 1 1.2 };

//Draw a cumulative area plot of the columns of 'y'
plotArea(x, y);
```

See Also

[plotLogX](#), [plotLogLog](#), [plotScatter](#)

plotBar

Purpose

Generates a bar graph.

Format

```
plotBar(myPlot, labels, height);
plotBar(labels, height);
```

Input

<i>myPlot</i>	A plotControl structure.
---------------	---------------------------------

<i>labels</i>	Nx1 numeric vector, or Nx1 string array containing the bar labels. If scalar 0, a sequence from 1 to rows (<i>height</i>) will be created.
<i>height</i>	NxK numeric vector, bar heights. K overlapping or side-by-side sets of N bars will be graphed.

Example

```
// Create data and labels
labels = "January" $| "June";
temp = { 68, 105 };

// Draw bar graph
plotBar(labels, temp);
```

Remarks

To control the color and texture of the bars as well as whether they are stacked or side by side:

If you are passing a **plotControl** structure to your graph, you may use the function **plotSetBar**.

If you are not passing a **plotControl** structure, these properties are set in the Preferences. To access the Graphics Preferences, select **Tools->Preferences** from the **GAUSS** main menu. Select "Graphics" on the left side of the preferences and then select "Bar" from the list of graph types.

See Also

[plotXY](#), [plotLogX](#), [plotHist](#)

plotBox

plotBox

Purpose

Graphs data using the box graph percentile method.

Format

```
plotBox(myPlot, group_id, y);  
plotBox(group_id, y);
```

Input

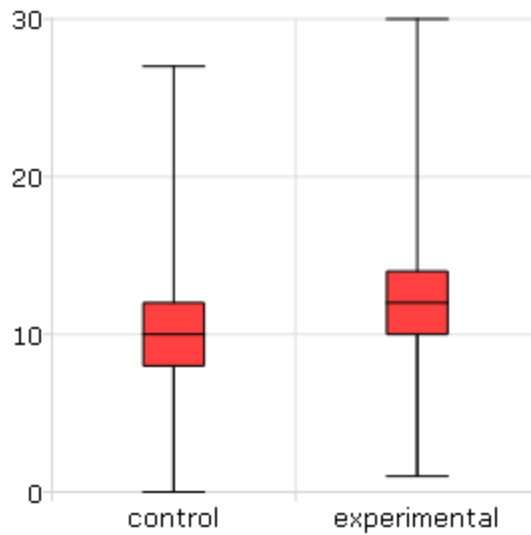
<i>myPlot</i>	A plotControl structure.
<i>group_id</i>	Mx1 vector for numeric labels or Mx1 string array for text labels. This contains the group numbers or string labels corresponding to each column of <i>y</i> data. If scalar 0, a sequence from 1 to cols (<i>y</i>) will be generated automatically for the X axis.
<i>y</i>	NxM matrix. Each column represents the set of <i>y</i> values for an individual percentiles box symbol.

Examples

Example 1: Using string labels

```
//Create two columns of random data  
nobs = 1e5;  
y_1 = rndPoisson(nobs, 1, 10);  
y_2 = rndPoisson(nobs, 1, 12);  
  
//Create a 2x1 string array, using the  
//horizontal string concatenation operator  
labels = "control" $| "experimental";
```

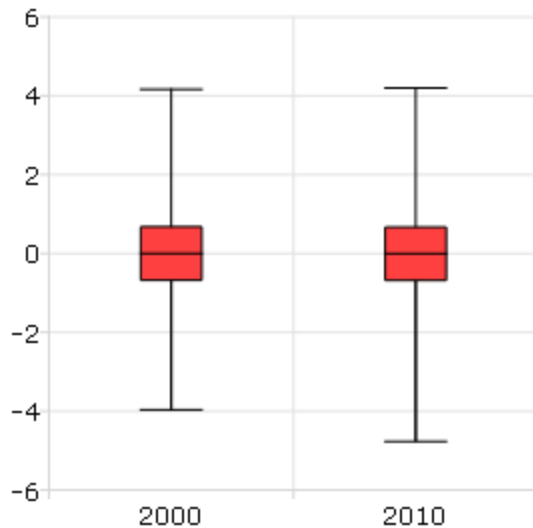
```
//Draw the two boxes  
plotBox(labels, y_1~y_2);
```



Example 2: Using numeric labels

```
//Create two columns of random data  
nobs = 1e5;  
y_1 = rndn(nobs, 1);  
y_2 = rndn(nobs, 1);  
  
//Create a 2x1 vector of numeric labels  
labels = { 2000, 2010 };  
  
//Draw the two boxes  
plotBox(labels, y_1~y_2);
```

plotCDFEmpirical



Remarks

Note that numeric labels added to the X-axis are only labels for the corresponding boxes. They do not imply any order or denote a particular X-axis location.

If missing values are encountered in the y data, they will be ignored during calculations and will not be plotted.

See Also

[plotHistP](#), [plotScatter](#)

plotCDFEmpirical

Purpose

Plots the cumulative distribution function (cdf) of the empirical distribution.

Format

```
plotCDFEmpirical(myPlot, x);  
plotCDFEmpirical(x);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	N x 1 vector.

Example

```
new ;  
cls ;  
rndseed 2223;  
  
// Create a random vector  
x = rndn(30,1);  
  
// Sort x for the first column  
x = sortc(x,1);  
  
// Get empirical cdf of x  
f = cdfEmpirical(x);  
  
// Add negative infinity (___INFN) for probability equal to  
0.  
print (___INFN|x)~f;  
  
// Plot empirical distribution  
plotCDFEmpirical(x);  
  
// Get normal cdf of x  
f2 = cdfN(x);
```

plotCDFEmpirical

```
// Plot theoretical distribution
plotADDXY(x, f2);
```

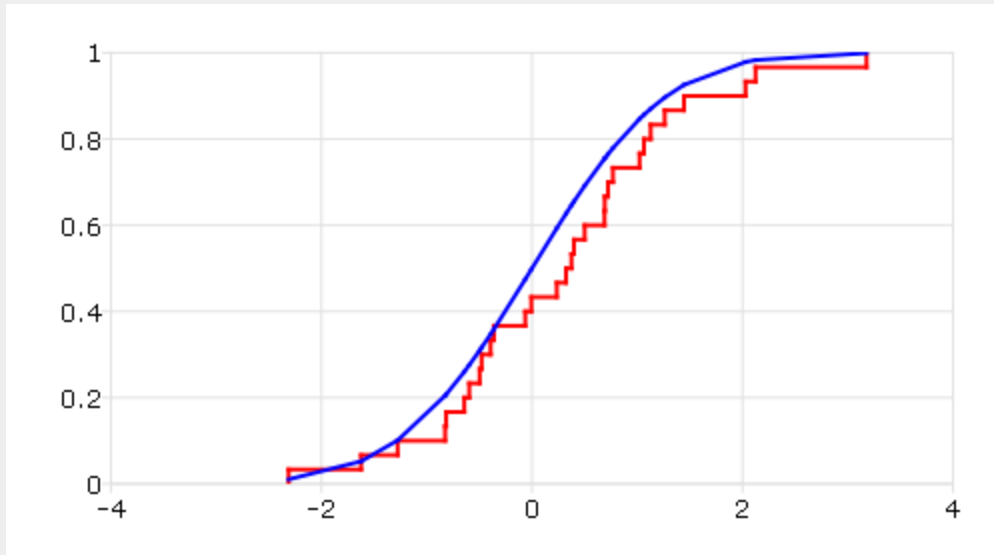
After above code,

-INF	0.00000000
-2.3124206	0.03333333
-1.6240227	0.06666667
-1.2763153	0.10000000
-0.82532512	0.13333333
-0.81574278	0.16666667
-0.64338729	0.20000000
-0.59625173	0.23333333
-0.49725006	0.26666667
-0.47855430	0.30000000
-0.39340284	0.33333333
-0.36201638	0.36666667
-0.063830011	0.40000000
-0.0064523646	0.43333333
0.23570074	0.46666667
0.32355136	0.50000000
0.37501508	0.53333333
0.39847826	0.56666667
0.50039685	0.60000000
0.68900341	0.63333333
0.69132515	0.66666667
0.72246796	0.70000000
0.76893134	0.73333333
1.0221019	0.76666667
1.0638924	0.80000000
1.1274880	0.83333333
1.2610791	0.86666667
1.4445086	0.90000000
2.0295113	0.93333333

p

2.1240430	0.96666667
3.1784008	1.0000000

The Plot is



See Also

[cdfEmpirical](#)

plotClearLayout

Purpose

Clears any previously set plot layouts.

Format

```
plotClearLayout();
```

plotClearLayout

Example

```
//Create a 1x2 Plot Layout and insert a percentage
//histogram of some random normal numbers in the first
//cell.
plotLayout(1, 2, 1);
plotHistP(rndn(1000, 1), 30);

//Insert gamma distributed random numbers into the second
//cell.
plotLayout(1, 2, 2);
plotHistP(rndGamma(1000, 1, 3, 2), 30);

//Display the image for 2 seconds
pause(2);

//Clear the 1x2 layout
plotClearLayout();

//Plot percentage histogram of beta distributed random
//numbers. This graph will take up the entire plot window
//since the 1x2 plot layout has been cleared.
plotHistP(rndBeta(1000, 1, 2, 1), 30);
```

Remarks

After calling this function all subsequent graphs will be drawn to fill the entire graph window.

See Also

[plotSetBar](#), [plotBar](#), [plotLayout](#), [plotCustomLayout](#)

plotContour

Purpose

Graphs a matrix of contour data.

Format

```
plotContour(myPlot, x, y, z);
plotContour(x, y, z);
```

Input

<i>myPlot</i>	Optional input: plotControl structure.
<i>x</i>	1xK vector, the X axis data.
<i>y</i>	Nx1 vector, the Y axis data.
<i>z</i>	NxK matrix, the matrix of height data to be plotted.

Examples

```
//Clear out variables in GAUSS workspace
new;

//Create contour data
x = seqa(-4,.125,161)';
y = seqa(-8,.125,161);
z = sin(x) .* cos(y) * .5;
z = z .* sin(x/3) .* cos(y/3);
z = z .* sin(x/5) + sin(y/2.5)/3 + sin(x/2.5)/3;

//Set up control structure with defaults
//for surface plots
struct plotControl myPlot;
```

plotContour

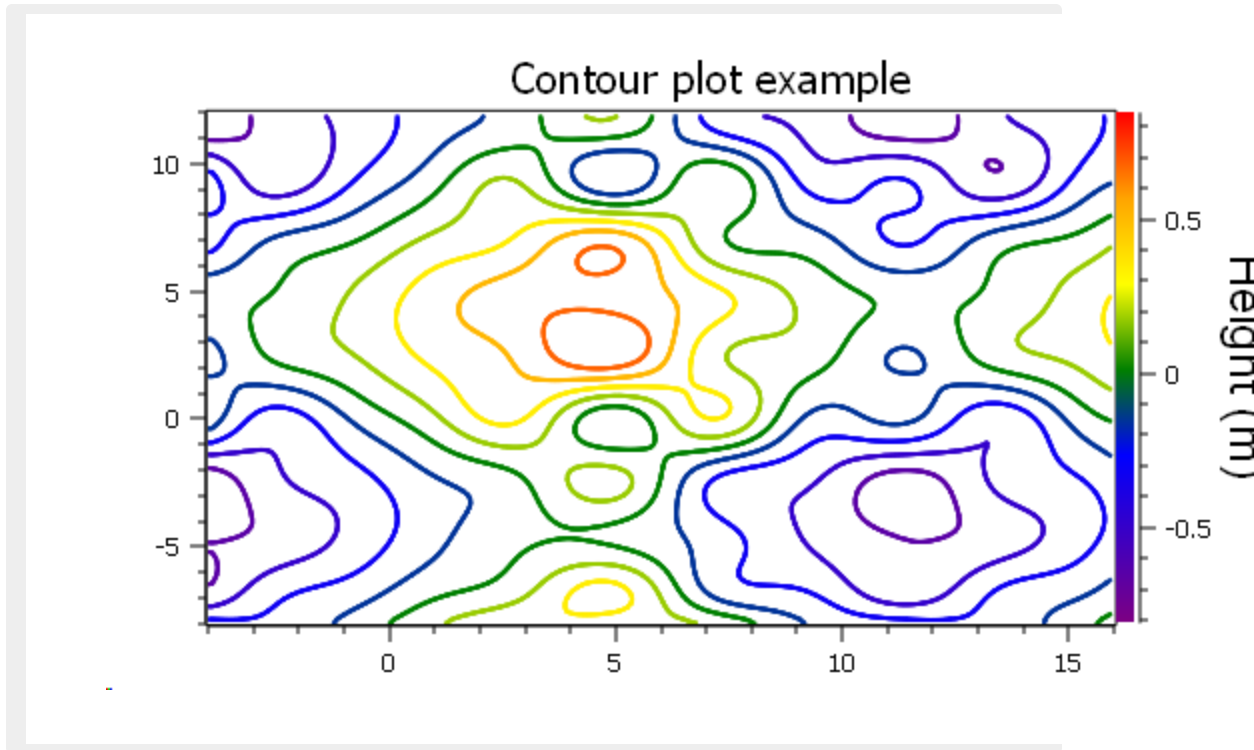
```
myPlot = plotGetDefaults("surface");

//Set title and Z axis label
plotSetTitle(&myPlot, "Contour plot example", "Courier
bold", 16, "black");
plotSetZLabel(&myPlot, "Height (m)", "Arial", 18);

//Turn off X and Y axis labels
plotSetXLabel(&myPlot, "");
plotSetYLabel(&myPlot, "");

//Set contour level colors
string rainbow = { "Red", "Orange", "Yellow", "Green",
"Blue", "Purple" };
plotSetLineColor(&myPlot, rainbow);

//Draw graph using plotcontrol structure
plotContour(myPlot, x, y, z);
```



Remarks

A vector of evenly spaced contour levels will be generated automatically from the `z` matrix data. Each contour level will be labeled. For unlabeled contours, use **`zticks`**.

To specify a vector of your own unequal contour levels, set the vector `_plev` before calling **`contour`**.

To specify your own evenly spaced contour levels, see **`zticks`**.

See Also

[plotSurface](#)

plotCustomLayout

plotCustomLayout

Purpose

Plots a graph of user-specified size at a user-specified location.

Format

```
plotCustomLayout(xStart, yStart, width, height);
```

Input

<i>xStart</i>	scalar, the distance from the left edge of the canvas to the left edge of the custom plot expressed as a number between 0 and 1.
<i>yStart</i>	scalar, the distance from the bottom edge of the canvas to the bottom edge of the custom plot expressed as a number between 0 and 1.
<i>width</i>	scalar, the width of the custom plot expressed as a number between 0 and 1.
<i>height</i>	scalar, the height of the custom plot expressed as a number between 0 and 1.

Example

```
//Create an additive sequence starting from -pi and moving
//forward in 0.1 increments
x = seqa(-pi, 0.1, 63);

//Plot the cosine of x
plotXY(x, cos(x));

//Create a custom section for the next graph starting 10%
```



```
//from the main graph's left edge, 10% from the bottom of
//the main graph, with a width and height both equalling
//30% of the width of the main graph.
plotCustomLayout(0.1, 0.1, 0.3, 0.3);

//Plot the next graph in the custom layout
plotXY(x[1:20], cos(x[1:20] ) ));

//Prevent the next graph from being drawn in this custom
//region
plotClearLayout();
```

Remarks

After calling this function all subsequent graphs will be plotted inside of the specified custom layout until the layout is reset with **plotLayout**, or the layout is cleared with **plotClearLayout**.

See Also

[plotSetBar](#), [plotBar](#), [plotHistP](#), [plotGetDefaults](#)

plotGetDefaults

Purpose

Gets default settings for plotting graphs.

Format

```
myPlot = plotGetDefaults(graph);
```

Input

<i>graph</i>	String, name of graph type: bar, box, hist, polar, scatter,
--------------	---

plotGetDefaults

surface or xy.

Output

myPlot A **plotControl** structure.

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure with defaults for an
//'xy' graph
myPlot = plotGetDefaults ("xy");

//Create some data to plot
x = seqa (-5, 0.1, 50);
y = pdfn (x);

//Make a desired change to the plotControl structure
plotSetTitle (&myPlot, "Default XY Settings");

//Plot the data using the plotControl structure
plotXY (myPlot, x, y);
```

Remarks

The **plotGetDefaults** function will use the default settings for the specified graph type. These may be accessed from the main menu bar: **Tools->Preferences->Graphics**.

See Also

[plotSetBkdColor](#), [plotSetLineColor](#), [plotSetLineSymbol](#)

plotHist

Purpose

Computes and graphs a frequency histogram for a vector. The actual frequencies are plotted for each category.

Format

```
plotHist(myPlot, x, v);  
plotHist(x, v);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Mx1 vector of data.
<i>v</i>	Nx1 vector, the breakpoints to be used to compute the frequencies - or - scalar, the number of categories

Example

```
//Create some data to plot  
x = randn(5000, 1);  
  
//Plot the data  
plotHist(x, 20);
```

See Also

[plotHistP](#), [plotHistF](#), [plotBar](#)

plotHistF

plotHistF

Purpose

Graphs a histogram given a vector of frequency counts.

Format

```
plotHistF(myPlot, f, c);  
plotHistF(f, c);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>f</i>	Nx1 vector, frequencies to be graphed.
<i>c</i>	Nx1 vector, numeric labels for categories. If this is a scalar 0, a sequence from 1 to rows (<i>f</i>) will be created.

Remarks

The axes are not automatically labeled. Use the functions **plotSetXLabel** and **plotSetYLabel**.

See Also

[plotHist](#), [plotBar](#), [plotSetXLabel](#)

plotHistP

Purpose

Computes and graphs a percent frequency histogram of a vector. The percentages in each category are plotted.

Format

```
plotHistP(myPlot, x, v);
plotHistP(x, v);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Mx1 vector of data.
<i>v</i>	Nx1 vector, the breakpoints to be used to compute the frequencies - or - scalar, the number of categories.

See Also

[plotHist](#), [plotHistF](#), [plotBar](#), [plotBox](#), [plotScatter](#)

plotLayout

Purpose

Divides a plot into a grid of subplots and assigns the cell location in which to draw the next created graph.

Format

```
plotLayout(gRows, gCols, ind);
```

Input

<i>gRows</i>	scalar, number of rows of the graph layout.
<i>gCols</i>	scalar, number of columns of the graph layout.

plotLayout

<i>ind</i>	scalar, cell location in which to place the next created graph.
------------	---

Example

```
//Create 10x4 matrix where each column is an additive
//sequence from 0.1 to 1.0
x = sega(0.1, 0.1, 10);
y = ones(10, 4).*x;

//Apply a function to each column of 'y'
y[:,1] = cos(x);
y[:,2] = sin(x);
y[:,3] = cdfn(x);
y[:,4] = exp(x);

for i(1, 4, 1);
    //Divide plot canvas into a 2x2 grid of subplot
    //locations and place each newly created graph in the
    //next available cell location.
    plotLayout(2, 2, i);

    //Plot each column of y in a separate subplot window.
    plotXY(x, y[:,i]);
endfor;

//Clear the layout so the next plot will not be inside this
//layout
plotClearLayout();
```

Remarks

After calling this function all subsequent graphs will be plotted inside of the specified layout until the layout is reset with **plotLayout**, or the layout is cleared with

`plotClearLayout.`

See Also

[plotBar](#), [plotClearLayout](#), [plotCustomLayout](#), [plotHist](#)

plotLogLog

Purpose

Graphs X vs. Y using log coordinates.

Format

```
plotLogLog(myPlot, x, y);
plotLogLog(x, y);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Nx1 or NxM matrix. Each column contains the X values for a particular line.
<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular line.

See Also

[plotXY](#), [plotLogX](#), [plotLogY](#)

plotLogX

Purpose

Graphs X vs. Y using log coordinates for the X axis.

plotLogY

Format

```
plotLogX(myPlot, x, y);  
plotLogX(x, y);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Nx1 or NxM matrix. Each column contains the X values for a particular line.
<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular line.

See Also

[plotXY](#), [plotLogY](#), [plotLogLog](#)

plotLogY

Purpose

Graphs X vs. Y using log coordinates for the Y axis.

Format

```
plotLogY(myPlot, x, y);  
plotLogY(x, y);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Nx1 or NxM matrix. Each column represents the X values for a particular line.

y	Nx1 or NxM matrix. Each column represents the Y values for a particular line.
-----	---

See Also

[plotXY](#), [plotLogX](#), [plotLogLog](#)

plotOpenWindow

Purpose

Opens a new, empty graphic window to be used by the next drawn graph.

Format

```
plotOpenWindow();
```

Example

```
//Create data
x = rndn(10000, 1);
x2 = rndn(10000, 1);
x3 = rndn(10000, 1);

//Plot first vector as a percentage histogram with 30 bins
plotHistP(x, 30);

//Plot second vector, drawing over the previously created
//graph.
plotHistP(x2, 30);

//Create a new graphic window and plot the second vector as
//a percentage histogram with 30 bins inside this new
//window.
plotOpenWindow();
```

plotPolar

```
//Draw the graph
plotHistP(x3, 30);
```

Remarks

To automatically open each new graph in a new graph window, use **plotSetNewWindow** or set the preference in the main applicaton menu. This may be found by selecting **Tools->Preferences** and then clicking on **Graphics** on the left side of the preferences window.

If you select the radio button next to "New Window" at the top of the graphics preferences window, each new graph will be automatically drawn in a new graphics window.

See Also

[plotSave](#), [plotCustomLayout](#), [plotSetLegend](#), [plotSetNewWindow](#)

plotPolar

Purpose

Graph data using polar coordinates.

Format

```
plotPolar(myPlot, radius, theta);
plotPolar(radius, theta);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>radius</i>	Nx1 or NxM matrix. Each column contains the

<i>theta</i>	magnitude for a particular line. Nx1 or NxM matrix. Each column represents the angle values for a particular line.
--------------	---

See Also

[plotXY](#), [plotLogX](#), [plotLayout](#), [plotSetXLabel](#)

plotSave

Purpose

Saves the last created graph to a user specified file type.

Format

```
plotSave(filename);
plotSave(filename, size);
plotSave(filename, size, units);
plotSave(filename, size, units, dpi);
```

Input

<i>filename</i>	String, name of the file to create with a file type extension. Available file extensions include: .jpg, .plot, .png, .pdf, .svg, .tiff (NOTE: Available file types may vary per system. A list of valid types can be found in the File->Export Graph file dialog window).
<i>size</i>	2x1 vector, dimensions of the saved graph in specified units. Default unit is centimeters. <i>size</i> is an optional input when saving a .plot file, but is required for all other file types.

plotSave

unit

Optional input, String, type of units dimension is specified in. This value is ignored if the filename extension is '.plot'. Valid options include:

- "cm" Centimeters (Default)
- "mm" Millimeters
- "in" Inches
- "px" Pixels

dpi

Optional input, scalar, requested dots per inch when saving file. Defaults to current system dpi. This value is ignored if the filename extension is '.plot'. dpi determines the number of pixels rendered when saving a file in terms of physical dimensions (cm, mm, in). Specifying the dpi parameter has no effect if the specified units are pixels (px).

e.g. if a printing requirement demanded 11"x8.5" (landscape) with 300 dpi then the plot could be made to fit those dimensions exactly with the line:

```
plotSave("file.pdf", 11|8.5, "in",  
300);
```

which would create an output of 3300x2550 pixels with the PDF page size set in the specified physical dimensions.

Examples

Example 1: Basic save in GAUSS .plot format

```
//Create data
x = sega(0.1, 0.1, 10);
y = cos(x);

//Plot the data
plotXY(x, y);

//Save the graph as a GAUSS .plot file
plotSave("mygraph.plot");
```

Example 2: Save as 640x480 PNG

```
//Create data
x = sega(0.1, 0.1, 10);
y = cos(x);

//Plot the data
plotXY(x, y);

//Save the graph as a 640 wide by 480 tall PNG file
plotSave("mygraph.png", 640 | 480, "px");
```

Example 3: Save as 11x8.5 inch PDF at 300 DPI

```
//Create data
x = sega(0.1, 0.1, 10);
y = cos(x);

//Plot the data
plotXY(x, y);

plotSave("mygraph.png", 11 | 8.5, "in", 300);
```

plotScatter

Technical Notes

The `.plot` file extension is a JSON file that is the native format used by **GAUSS** to save graphs.

Remarks

The font sizes in the graph will not be scaled with the size change. So make sure to set the font sizes to the correct size for the final graph dimensions.

See Also

[plotCustomLayout](#), [plotSetLegend](#)

plotScatter

Purpose

Creates a 2-dimensional scatter plot.

Format

```
plotScatter(myPlot, x, y);  
plotScatter(x, y);
```

Input

p	<i>myPlot</i>	A plotControl structure.
	<i>x</i>	Nx1 or NxM matrix. Each column contains the X values for a particular data point.
	<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular data point.

Example

```
//Create random normal data
x = rndn(50, 1);

//Reverse the order of 'x' and set it to be the 'y' value
y = rev(x);

//Plot the data
plotScatter(x, y);
```

See Also

[plotXY](#), [plotLogLog](#), [plotBox](#), [plotHistP](#)

plotSetAxesPen

Purpose

Sets the color for the axes line.

Format

```
plotSetAxesPen(&myPlot, thickness);
plotSetAxesPen(&myPlot, thickness, clr);
```

Input

<i>&myPlot</i>	A plotControl structure pointer
<i>thickness</i>	Scalar, the thickness of the axis line in pixels
<i>clr</i>	String, name or rgb value of the new color for the axes

plotSetBar

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("xy");

//Set axis to be 2 pixels wide and black
plotSetAxesPen(&myPlot, 2, "black");

//Create data
x = seqa(0.1, 0.1, 50);
y = sin(x)~cos(x);

//Plot the data with the new line colors
plotXY(myPlot, x, y);
```

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS, CHAPTER 1**, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetLineSymbol](#)

plotSetBar

Purpose

Sets the fill style and format of bars in a histogram or bar graph.

Format

```
plotSetBar(&myPlot, fillType, barStacked);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>fillType</i>	Nx1 vector, where N is the number of bar styles to set.
	0 Solid, beveled edge
	1 Solid
	2 Dense 1
	3 Dense 2
	4 Dense 3
	5 Dense 4
	6 Dense 5
	7 Dense 6
	8 Horizontal lines
	9 Vertical lines
	10 Cross pattern
	11 B diagonal pattern
	12 F diagonal pattern
	13 Diagonal Cross
<i>barStacked</i>	Scalar, 1 for stacked or 0 for side-by-side bars.

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
```

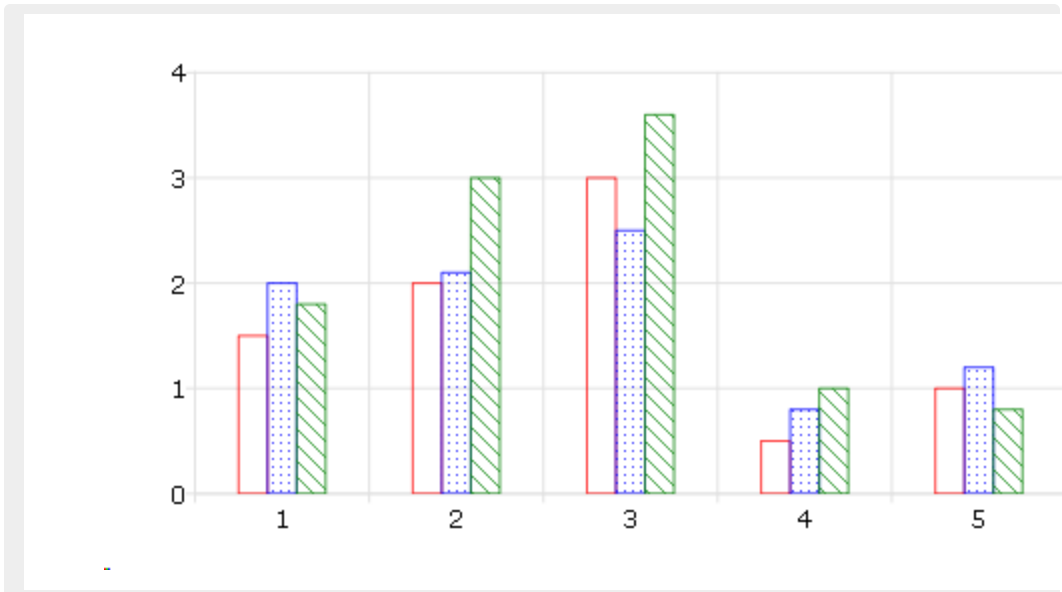
plotSetBar

```
myPlot = plotGetDefaults("bar");

//Set the first set of bars to have a solid-fill, the
//second set to have a fill of horizontal lines, the third
//to have a diagonal cross fill and set the bars to be
//side-by-side.
textures = { 0, 8, 13 };
plotSetBar(&myPlot, textures, 0);

//Create data
x = seqa(1, 1, 5);
y = {
1.5      2      1.8,
2        2.1    3,
3        2.5    3.6,
0.5      0.8    1,
1        1.2    0.8
};

//Draw bar graph
plotBar(myPlot, x, y);
```



Remarks

When graphing without the use of a **plotControl** structure, these settings may be chosen through the **Tools->Preferences->Graphics** menu, after selecting the Bar radio button. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotBar](#), [plotGetDefaults](#), [plotHist](#)

plotSetBkdColor

Purpose

Sets the background color of a graph.

plotSetBkdColor

Format

```
plotSetBkdColor(&myPlot, color);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>color</i>	String, name or rgb value of the new color.

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("polar");

//Set new background color to light grey
plotSetBkdColor(&myPlot, "light grey");

//Create data
x = seqa(0.1, 0.1, 200);
y = x;

//Create a polar plot of the data with the new background
//color
plotPolar(myPlot, x, y);
```

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetLineColor](#), [plotSetLineStyle](#)

plotSetColorMap

Purpose

Sets the color maps for a surface or contour plot.

Format

```
plotSetColorMap(&myPlot, color_type);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>color_type</i>	String, name of color maps: "viridis", "magma", "inferno", "plasma".

Example

```
//Clear out variables in GAUSS workspace
new;

//Create data
x = seqa(-4,.125,161)';
y = seqa(-8,.125,161);
z = sin(x) .* cos(y) * .5;
z = z .* sin(x/3) .* cos(y/3);
z = z .* sin(x/5) + sin(y/2.5)/3 + sin(x/2.5)/3;

//Set up control structure with defaults
//for surface plots
```

plotSetColorMap

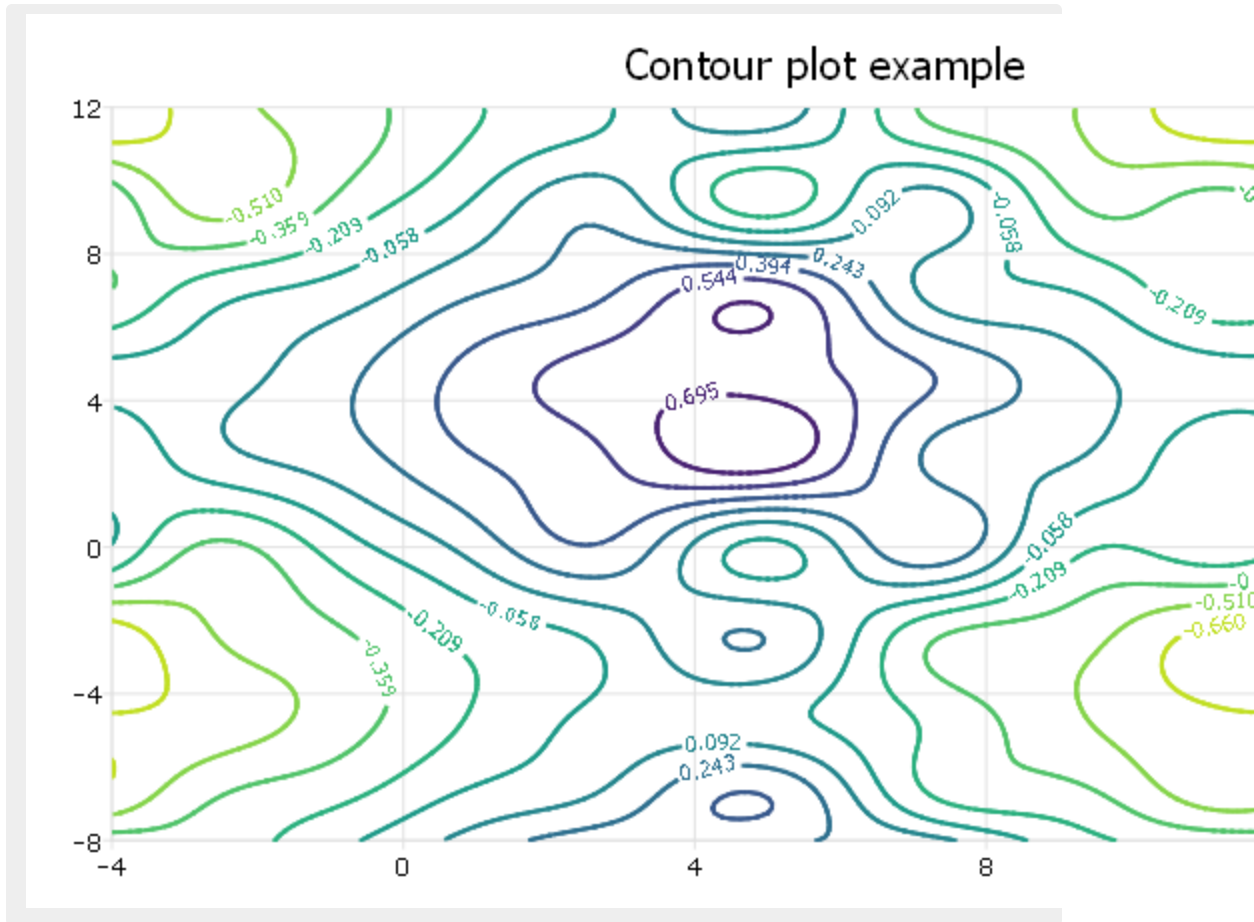
```
struct plotControl myPlot;
myPlot = plotGetDefaults("surface");

//Set title and Z axis label
plotSetTitle(&myPlot, "Contour plot example", "arial", 16,
"black");

//Set color map for contour
plotSetColorMap(&myplot, "viridis");

//Draw graph using plotcontrol structure
plotContour(myPlot, x, y, z);
```

The Plot is



Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetContourLabels](#), [plotSetZLevels](#)

plotSetContourLabels

plotSetContourLabels

Purpose

Sets the contour label for a graph.

Format

```
plotSetContourLabels(&myPlot, show_labels);  
plotSetContourLabels(&myPlot, show_labels, label_format);  
plotSetContourLabels(&myPlot, show_labels, label_format,  
label_precision);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>show_labels</i>	Scalar, 0 or 1. The flag of show labels on contours.
<i>label_format</i>	String, format's option: "D", e.g. 1.234567; "E", e.g. 1.23E1; "G". "G" is either "D" or "E", whichever is more compact
<i>label_precision</i>	Scalar, precision of contour label.

Example

```
//Clear out variables in GAUSS workspace  
new;
```



```
//Create data
x = seqa(-4,.125,161)';
y = seqa(-8,.125,161);
z = sin(x) .* cos(y) * .5;
z = z .* sin(x/3) .* cos(y/3);
z = z .* sin(x/5) + sin(y/2.5)/3 + sin(x/2.5)/3;

//Set up control structure with defaults
//for surface plots
struct plotControl myPlot;
myPlot = plotGetDefaults("surface");

//Set title and Z axis label
plotSetTitle(&myPlot, "Contour plot example", "arial", 16,
"black");

//Set color map for contour
plotSetColorMap(&myplot, "viridis");

//Set up flag of show label for contour
showLabels = 1;

//Set up format for contour label
label_format = "G";

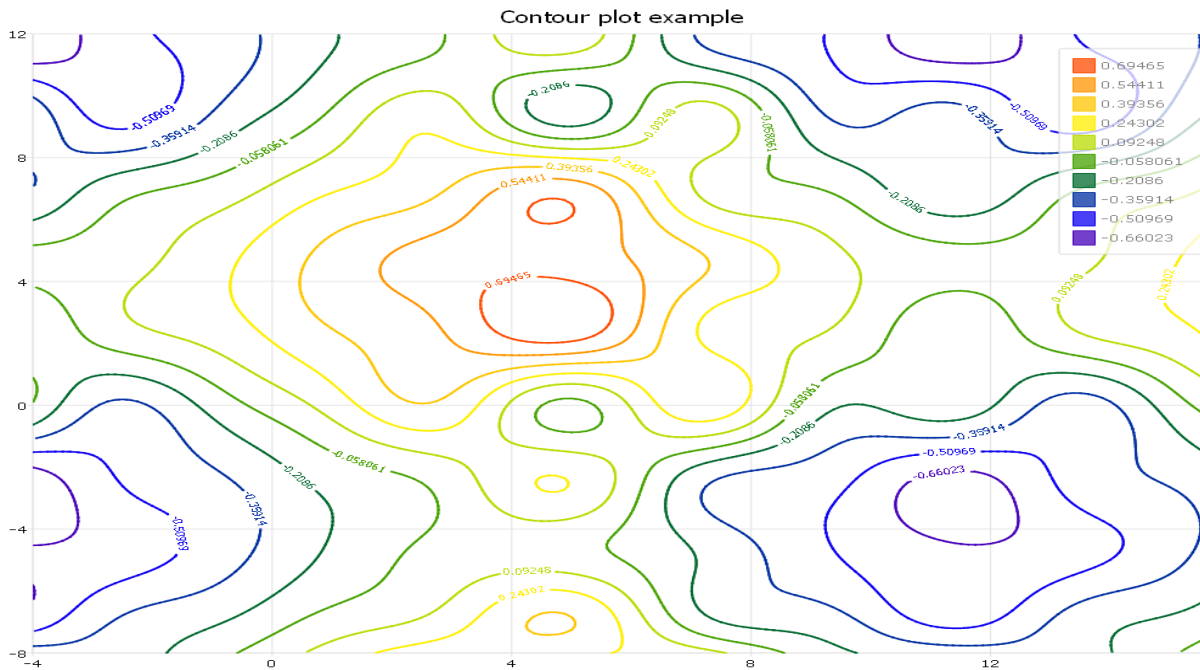
//Set up precision for contour label
label_precision = 5;

//Use function to set up contour labels
plotSetContourLabels(&myplot, showlabels, label_format,
label_precision);

//Draw graph using plotcontrol structure
plotContour(myPlot, x, y, z);
```

plotSetContourLabels

The Plot is



Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetColorMap](#), [plotSetZLevels](#)

plotSetFill

Purpose

Sets the fill style, transparency and color for area plots, histograms and bar graphs.

Format

```
plotSetFill(&myPlot, fillType, transparency_pct, colors);
plotSetFill(&myPlot, fillType, transparency_pct);
plotSetFill(&myPlot, fillType);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>fillType</i>	Nx1 vector, where N is the number of bar styles to set.
0	No fill
1	Solid
2	Dense 1
3	Dense 2
4	Dense 3
5	Dense 4
6	Dense 5
7	Dense 6
8	Horizontal lines
9	Vertical lines
10	Cross pattern
11	B diagonal pattern

plotSetFill

	12	F diagonal pattern
	13	Diagonal Cross
<i>transparency_</i> <i>pct</i>	Scalar, between 0 and 1. The percent opacity of the fill.	
<i>colors</i>	String array, color names or HTML hex value colors.	

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("bar");

//Set all bars to have a solid, blue, fill, with 50% opa-
city
textures = 1;
plotSetFill(&myPlot, textures, 0.5, "blue");

//Create data
x = seqa(1, 1, 5);
y = { 1.5, 2, 3, 0.5, 1 };

//Draw bar graph
plotBar(myPlot, x, y);
```

Remarks

When graphing without the use of a **plotControl** structure, these settings may be chosen through the **Tools->Preferences->Graphics** menu, after selecting the Bar radio button. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotBar](#), [plotGetDefaults](#), [plotHist](#)

plotSetGrid

Purpose

Controls the settings for the background grid of a plot.

Format

```
plotSetGrid(&myPlot, ticStyle, color);  
plotSetGrid(&myPlot, ticStyle);  
plotSetGrid(&myPlot, onOff);
```

Input

<i>ticStyle</i>	String, specifies whether grid marks should be drawn on major tic marks. Options: "major"
<i>color</i>	String, name or rgb value of the new color.
<i>onOff</i>	String, turns the grid on or off. Options: "on" or "off." If used, this must be the only argument passed to the function besides the plotControl structure pointer.

Example

```
//Declare plotControl structure  
struct plotControl myPlot;  
  
//Initialize plotControl structure  
myPlot = plotGetDefaults("scatter");  
  
//Set grid to be black and on the major tics only
```

plotSetLegend

```
plotSetGrid(&myPlot, "major", "black");

//Create a scatter plot of random data
plotScatter(myPlot, seqa(1, 1, 10 ), rndn(10, 1));

//Turn off the grid
plotSetGrid(&myPlot, "off");
```

See Also

[plotCustomLayout](#), [plotSetTitle](#)

plotSetLegend

Purpose

Adds a legend to a graph.

Format

```
plotSetLegend(&myPlot, label, location, orientation);
plotSetLegend(&myPlot, label, location);
plotSetLegend(&myPlot, label);
plotSetLegend(&myPlot, turn_off);
```

Input

p

<i>&myPlot</i>	A plotControl structure pointer.
<i>label</i>	String array, names of the line labels.
<i>location</i>	String, the location to place the legend. The location string may contain up to three tokens, or words. <ol style="list-style-type: none">1. Vertical location: top (default), vcenter or bottom.

(Note: for backwards compatibility middle may still be used for vcenter. However, new programs should use vcenter).

2. Horizontal location: left, hcenter or right (default). (Note: for backwards compatibility center may still be used for hcenter. However, new programs should use hcenter.

3. Inside/Outside location: inside (default), below or outside.

<i>orientation</i>	scalar, 0 for a horizontal legend or 1 for a vertical legend.
<i>turn_off</i>	string, "off" will disable the legend.

Technical Notes

- The location parameter is a string with up to three tokens or words that are separated by a space. For example,

```
location = "top right";
location = "right top";
location = "inside top right";
```

- Use **plotSetLegendFont** to control the legend font family, size and color.
- See **plotSetTextInterpreter**, for instructions on using Latex, or HTML in the legend labels.

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
```

plotSetLegendFont

```
myPlot = plotGetDefaults("scatter");

//Set labels, location, and orientation of legend
label = "sample A"|"sample B";
location = "top right";
orientation = 0;
plotSetLegend(&myPlot, label, location, orientation);

//Create data
x = rndn(30, 2);
y = rndn(30, 2);

//Plot the data with the legend settings
plotScatter(myplot, x, y);
```

See Also

[plotSetLegendFont](#), [plotSetTextInterpreter](#)

plotSetLegendFont

Purpose

Controls the settings for the legend on a graph.

Format

```
plotSetLegendFont(&myPlot, font);
plotSetLegendFont(&myPlot, font, font_size);
plotSetLegendFont(&myPlot, font, font_size, font_color);
```

Input

<i>myPlot</i>	A plotControl structure pointer.
---------------	---

<i>font</i>	String, font or font family name.
<i>font_size</i>	Scalar, font size in points.
<i>font_color</i>	String, named color or RGB value.

Example

```
new;
cls;

//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("scatter");

//Set labels, location, and orientation of legend
label = "sample A"|"sample B";
location = "top right";
orientation = 0;
plotSetLegend(&myPlot, label, location, orientation);

//Set font of legend
plotSetLegendFont(&myPlot, "arial", 18, "dark grey");

//Create data
x = randn(30, 2);
y = randn(30, 2);

//Plot the data with the legend settings
plotScatter(myplot, x, y);
```

See Also

[plotSetLegend](#)

plotSetLineColor

plotSetLineColor

Purpose

Sets the line colors for a graph.

Format

```
plotSetLineColor(&myPlot, colors);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>colors</i>	String array, name or rgb value of the new colors.

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("xy");

//Set new line colors to aqua and midnight blue
clrs = "aqua"$|"midnight blue";
plotSetLineColor(&myPlot, clrs);

//Create data
x = seqa(0.1, 1, 50);
y = sin(x)~cos(x);

//Plot the data with the new line colors
plotXY(myPlot, x, y);
```

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetLineStyleSymbol](#)

plotSetLineStyle

Purpose

Sets the line styles for a graph.

Format

```
plotSetLineStyle(&myPlot, newStyle);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>newStyle</i>	Matrix, new line styles. Options include:
1	Solid line.
2	Dash line.
3	Dot line.
4	Dash-Dot line.
5	Dash-Dot-Dot line.

plotSetLineStyle

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("xy");

//Set line 1 as a solid line,
//set line 2 as a dash line, etc.
newStyle = { 1, 2, 3, 4, 5 };
plotSetLineStyle(&myPlot, newStyle);

//Create data
x = seqa(0.1, 1, 50);
y = sin(x)~cos(x);

//Plot the data with the new line styles
plotXY(myPlot, x, y);
```

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetTitle](#), [plotSetLineStyle](#)

plotSetLineStyle

Purpose

Sets the symbols displayed on the plotted points of a graph.

Format

```
plotSetLineSymbol(&myPlot, newSymbol, symbolWidth);
plotSetLineSymbol(&myPlot, newSymbol);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>newSymbol</i>	Matrix, new line symbol settings. Options include: <ul style="list-style-type: none"> -1 None. 0 Ellipse. 1 Rectangle. 2 Diamond. 3 Upward pointing triangle. 4 Downward pointing triangle. 5 Triangle. 6 Leftward pointing triangle. 7 Rightward pointing triangle. 8 Cross. 9 Diagonal cross. 10 Horizontal line. 11 Vertical line. 12 Star 1. 13 Star 2. 14 Hexagon.
<i>symbolWidth</i>	Scalar, width to draw line symbols.

plotSetLineSymbol

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("xy");

//Set line 1 to have no symbol
//Set line 2 to display an ellipse at each plotted point.
newSymbol = { -1, 0 };
symbolWidth = 5;
plotSetLineSymbol(&myPlot, newSymbol, symbolWidth);

//Create data
x = seqa(0.1, 0.1, 50);
y = sin(x)~cos(x);

//Plot the data with the new line symbols
plotXY(myPlot, x, y);
```

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetXLabel](#), [plotSetLineColor](#)

plotSetLineThickness

Purpose

Sets the thickness of the lines on a graph.

Format

```
plotSetLineThickness(&myPlot, newTh);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>newTh</i>	1 x N matrix, new line thickness settings.

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("xy");

//Set all lines to have a thickness of 2
newTh = 2;
plotSetLineThickness(&myPlot, newTh);

//Create data
x = seqa(0.1, 1, 50);
y = sin(x)~cos(x);

//Plot the data with the new line thickness settings
plotXY(myPlot, x, y);
```

plotSetNewWindow

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS, CHAPTER 1**, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotLayout](#), [plotSetTitle](#)

plotSetNewWindow

Purpose

Determines whether each new graph is drawn in a new graph tab or re-uses a pre-existing graph tab.

Format

```
plotSetNewWindow(&myPlot, newW);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>newW</i>	Scalar, 1 to create a new graph tab or 0 to re-use.

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
```



```
myPlot = plotGetDefaults("xy");

//Set graph to create a new graph tab
newW = 1;
plotSetNewWindow(&myPlot, newW);

//Create data
x = seqa(0.1, 1, 50);
y = sin(x)~cos(x);

//Plot the data in a new graph tab window
plotXY(myPlot, x, y);
```

Remarks

To open a new graph window once, use **plotOpenWindow**. This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible in the main application window from the **Tools->Graphics>Preferences** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotOpenWindow](#), [plotSetTitle](#), [plotSetLineColor](#)

plotSetTextInterpreter

Purpose

Controls the text interpreter settings for a graph.

plotSetTextInterpreter

Format

```
plotSetTextInterpreter(&myPlot, interpreter);  
plotSetTextInterpreter(&myPlot, interpreter, location);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>interpreter</i>	String, "html", "plain", "latex".
<i>location</i>	String, "all", "legend", "title" or "axes". Default is "all".

Example

Example 1: Plain interpreter

```
new;  
  
//Declare plotControl structure  
struct plotControl myPlot;  
  
//Initialize plotControl structure  
myPlot = plotGetDefaults("hist");  
  
//Set the interpreter of axes  
plotSetTextInterpreter(&myPlot, "plain", "axes");  
  
//Set the X-axis label, using the > character which would  
//would fail with the default HTML interpreter.  
plotSetXLabel(&myPlot, "Weight > 50 Kg");  
  
//Create data  
x = rndn(1e5,1);  
  
//Plot a histogram of the x data spread over 50 bins
```

```
plotHist(myPlot, x, 50);
```

Example 2: HTML interpreter

You may add Greek letters, mathematical symbols, subscript and superscript to your title, axes and legend using HTML. To add HTML to a label, you can use **plotSetTextInterpreter** to set "html" for the text to be interpreted as HTML.

```
//Set the interpreter of axes
plotSetTextInterpreter(&myPlot, "html", "axes");

label_string = "&beta;";

//Set the X-axis label
plotSetXLabel(&myPlot, label_string);
```

The code above will add the letter β to the graph title. The HTML 'sup' tag will create superscript and the 'sub' tag will create subscript. For example:

```
label_string = "&sigma;<sup>2</sup>";

//Set the X-axis label
plotSetXLabel(&myPlot, label_string);
```

will add σ^2 to your title. While,

```
label_string = "Y<sub>t-1</sub>";

//Set the X-axis label
plotSetXLabel(&myPlot, label_string);
```

will create Y_{t-1} .

Example 3: Latex Interpreter

plotSetTextInterpreter

You can also use Latex to add complex math expression, or non-Latin scripts to your title, axes, and legend. You can use **plotSetTextInterpreter** to set "latex" for the text to be interpreted as Latex.

```
new;

// Declare plotControl structure
struct plotControl myPlot;

// Initialize plotControl structure
myPlot = plotGetDefaults("xy");

// Set up text interpreter
plotSetTextInterpreter(&myPlot, "latex", "all");

// Set up X-axis label
label_string = "x";
plotSetXLabel(&myPlot, label_string, "arial", 20);

// Set up legend in LaTeX format
string legend_string = {
    "y_1 = \\cos{(x)}",
    "y_2 = \\sin{(\\frac{x}{2})} = \\pm \\sqrt{\\frac{1-\\cos{(x)}}{2}}",
    "y_3 = \\cos{(\\frac{x}{2})} = \\pm \\sqrt{\\frac{1+\\cos{(x)}}{2}}";

plotSetLegend(&myPlot, legend_string, "bottom", 1);
plotSetLegendFont(&myPlot, "arial", 20);

// Set up title
title_string = "Trigonometric\\ Functions";
plotSetTitle(&myPlot, title_string, "arial", 24);

// Create data
```

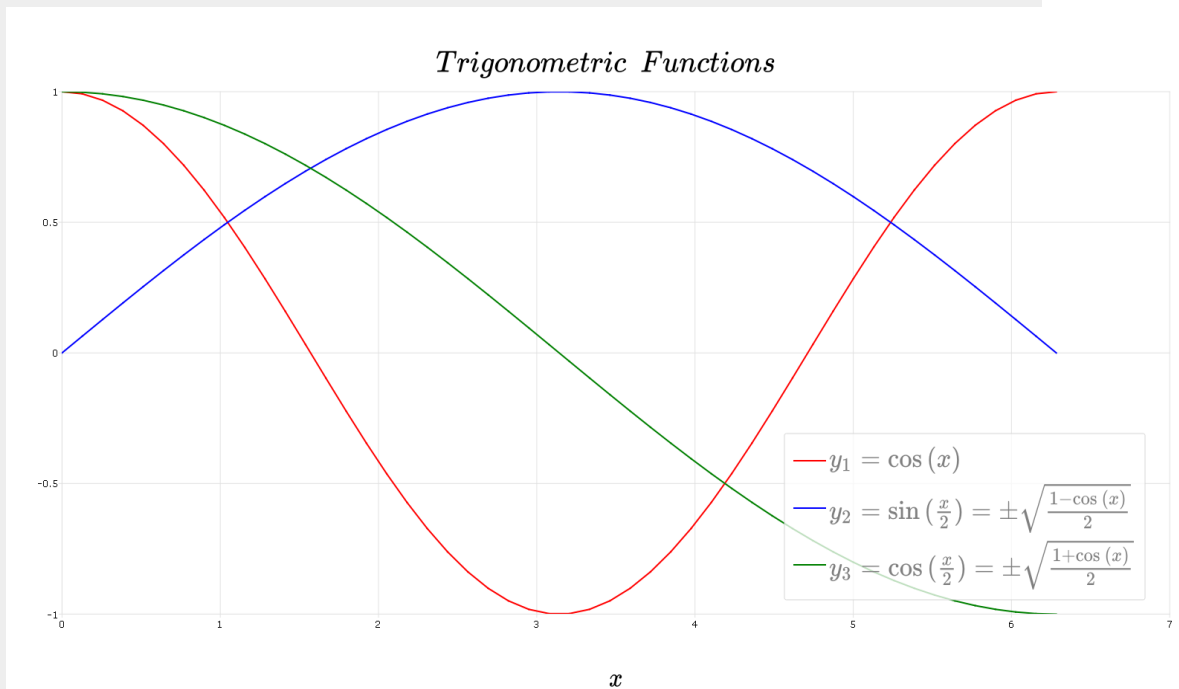
```

n = 50;
x = seqa(0, (2*pi)/(n-1), n);

// Plot
plotXY(myPlot, x, cos(x)~sin(x/2)~cos(x/2));

```

The plot is



Remarks

When the text interpreter is set to use Latex:

- Since backslashes inside of a string represent the escaping of a character, use double backslashes to represent a backslash.

plotSetTitle

- The default mode is that of an in-line equation. To add a section of strictly text, wrap the text only section in `\text{}`. For example:

```
"\\text{The formula is } \\alpha + \\beta_1 X + \\epsilon"
```

- Text outside of a `\text{}` section will use the TeX font. Text inside of a `\text{}` section will use whatever font was specified for the label.

The 'plain' text interpreter will allow you to pass in characters that would be invalid HTML, such as the symbols '<' and '>'.

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[LaTeX in graphics labels](#), [plotGetDefaults](#), [plotSetYLabel](#), [plotSetXLabel](#), [plotSetTitle](#), [plotSetLegend](#)

plotSetTitle

Purpose

Controls the settings for the title for a graph.

Format

```
plotSetTitle(&myPlot, title, font, fontSize, fontColor);  
plotSetTitle(&myPlot, title, font);  
plotSetTitle(&myPlot, title);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>title</i>	String, the new title. This may contain HTML for the creation of Greek letters, mathematical symbols and text formatting.
<i>font</i>	String, font or font family name.
<i>fontSize</i>	Scalar, font size in points.
<i>fontColor</i>	String, named color or RGB value.

Example

Example 1

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("hist");

//Set the title, title font and title font size
plotSetTitle(&myPlot, "GAUSS Example Graph", "verdana",
10);

//Create data
x = rndn(1e5,1);

//Plot a histogram of the x data spread over 50 bins
plotHist(myPlot, x, 50);
```

Example 2

plotSetTitle

You may add Greek letters, mathematical symbols, subscript and superscript to your title using HTML. To add HTML to a label, you need to wrap the text to be interpreted as HTML in HTML tags.

```
label_string = "<html>&beta;</html>";  
plotSetTitle(&myPlot, label_string);
```

The code above will add the letter β to the graph title. The HTML 'sup' tag will create superscript and the 'sub' tag will create subscript. For example:

```
label_string = "<html>&sigma;<sup>2</sup></html>";  
plotSetTitle(&myPlot, label_string);
```

will add σ^2 to your title. While,

```
label_string = "<html>Y<sub>t-1</sub></html>";  
plotSetTitle(&myPlot, label_string);
```

will create Y_{t-1}

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetYLabel](#), [plotSetLineColor](#), [plotSetGrid](#)

plotSetWhichYAxis

Purpose

Assigns curves to the right or left Y-axis.

Format

```
plotSetWhichYAxis(&myPlot, which);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>which</i>	String or Nx1 string array, where each element contains either "right" or "left".

Example

```
//Create data
x = sega(0.1, 0.1, 50);

//Data with y-range of -1 to 1
y1 = sin(x);

//Data with y-range of 0 to 150
y2 = exp(x);

//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("xy");

//Set the first curve, 'y1' to the left Y-axis
//Set the second curve 'y2' to the right Y-axis
```

plotSetXLabel

```
string which = { "left", "right" };  
plotSetWhichYAxis(&myPlot, which);  
  
//Plot the data  
plotXY(myPlot, x, y1~y2);
```

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS, CHAPTER 1**, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetLineSymbol](#)

plotSetXLabel

Purpose

Controls the settings for the X-axis label on a graph.

Format

```
plotSetXLabel(&myPlot, label, font, fontSize,  
fontColor);  
plotSetXLabel(&myPlot, label, font, fontSize);  
plotSetXLabel(&myPlot, label, font);  
plotSetXLabel(&myPlot, label);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
--------------------	---

<i>label</i>	String, the new label. This may contain HTML for the creation of Greek letters, mathematical symbols and text formatting.
<i>font</i>	String, font or font family name.
<i>fontSize</i>	Scalar, font size in points.
<i>fontColor</i>	String, named color or RGB value.

Example

Example 1

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("hist");

//Set the X-axis label, label font, label font size, and
//label color
plotSetXLabel(&myPlot, "Time (sec)", "verdana", 10,
"black");

//Create data
x = rndn(1e5,1);

//Plot a histogram of the x data spread over 50 bins
plotHist(myPlot, x, 50);
```

Example 2

You may add Greek letters, mathematical symbols, subscript and superscript to your axis labels using HTML. To add HTML to a label, you need to wrap the text to be interpreted as HTML in HTML tags.

plotSetXLabel

```
label_string = "<html>&beta;</html>";  
plotSetXLabel(&myPlot, label_string);
```

The code above will add the letter β to the x-axis label. The HTML 'sup' tag will create superscript and the 'sub' tag will create subscript. For example:

```
label_string = "<html>&sigma;<sup>2</sup></html>";  
plotSetXLabel(&myPlot, label_string);
```

will add σ^2 to your x-axis label. While,

```
label_string = "<html>Y<sub>t-1</sub></html>";  
plotSetXLabel(&myPlot, label_string);
```

will create Y_{t-1}

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetXTicInterval](#), [plotSetXTicLabel](#), [plotSetYLabel](#), [plotSetZLabel](#), [plotSetLineColor](#), [plotSetGrid](#)

plotSetXRange

Purpose

Sets the range for the X-axis.

Format

```
plotSetXRange(&myPlot, x_min, x_max);
```

Input

<i>myPlot</i>	A plotControl structure pointer.
<i>x_min</i>	Scalar, minimum limit of the x-axis.
<i>x_max</i>	Scalar, maximum limit of the x-axis.

Example

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("scatter");

//Set X-axis to to range from -5 to +5
plotSetXRange(&myPlot, -5, 5);

//Create and plot data using our x-range
x_1 = rndn(100, 1);
x_2 = rndn(100, 1);

plotScatter(myPlot, x_1, x_2);
```

plotSetXTicCount

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS, CHAPTER 1**, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetLineSymbol](#)

plotSetXTicCount

Purpose

Controls the number of major tics on the X-axis of a 2-D plot.

Format

```
plotSetXTicCount(&myPlot, num_tics);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>num_tics</i>	Scalar, the number of major tics to place on the X-axis.

Example

```
//Create some data to plot
x = seqa(-3, 0.1, 61);
y = x.^3 + rndn(rows(x), 1);

//Plot the data
plotXY(x, y);
```

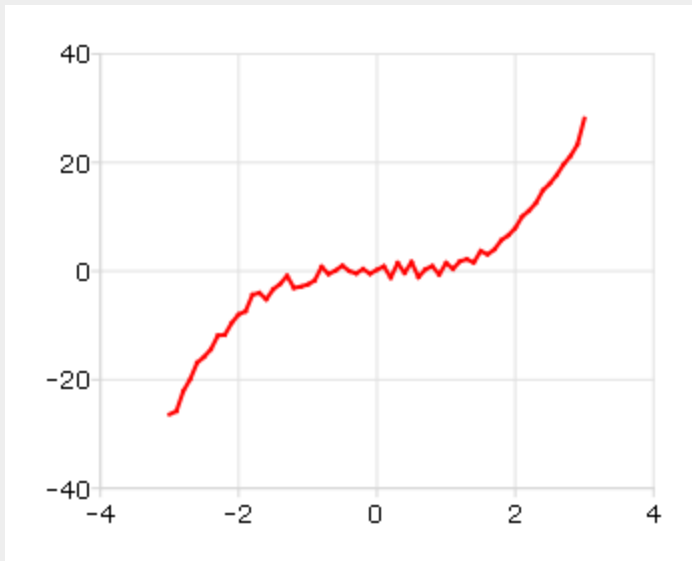


Figure 35.2: 5 tic marks

will produce a graph that looks similar to the one above, with 5 major tic marks on the x-axis. If we use 8 tic marks, there will be one major tic for every integer on the x-axis. We can make that change like this:

```
//Declare and initialize plotControl structure
struct plotControl myPlot;
myPlot = plotGetDefaults("xy");

//Set the x-axis to have 8 tic marks
plotSetXTicCount(&myPlot, 8);

//Plot the data, using the plotControl structure
plotXY(myPlot, x, y);
```

plotSetXTicCount

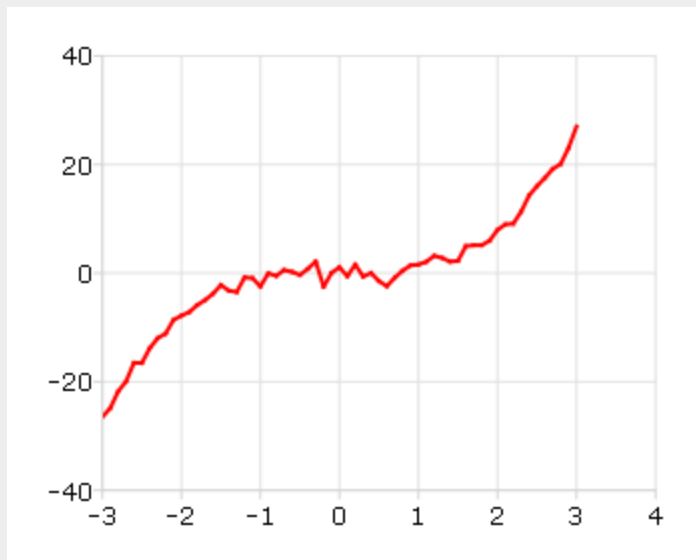


Figure 35.3: 8 tic marks

Remarks

Note that **plotSetXTicInterval** does not provide complete control over the x-axis ticks. If the number of x-ticks requested would cause an odd x-tic interval, GAUSS will create a number of ticks that will provide more even spacing. For instance, in the example above, 8 ticks gave a space between ticks of 1. If we chose 9 ticks, the spacing between ticks would be 0.889. In that case, GAUSS would instead draw 8 ticks for a more even appearance.

For more control over the x-axis of time series plots, use **plotSetXTicInterval** instead.

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotSetXTicInterval](#), [plotSetXLabel](#)

plotSetXTicInterval

Purpose

Controls the interval between X-axis tic labels and also allows the user to specify the first tic to be labeled for 2-D time series graphs.

Format

```
plotSetXTicInterval(&myPlot, ticInterval,  
firstLabeled);  
plotSetXTicInterval(&myPlot, ticInterval);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>ticInterval</i>	Scalar, the number of X-values between X-axis tic labels.
<i>firstLabeled</i>	Scalar, the value of the first X-value on which to place a tic label.

Example

Time Series Example

```
//Declare and initialize plotControl structure  
struct plotControl myPlot;  
myPlot = plotGetDefaults("xy");  
  
//Place one tic label every 4 x-values
```

plotSetXTicInterval

```
ticInterval = 4;  
plotSetXTicInterval(&myPlot, ticInterval);  
  
//Start the time series in April of 2008  
dtstart = 200804;  
  
//Specify quarterly data  
frequency = 4;  
  
//Create the multiplicative sequence 1, 2, 4, 8...  
y = seqm(1, 2, 10);  
  
//Create a time series plot of the data.  
plotTS(myPlot, dtstart, frequency, y);
```

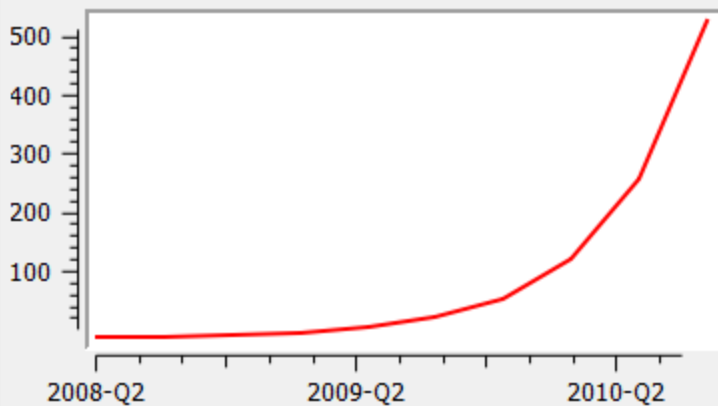


Figure 35.4: Time series plot example

If you would like to change the tic labels so that they start on the first full year, 2009, continuing with the example from above, execute the following lines:

```
//Set the optional 'firstLabeled' parameter  
plotSetXTicInterval(&myPlot, ticInterval, 2009);  
plotTS(myPlot, dtstart, frequency, y);
```

This new plot should now have tic labels only on the first quarters of each year:

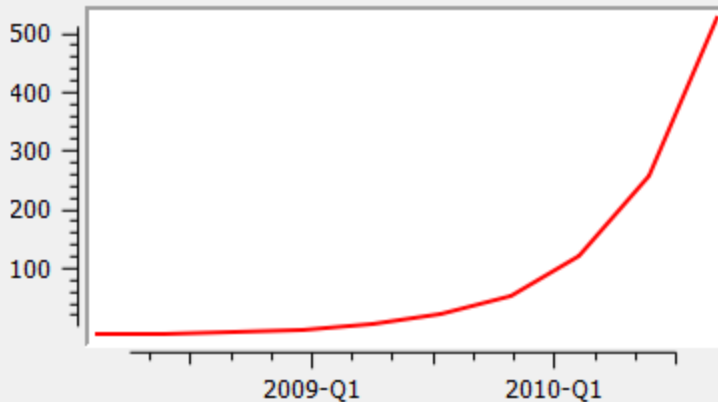


Figure 35.5: Revised time series example

Remarks

plotSetXTicInterval is currently only supported for use with time series plots. It is ignored by other plot types.

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[dttostr](#), [strtodt](#), [plotSetXLabel](#), [plotSetXTicLabel](#)

plotSetXTicLabel

plotSetXTicLabel

Purpose

Controls the formatting and angle of X-axis tic labels for 2-D time series graphs.

Format

```
plotSetXTicLabel(&myPlot, fmt, angle);  
plotSetXTicLabel(&myPlot, fmt);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>fmt</i>	String, the desired formatting for the X-axis tic labels. <ul style="list-style-type: none">• Time series graphs use the same formatting type as function dttostr• Other graph types use a sprintf style formatting string.
<i>angle</i>	Scalar, the angle in degrees at which to display the X-axis tic labels.

Example

Example 1: Time series

```
//Declare and initialize plotControl structure  
struct plotControl myPlot;  
myPlot = plotGetDefaults("xy");  
  
y = rndn(5, 1);  
  
plotSetXTicLabel(&myPlot, "YYYY-MO");
```

```
//Start the series in January 1982
dtstart = 198201;

//Specify the data to be monthly
frequency = 12;

//Draw the time series plot
plotTS(myPlot, dtstart, frequency, y);
```

The code above produces a graph with X-tic labels like the image below:

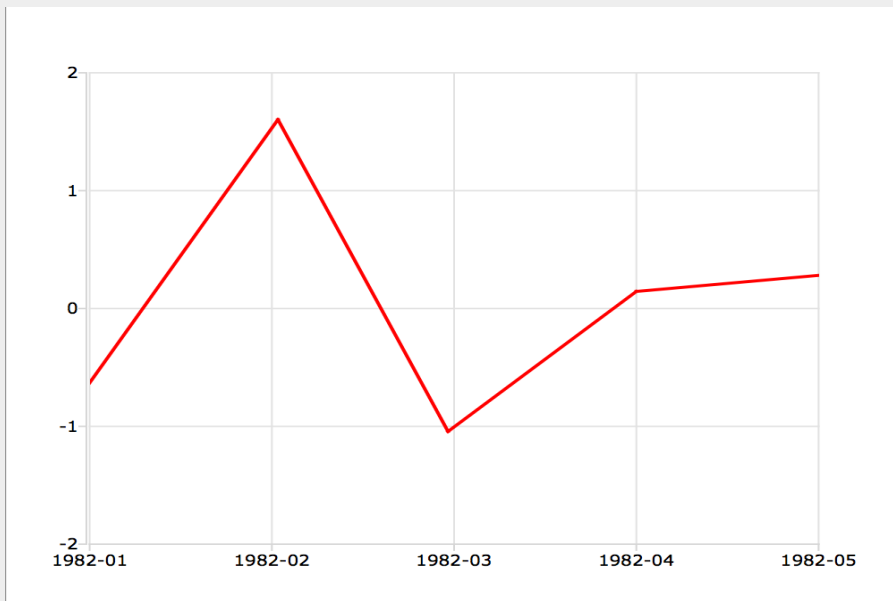


Figure 35.6: X-tic labels

Changing to format string to "MO/YYYY" will change the labels to appear like this:

plotSetXTicLabel

01/1982

Changing to format string to "YYYY-QQ" will change the the labels to appear like this:

1982-Q1

Example 2: Other than time series

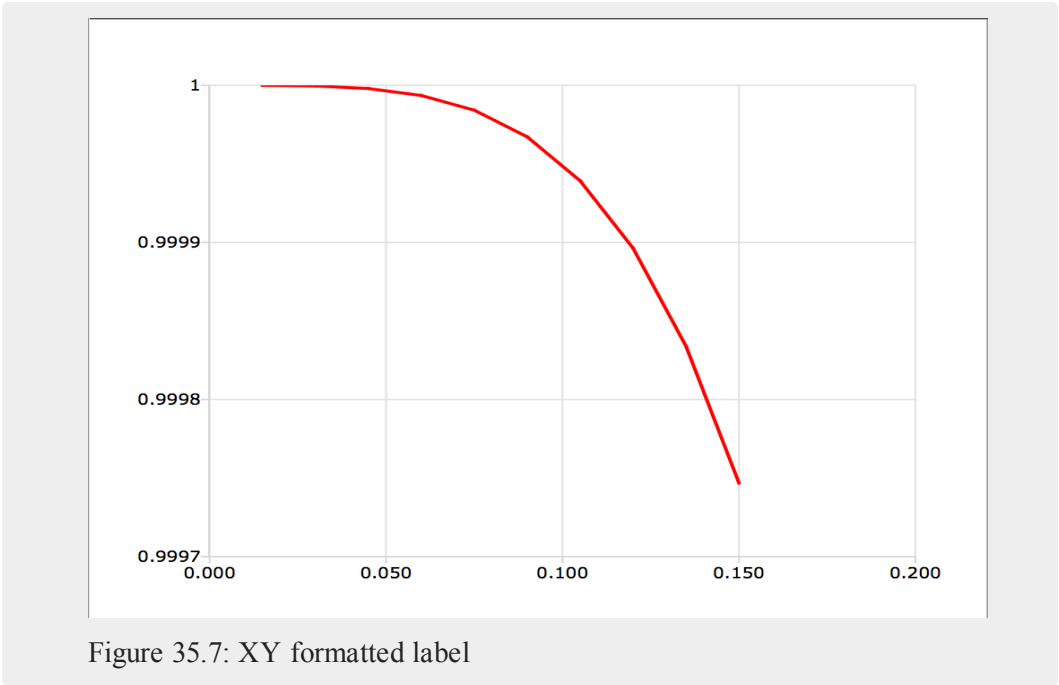
```
//Declare and initialize plotControl structure
struct plotControl myPlot;
myPlot = plotGetDefaults("xy");

x = seqa(0.015, 0.015, 10);
y = cos(x.^2);

plotSetXTicLabel(&myPlot, "%.3f");

//Draw the graph, using our format specifier
plotXY(myPlot, x, y);
```

The code above produces a graph with X-tic labels like the image below:



Remarks

Time series format specifier

The following format specifiers are supported for the second input to `plotSetXTicLabel`, *fmt* when the plot will be drawn with `plotTS`:

YYYY	4 digit year
YR	Last two digits of year
QQ	Quarter of the year. This is calculated from the month number.
MO	Number of month, 01-12

plotSetXTicLabel

DD	Day of month, 01-31
HH	Hour of day, 00-23
MI	Minute of hour, 00-59
SS	Second of minute, 00-59

To learn more about DT scalar format, see **Date and Time Formats**, Section 1.0.1, or the functions [dttostr](#) and [strtodt](#).

Format specifier for other graph types

Graph types other than time series use a **sprintf** style format string. The parameters of the format string are:

- precision - a literal scalar numeral, the number of digits after the decimal point
- format type - one of the following letters:
 - d - integer
 - e - scientific notation
 - f - floating point/decimal
 - g - either scientific or decimal, whichever is most compact

The format of the format string is:

```
"%.<precision><format type>"
```

Below are some examples of different format strings and how they would represent *pi*

- "%.2f" - 3.14
- "%.4f" - 3.1415
- "%.2e" - 3.14e+00
- "%.2g" - 3.14
- "%d" - 3

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the

Tools->Preferences->Graphics menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[dttostr](#), [strtodt](#), [plotSetXLabel](#), [plotSetXTicInterval](#)

plotSetYLabel

Purpose

Controls the settings for the Y-axis label on a graph.

Format

```
plotSetYLabel(&myPlot, label, font, fontSize,
              fontColor);
plotSetYLabel(&myPlot, label, font, fontSize);
plotSetYLabel(&myPlot, label, font);
plotSetYLabel(&myPlot, label);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>label</i>	String or 2x1 string array, the new label or labels. If you are using more than one Y-axis, the first element of the 2x1 label string array will set the label for the left Y-axis and the second element will set the label for the right Y-axis. This may contain HTML for the creation of Greek letters, mathematical symbols and text formatting.
<i>font</i>	String, font or font family name.

plotSetYLabel

<i>fontSize</i>	Scalar, font size in points.
<i>fontColor</i>	String, named color or RGB value.

Example

Example 1: Basic usage

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("hist");

//Set the Y-axis label, label font, font size and color
plotSetYLabel(&myPlot, "Time (sec)", "verdana", 10,
"black");

//Create data
x = rndn(1e5,1);

//Plot a histogram of the x data spread over 50 bins
plotHist(myPlot, x, 50);
```

Example 2: Setting both Y-axes

```
//Create with different Y-ranges
x = seqa(1,1,5);
y = { 98 1.5,
      92 0.9,
      97 1.3,
      94 2.1,
      95 2.4 };

//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
```

```

myPlot = plotGetDefaults("xy");

//Set the first curve to use the left Y-axis and the second
curve to use the right
plotSetWhichYAxis(&myPlot, "left" $| "right");

//Set the left and right Y-axis labels
plotSetYLabel(&myPlot, "Number of subjects", "Percent clas-
sified");

//Plot the data
plotXY(myPlot, x, y);

```

Example 3

You may add Greek letters, mathematical symbols, subscript and superscript to your axis labels using HTML. To add HTML to a label, you need to wrap the text to be interpreted as HTML in HTML tags.

```

label_string = "<html>&beta;</html>";
plotSetYLabel(&myPlot, label_string);

```

The code above will add the letter β to the y-axis label. The HTML 'sup' tag will create superscript and the 'sub' tag will create subscript. For example:

```

label_string = "<html>&sigma;<sup>2</sup></html>";
plotSetYLabel(&myPlot, label_string);

```

will add σ^2 to your y-axis label. While,

```

label_string = "<html>Y<sub>t-1</sub></html>";
plotSetYLabel(&myPlot, label_string);

```

will create Y_{t-1}

plotSetYRange

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetXLabel](#), [plotSetXTicInterval](#), [plotSetXTicLabel](#), [plotSetZLabel](#), [plotSetLineColor](#), [plotSetGrid](#)

plotSetYRange

Purpose

Sets the range for the y-axis.

Format

```
plotSetYRange(&myPlot, y_min, y_max);
```

Input

p

<i>&myPlot</i>	A plotControl structure pointer.
<i>y_min</i>	Scalar, minimum limit of the y-axis.
<i>y_max</i>	Scalar, maximum limit of the y-axis.

Example

```
//Declare plotControl structure
struct plotControl myPlot;
```

```
//Initialize plotControl structure
myPlot = plotGetDefaults("scatter");

//Set y-axis to to range from 0 to 2
plotSetYRange(&myPlot, 0, 2);

//Create and plot data using our x-range
x = rndu(100, 1);
y = rndu(100, 1);

plotScatter(myPlot, x, y);
```

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS, CHAPTER 1**, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetLineSymbol](#)

plotSetYTicCount

Purpose

Controls the number of major tics on the y-axis of a 2-D plot.

Format

```
plotSetYTicCount(&myPlot, num_tics);
```

plotSetYTicCount

Input

<code>&myPlot</code>	A plotControl structure pointer.
<code>num_tics</code>	Scalar, the number of major tics to place on the y-axis.

Example

```
//Create some data to plot
x = seqa(-3, 0.1, 61);
y = x.^3 + rndn(rows(x), 1);

//Plot the data
plotXY(x, y);
```

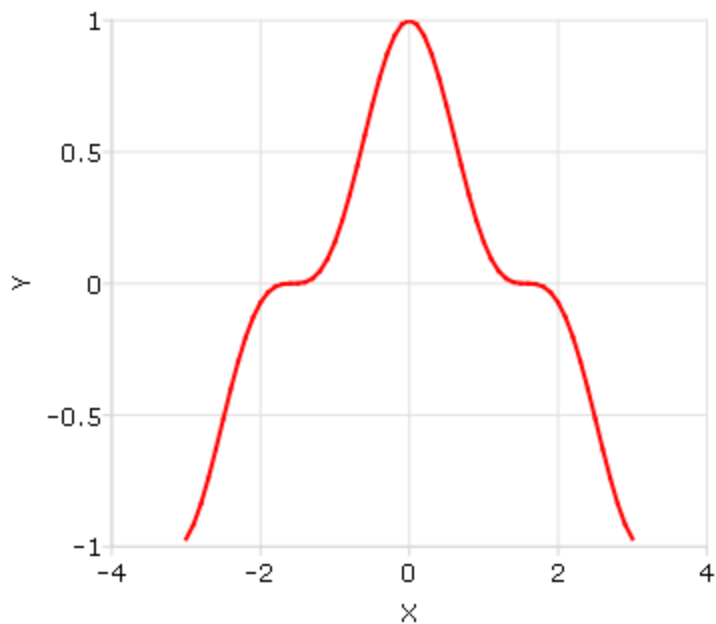


Figure 35.8: 5 tic marks

will produce a graph that looks similar to the one above, with 5 major tic marks on the y-axis. If we use 11 tic marks, there will be one major tic for 0.2 on the y-axis. We can make that change like this:

```
//Declare and initialize plotControl structure
struct plotControl myPlot;
myPlot = plotGetDefaults("xy");

//Set the y-axis to have 11 tic marks
plotSetYTicCount(&myPlot, 11);

//Plot the data, using the plotControl structure
plotXY(myPlot, x, y);
```

plotSetYTicCount

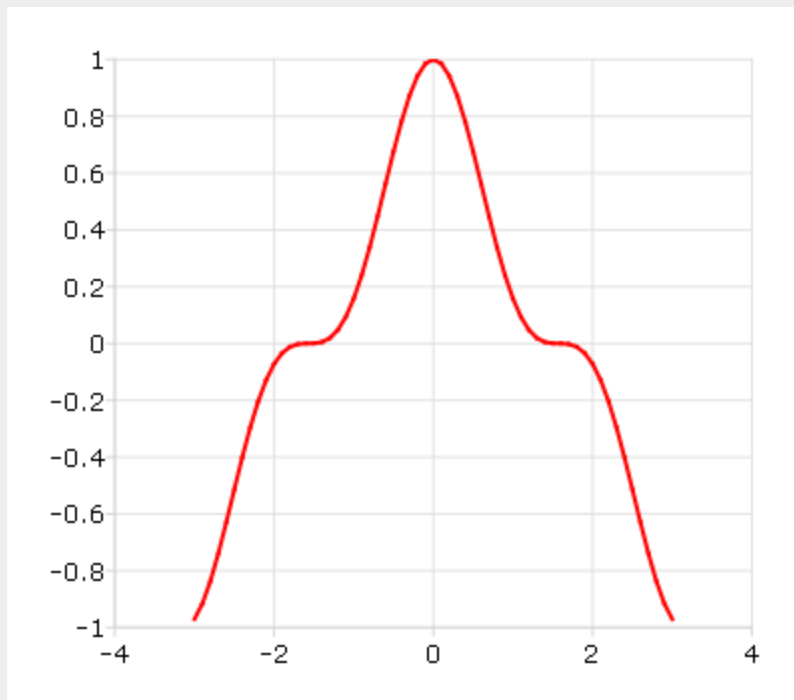


Figure 35.9: 11 tic marks

Remarks

Note that **plotSetYTicCount** does not provide complete control over the y-axis ticks. If the number of y-tics requested would cause an odd tic interval, GAUSS will create a number of ticks that will provide more even spacing. For instance, in the example above, 11 tics gave a space between tics of 0.2. If we chose 10 tics, the spacing between tics would be 0.222. In that case, GAUSS would instead draw 10 tics for a more even appearance.

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the

Tools->Preferences->Graphics menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotSetXTicInterval](#), [plotSetXLabel](#)

plotSetZLabel

Purpose

Controls the settings for the Z-axis label on a surface plot.

Format

```
plotSetZLabel(&myPlot, label, font, fontSize,
              fontColor);
plotSetZLabel(&myPlot, label, font, fontSize);
plotSetZLabel(&myPlot, label, font);
plotSetZLabel(&myPlot, label);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>label</i>	String, the new label.
<i>font</i>	String, font or font family name.
<i>fontSize</i>	Scalar, font size in points.
<i>fontColor</i>	String, named color or RGB value.

plotSetZLabel

Example

Example 1

```
//Declare plotControl structure
struct plotControl myPlot;

//Initialize plotControl structure
myPlot = plotGetDefaults("surface");

//Set the Z-axis label, label font, font size, and color
plotSetZLabel(&myPlot, "Depth", "verdana", 10, "black");

//Create data
x = seqa(-10.6, .3, 71)';
y = seqa(-12.4, .35, 71);
z = sin(sqrt((x/2)^2+(y/2)^2)) ./ sqrt(x^2+y^4);
z = z .* sin(x/3);

//Plot the data
plotSurface(myPlot, x, y, z);
```

Remarks

This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetXLabel](#), [plotSetXTicInterval](#), [plotSetXTicLabel](#), [plotSetYLabel](#), [plotSetLineColor](#), [plotSetGrid](#)

plotSetZLevels

Purpose

Controls the heights at which lines are drawn on a contour plot.

Format

```
plotSetZLevels(&myPlot, zlevels);
```

Input

<i>&myPlot</i>	A plotControl structure pointer.
<i>zlevels</i>	Scalar, or Nx1 vector. If <i>zlevels</i> is a scalar, <i>zlevels</i> represents the number of different heights at which to draw contour lines (from MIN(Z) to MAX(Z)). If <i>zlevels</i> is an Nx1 vector, <i>zlevels</i> indicates the heights at which to draw the contour lines.

Examples

Example 1: Basic example setting number of lines

```
//Linear sequence from -3 to +3
y = seqa(-3, 0.1, 61);
x = y';

//Joint probability of 'x' and 'y'
z = pdfn(x) .* pdfn(y);

//Declare plotControl structure
//and fill in with defaults for surface/contour
struct plotControl myPlot;
```

plotSetZLevels

```
myPlot = plotGetDefaults("surface");

//Set number of lines to be drawn
//MIN(Z) < Line 1 < Line 2 < Line 3 < Line 4 < MAX(Z)
plotSetZLevels(&myPlot, 4);

//Draw contour plot with 4 lines
plotContour(myPlot, x,y,z);
```

Example 2: Basic example setting heights of lines

Continuing with the data and plotControl structure created in the example above:

```
//Draw lines at z == 0.05 and when z == 0.1
plotSetZLevels(&myPlot, 0.05 | 0.1);

//Draw contour plot with lines at 0.05 and 0.1
plotContour(myPlot, x,y,z);
```

Example 3

```
//Clear out variables in GAUSS workspace
new;

//Create data
x = seqa(-4,.125,161)';
y = seqa(-8,.125,161);
z = sin(x) .* cos(y) * .5;
z = z .* sin(x/3) .* cos(y/3);
z = z .* sin(x/5) + sin(y/2.5)/3 + sin(x/2.5)/3;

//Set up control structure with defaults
//for surface plots
struct plotControl myPlot;
myPlot = plotGetDefaults("surface");
```

```
//Set title and Z axis label
plotSetTitle(&myPlot, "Contour plot example");

//Set color map for contour
plotSetColorMap(&myplot, "plasma");

//Set up flag of show label for contour
showLabels = 1;

//Set up format for contour label
labelFormat = "g";

//Set up precision for contour label
labelPrecision = 5;

//Use function to set up contour labels
plotSetContourLabels(&myplot, showLabels, labelFormat,
labelPrecision);

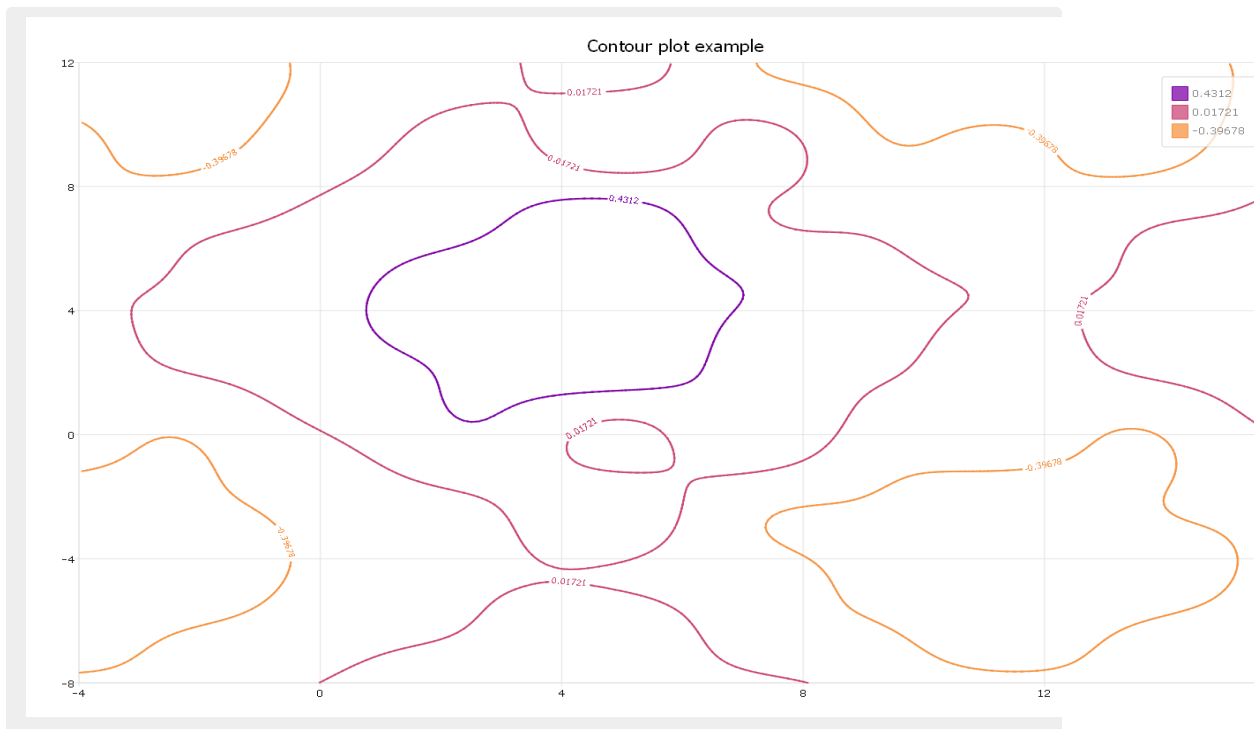
//Use function to set up contour z levels = 3
plotSetZLevels(&myplot, 3);

//Draw graph using plotcontrol structure
plotContour(myPlot, x, y, z);
```

The Plot is

d

plotSetZLevels



Remarks

- To indicate a single Z-level, pass in a 2x1 vector with the same value for both elements:

```
single_level = { 1.5, 1.5 };  
plotSetLevels(&myPlot, single_level);
```

- This function sets an attribute in a **plotControl** structure. It does not affect an existing graph, or a new graph drawn using the default settings that are accessible from the **Tools->Preferences->Graphics** menu. See **GAUSS GRAPHICS**, CHAPTER 1, for more information on the methods available for customizing your graphs.

See Also

[plotGetDefaults](#), [plotSetColorMap](#), [plotSetContourLabels](#)

plotSurface

Purpose

Graphs a 3-D surface.

Format

```
plotSurface(myPlot, x, y, z);  
plotSurface(x, y, z);
```

Input

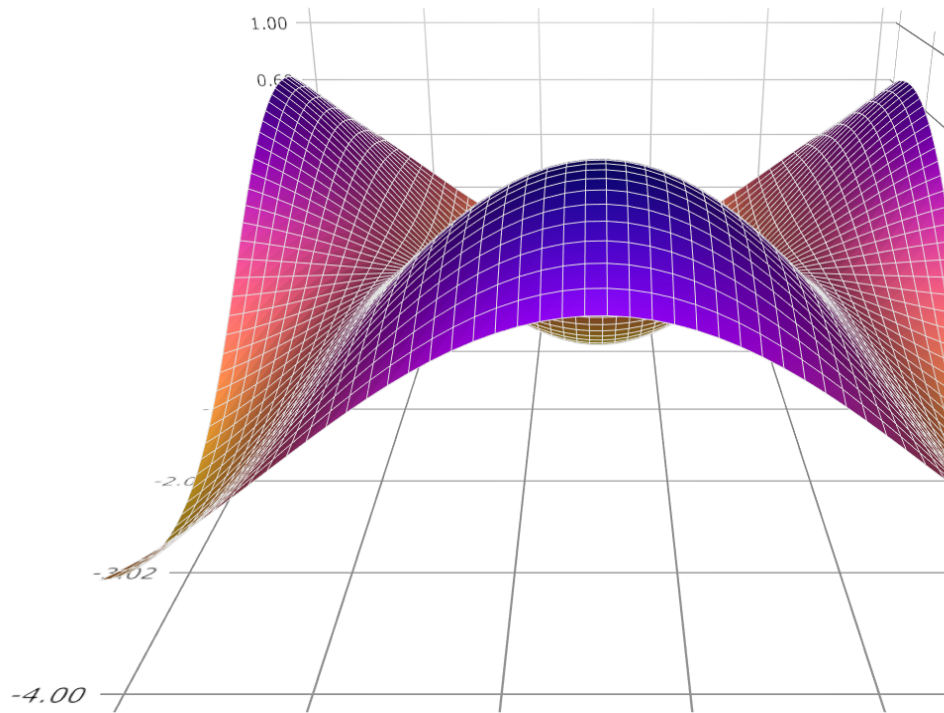
<i>myPlot</i>	A plotControl structure.
<i>x</i>	1xK vector, the X axis data.
<i>y</i>	Nx1 vector, the Y axis data.
<i>z</i>	NxK matrix, the matrix of height data to be plotted.

Examples

```
//Clear out variables in GAUSS workspace  
new;  
  
//Create contour data  
x = sega(-4, 0.1, 50)';  
y = x';  
z = sin(y) .* cos(x) ;  
  
//Set up control structure with defaults  
//for surface plots
```

plotSurface

```
struct plotControl myPlot;  
myPlot = plotGetDefaults("surface");  
  
//Set color map  
plotSetColormap(&myPlot, "plasma");  
  
//Draw graph using plotcontrol structure  
plotSurface(myPlot, x, y, z);
```



See Also

[plotSetColorMap](#), [plotContour](#), [plotSetBkdColor](#)

plotTS

Purpose

Creates a graph of time series data.

Format

```
plotTS(myPlot, dtstart, frequency, y);
plotTS(dtstart, frequency, y);
```

Input

<i>myPlot</i>	A plotControl structure.						
<i>dtstart</i>	Scalar, starting date in DT scalar format.						
<i>frequency</i>	Scalar, frequency of the data per year. Valid options include: <table> <tr> <td>1</td><td>Yearly</td></tr> <tr> <td>4</td><td>Quarterly</td></tr> <tr> <td>12</td><td>Monthly</td></tr> </table>	1	Yearly	4	Quarterly	12	Monthly
1	Yearly						
4	Quarterly						
12	Monthly						
<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular line.						

Examples

Example 1

```
//Create some data to plot
y = rndn(100, 1);

//The first input starts the series in January of 1982
//The second input specifies the data to be monthly
plotTS(1982, 12, y);
```

Example 2

```
//Create file name with full path
file = getGAUSSHome() $+ "examples/tbill_3mo.xlsx";

//Load starting date
date_1 = xlsReadM(file, "A2:A2");

//Load the first 20 observations from the time series
y = xlsReadM(file, "B2:B20");

//Specify the data is monthly
freq = 12;

//Draw the time series plot
plotTS(date_1, freq, y);
```

Example 3: Time Series Plot With Custom X-tics

```
//Create file name with full path
file = getGAUSSHome() $+ "examples/tbill_3mo.xlsx";

//Load date of observation 20 (header is row 1)
date_1 = xlsReadM(file, "A21:A21");
```

```
//Load 28 observations
y = xlsReadM(file, "B21:B49");

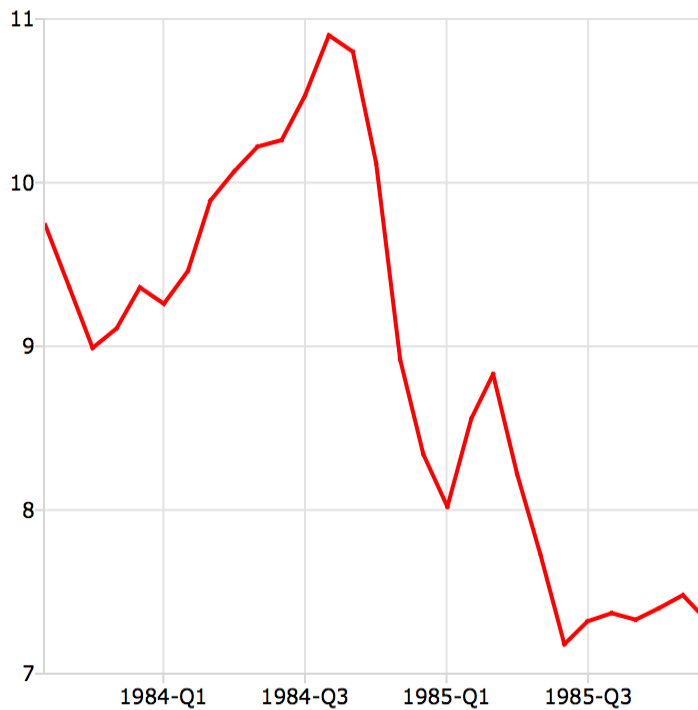
//Declare 'myPlot' to be a plotControl structure
//and fill it with 'xy' default settings
struct plotControl myPlot;
myPlot = plotGetDefaults("xy");

//Place first 'X' tic mark at 1984 month 1 and draw one
every 6 months
plotSetXTicInterval(&myPlot, 6, 1984);

//Display only 4 digit year on 'X' tic labels
plotSetXTicLabel(&myPlot, "YYYY-QQ");

//Draw time series plot, using settings in 'myPlot'
plotTS(myPlot, date_1, 12, y);
```

plotTS



Example 4

In DT Scalar format, quarters are represented by supplying the first month of the quarter for the sixth and seventh leading digits. As we see below, 200504 represents April of 2005, but it also represents the second quarter of April 2005.

```
//The first input starts the series in the second quarter  
of 2005  
//The second input specifies the data to be quarterly  
plotTS(200504, 4, y);
```

Remarks

Formatting for the X-tic labels can be set with the function `plotSetXTicLabel`. If a `plotControl` structure is not passed in to `plotTS`, or the format specifier is not set with `plotSetXTicLabel` the default formatting: for annual data is "YYYY", for quarterly data "YYYY-QQ" and for monthly data is "YYYY-MO".

By default missing values in the *y* variable will be represented as gaps in the line.

See Also

[plotSetXTicLabel](#), [plotSetXTicInterval](#), [plotScatter](#)

plotXY

Purpose

Graphs X vs. Y using Cartesian coordinates.

Format

```
plotXY(myPlot, x, y);
plotXY(x, y);
```

Input

<i>myPlot</i>	A plotControl structure.
<i>x</i>	Nx1 or NxM matrix. Each column contains the X values for a particular line.
<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular line.

Remarks

By default missing values in the *y* variable will be represented as gaps in the line.

polar

See Also

[plotLogX](#), [plotLogLog](#), [plotScatter](#)

polar

Purpose

Graph data using polar coordinates. NOTE: This function is for use only with the deprecated PQG graphics.

Library

pgraph

Format

```
polar(radius, theta);
```

Input

<i>radius</i>	Nx1 or NxM matrix. Each column contains the magnitude for a particular line.
<i>theta</i>	Nx1 or NxM matrix. Each column represents the angle values for a particular line.

Source

polar.src

See Also

[xy](#), [logx](#), [logy](#), [loglog](#), [scale](#), [xtics](#), [ytics](#)

polychar

Purpose

Computes the characteristic polynomial of a square matrix.

Format

$c = \text{polychar}(x);$

Input

x $N \times N$ matrix.

Output

c $(N+1) \times 1$ vector of coefficients of the N th order characteristic polynomial of x :

$$p(x) = c[1] * x^n + c[2] * x^{(n-1)} + \dots + c[n] * x + c[n+1];$$

Remarks

The coefficient of x^n is set to unity ($c[1]=1$).

Source

poly.src

See Also

[polymake](#), [polymult](#), [polyroot](#), [polyeval](#)

polyeval

polyeval

Purpose

Evaluates polynomials. Can either be one or more scalar polynomials or a single matrix polynomial.

Format

```
y = polyeval(x, coeffs);
```

Input

x	1xK or NxN; that is, x can either represent K separate scalar values at which to evaluate the (scalar) polynomial(s), or it can represent a single NxN matrix.
c	(P+1)xK or (P+1)x1 matrix of coefficients of polynomials to evaluate. If x is 1xK, then c must be (P+1)xK. If x is NxN, c must be (P+1)x1. That is, if x is a matrix, it can only be evaluated at a single set of coefficients.

Output

y	Kx1 vector (if c is (P+1)xK) or NxN matrix (if c is (P+1)x1 and x is NxN): <div>$y = (\begin{bmatrix} c[1, \cdot] \cdot x^p + c[2, \cdot] \cdot x^{(p-1)} + \dots + c[p+1, \cdot] \end{bmatrix})';$</div>
---	--

Remarks

In both the scalar and the matrix case, Horner's rule is used to do the evaluation. In the

scalar case, the function **recsercp** is called (this implements an elaboration of Horner's rule).

Example

Scalar example 1

```
//Evaluate 2^4 + 2^3 + 2^1 + 2^0
x = 2;
coefs = { 1, 1, 0, 1, 1 };
y = polyeval(x,coefs);
```

The result is 27. Note that this is the decimal value of the binary number 11011.

Scalar example 2

```
//Evaluate 7*2^3 + 2^1 + 2^0
x = 2;
coefs = { 7, 0, 1, 1 };
y = polyeval(x,coefs);
```

The result is 59.

Matrix example 1

```
//Evaluate A*A*A + 2*A
A = { 2 6,
      4 8 };
coefs = { 1, 0, 2, 0 };
A_3 = polyeval(A, coefs);
```

The above code will set A_3 equal to:

```
300    660
440    960
```

Matrix example 2

polygamma

```
//Evaluate A*A*A*A
A = { 1.2 3.1,
      1.7 0.8 };
coefs = 1|zeros(4,1);
y = polyeval(A,coefs);
```

You can raise a matrix to the n'th power with the command:

```
A_n = polyeval(A, 1|zeros(n,1));
(e.g: A*A*A*A*...*A).
```

Source

poly.src

See Also

[polymake](#), [polychar](#), [polymult](#), [polyroot](#)

polygamma

Purpose

Computes the polygamma function.

Format

```
f = polygamma(z,n);
```

Input

z	NxK matrix; z may be complex.
n	The order of the function. If n is 0 then f will be the Digamma function. If $n = 1,2,3$, etc., then f will be the

tri-, tetra-, penta-, s-, etc., Gamma function. Real (n) must be positive.

Output

f NxK matrix; f may be complex.

Examples

Example 1

```
//Both calls are equivalent  
f = digamma(1);  
f2 = polygamma(1, 0);
```

After the code above, both f and $f2$ should be equal to $-\gamma$, where γ represents the Euler-Mascheroni constant:

```
-0.57721566
```

Example 2: Compute the pentagamma function

```
f = polygamma(1.5, 4);
```

After the code above, f should be equal to:

```
-3.47425
```

Example 3: Complex input

```
//Set 'z' equal to complex number -45.6-29.4i  
z = { -45.6 - 29.4i };  
polygamma(z, 101);
```

```
12.501909 + 9.0829590i
```

Example 4

polygamma

```
z = { -11.5 - 0.577007813568142i };  
polygamma(z,10);
```

will return the value:

```
-4.984e-06 + 8.217e-07i
```

Remarks

The polygamma function of order n is defined by the equation:

$$\psi^{(n)}(z) = \frac{d^n}{dz^n} \psi(z) = \frac{d^{n+1}}{dz^{n+1}} \ln \Gamma(z)$$

This program uses the partial fraction expansion of the derivative of the log of the Lanczos series approximation for the Gamma function. Accurate to about 12 digits.

References

1. C. Lanczos, SIAM JNA 1, 1964. pp. 86-96.
2. Y. Luke, "The Special ... approximations," 1969 pp. 29-31.
3. Y. Luke, "Algorithms ... functions," 1977.
4. J. Spouge, SIAM JNA 31, 1994. pp. 931.
5. W. Press, "Numerical Recipes."
6. S. Chang, "Computation of special functions," 1996.
7. Abramowitz & Stegun, section eq 6.4.6
8. Original code by Paul Godfrey

polyint

Purpose

Calculates an Nth order polynomial interpolation.

Format

```
y = polyint(xa, ya, x);
```

Input

<i>xa</i>	Nx1 vector, <i>x</i> values.
<i>ya</i>	Nx1 vector, <i>y</i> values.
<i>x</i>	scalar, <i>x</i> value to solve for.

Global Input

<i>_poldeg</i>	scalar, the degree of polynomial required, default 6.
----------------	---

Output

<i>y</i>	result of interpolation or extrapolation.
----------	---

Global Output

<i>_polerr</i>	scalar, interpolation error.
----------------	------------------------------

Remarks

Calculates an Nth order polynomial interpolation or extrapolation of *x* on *y* given the vectors *xa* and *ya* and the scalar *x*. The procedure uses Neville's algorithm to

polymake

determine an up to Nth order polynomial and an error estimate.

Polynomials above degree 6 are not likely to increase the accuracy for most data. Test `_polerr` to determine the required `_poldeg` for your problem.

Source

polyint.src

Technical Notes

Press, W.P., B.P. Flannery, S.A. Teukolsky, and W.T. Vetterling. *Numerical Recipes: The Art of Scientific Computing*. NY: Cambridge Press, 1986.

polymake

Purpose

Computes the coefficients of a polynomial given the roots.

Format

`c = polymake(r);`

Input

`r` Nx1 vector containing roots of the desired polynomial.

Output

`c` (N+1)x1 vector containing the coefficients of the Nth order polynomial with roots `r`:

$$p(z) = c[1] * z^N + c[2] * z^{(N-1)} + \dots c[n] * z + c[n+1]$$

p

Remarks

The coefficient of z^n is set to unity ($c[1]=1$).

Example

```
//Assign values for the roots of the polynomial
r = { 2, 1, 3 };

//Calculate the coefficients
c = polymake(r);

//Print 3 spaces for each number and 1 digit after the
//decimal place
format /rd 3,1;

//Iterate through each root in 'r'
for i(1, 3, 1);
    rtmp = r[i];
    //Calculate the polynomial
    rout = c[1]*rtmp^3 + c[2]*rtmp^2 + c[3]*rtmp + c[4];
    print "rtmp = " rtmp "rout = " rout;
endfor;
```

Since the values of r are roots for this polynomial, $rout$ should equal 0. Thus the code above gives the following output:

```
rtmp = 2.0 rout = 0.0
rtmp = 1.0 rout = 0.0
rtmp = 3.0 rout = 0.0
```

This example assigns c to be equal to:

```
1.0
c = -6.0
```

polymat

```
11.0
-6.0
```

This represents the polynomial:

$$x^3 - 6x^2 + 11x - 6$$

Source

poly.src

See Also

[polychar](#), [polymult](#), [polyroot](#), [polyeval](#)

polymat

Purpose

Returns a matrix containing the powers of the elements of x from 1 to p .

Format

```
y = polymat(x, p);
```

Input

p

x	$N \times K$ matrix.
p	scalar, positive integer.

Output

y	$N \times (p * K)$ matrix containing powers of the elements of
-----	--

x from 1 to p . The first K columns will contain first powers, the second K columns second powers, and so on.

Remarks

To do polynomial regression use `ols`:

```
{ vnam,m,b,stb,vc,stderr,sigma,cx,rsq,resid,dwstat } = ols
(0,y, polymat(x,p));
```

Source

polymat.src

polymroot

Purpose

Computes the roots of the determinant of a matrix polynomial.

Format

```
 $r$  = polymroot( $c$ );
```

Input

c	$(N+1)*K \times K$ matrix of coefficients of an N th order polynomial of rank K .
-----	---

Output

r	$K \times N$ vector containing the roots of the determinantal
-----	---

p

polymroot

equation.

Remarks

c is constructed of $N+1$ $K \times K$ coefficient matrices stacked vertically with the coefficient matrix of the t^n at the top, $t^{(n-1)}$ next, down to the t^0 matrix at the bottom.

Note that this procedure solves the scalar problem as well, that is, the one that POLYROOT solves.

Example

Solve

$$\det(A_2 t^2 + A_1 t + A_0) = 0$$

where:

$$A_2 = \begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \quad A_1 = \begin{bmatrix} 5 & 8 \\ 10 & 7 \end{bmatrix} \quad A_0 = \begin{bmatrix} 3 & 4 \\ 6 & 5 \end{bmatrix}$$

```
a2 = { 1 2, 2 1 };  
a1 = { 5 8, 10 7 };  
a0 = { 3 4, 6 5 };
```

```
//The pipe operator '|' provides vertical concatenation  
print polymroot(a2|a1|a0);
```

```
-4.3027756  
-.69722436  
-2.6180340  
-.38196601
```

polymult

Purpose

Multiplies polynomials.

Format

```
c = polymult(c1, c2);
```

Input

$c1$	$(D1+1) \times 1$ vector containing the coefficients of the first polynomial.
$c2$	$(D2+1) \times 1$ vector containing the coefficients of the second polynomial.

Output

c	$(D1+D2) \times 1$ vector containing the coefficients of the product of the two polynomials.
-----	--

Example

This example multiplies the polynomials:

$$(2x + 1)(2x^2 + 1)$$

and returns the answer:

$$4x^3 + 2x^2 + 2x + 1$$

```
//Assign c1 to represent 2x + 1  
c1 = { 2, 1 };
```

polymult

```
//Assign c2 to represent  $2x^2 + 1$ 
c2 = { 2, 0, 1 };
c = polymult(c1,c2);
```

After the code above:

```
      4
c = 2
      2
      1
```

Technical Notes

If the degree of $c1$ is $D1$ (e.g., if $D1=3$, then the polynomial corresponding to $c1$ is cubic), then there must be $D1+1$ elements in $c1$ (e.g., 4 elements for a cubic). Thus, for instance the coefficients for the polynomial

$$5x^3 + 6x + 3$$

would be:

```
//Using the pipe operator for vertical concatenation
c1 = 5|0|6|3;
```

or

```
//Using an array assignment
c1 = { 5, 0, 6, 3 };
```

(Note that zeros must be explicitly given if there are powers of x missing.)

Source

poly.src

See Also

[polymake](#), [polychar](#), [polyroot](#), [polyeval](#)

polyroot

Purpose

Computes the roots of a polynomial given the coefficients.

Format

```
y = polyroot(c);
```

Input

c (N+1)x1 vector of coefficients of an Nth order polynomial:

$$p(z) = c[1]*z^n + c[2]*z^{n-1} + \dots + c[n]*z + c[n+1]$$

Output

y Nx1 vector, the roots of *c*.

Remarks

Zero leading terms will be stripped from *c*. When that occurs the order of *y* will be the order of the polynomial after the leading zeros have been stripped.

c[1] need not be normalized to unity.

Source

poly.src

pop

See Also

[polymake](#), [polychar](#), [polymult](#), [polyeval](#)

pop

Purpose

Provides access to a last-in, first-out stack for matrices.

Format

```
pop b;  
pop a;
```

Remarks

This is used with [gosub](#), [goto](#), and [return](#) statements with parameters. It permits passing parameters to subroutines or labels, and returning parameters from subroutines.

The [gosub](#) syntax allows an implicit [push](#) statement. This syntax is almost the same as that of a standard [gosub](#), except that the matrices to be [push](#)'ed "into the subroutine" are in parentheses following the label name. The matrices to be [push](#)'ed back to the main body of the program are in parentheses following the [return](#) statement. The only limit on the number of matrices that can be passed to and from subroutines in this way is the amount of room on the stack.

No matrix expressions can be executed between the (implicit) [push](#) and the [pop](#). Execution of such expressions will alter what is on the stack.

Matrices must be [pop](#)'ped in the reverse order that they are [push](#)'ed, therefore in the statements:

```
goto label(x,y,z);  
.  
.  
.  
label:  
pop c;  
pop b;  
pop a;
```

After the code above:

```
c = z  
b = y  
a = x
```

Note that there must be a separate `pop` statement for each matrix popped.

See Also

[gosub](#), [goto](#), [return](#)

powerM

Purpose

Returns the power n of a matrix A , as the matrix product of n copies of A .

Format

$B = \text{powerM}(A, n);$

Input

A	$N \times N$ square matrix
n	Scalar, the power or exponent.

powerM

Output

B	$N \times N$ square matrix, the power of a matrix A
-----	---

Remarks

If $n = 0$, then the function will return an identity matrix.

Example

```
A = { 1 2,
      3 4 };

//Compute power matrix
B = powerM(A, 3);

print "A = " A;
print ;
print "B = " B;
```

After the code above:

A=	
1.0000000	2.0000000
3.0000000	4.0000000
B=	
37.000000	54.000000
81.000000	118.00000

See Also

[crossprd](#)

pqqgwin

Purpose

Sets the graphics viewer mode. NOTE: This function is for use only with the deprecated PQG graphics.

Library

pgraph

Format

```
pqqgwin one;  
pqqgwin many;
```

Remarks

If you call:

```
pqqgwin one
```

only a single viewer will be used. If you call

```
pqqgwin many
```

a new viewer will be used for each graph.

pqqgwin manual and **pqqgwin auto** are supported for backwards compatibility, **manual = one**, **auto = many**.

Example

```
pqqgwin many;
```

Source

pgraph.src

previousindex

See Also

[setvwrmode](#)

previousindex

Purpose

Returns the index of the previous element or subarray in an array.

Format

pi = **previousindex**(i , o);

Input

i	Mx1 vector of indices into an array, where $M \leq N$.
o	Nx1 vector of orders of an N-dimensional array.

Output

pi	Mx1 vector of indices, the index of the previous element or subarray in the array corresponding to o .
------	--

Remarks

previousindex will return a scalar error code if the index cannot be decremented.

Example

```
orders = {3,4,5,6,7};  
a = areshape(1,orders);
```



```
orders = getorders(a);
ind = { 2,3,1 };
ind = previousindex(ind,orders);
```

After the code above, *ind* is equal to:

```
      2
ind = 2
      5
```

In this example, **previousindex** decremented *ind* to index the previous 6x7 subarray in array *a*.

See Also

[nextindex](#), [loopnextindex](#), [walkindex](#)

princomp

Purpose

Computes principal components of a data matrix.

Format

```
{ p, v, a } = princomp(x, j);
```

Input

<i>x</i>	NxK data matrix, N>K, full rank.
<i>j</i>	scalar, number of principal components to be computed (<i>j</i> ≤ K).

p

princomp

Output

p	$N \times J$ matrix of the first j principal components of x in descending order of amount of variance explained.
v	$J \times 1$ vector of fractions of variance explained.
a	$J \times K$ matrix of factor loadings, such that:

$$x = p \cdot a + \text{error}.$$

Exampe

```
//Create matrix with percent return
//of 4 stocks over 11 time periods
pcnt_return = { 0.0646  1.2326  0.0508 -0.0346,
                -0.1632  0.1806  0.1104  0.1276,
                1.3477  1.3347  0.1424  0.0159,
                -0.4465 -0.5691 -0.1524 -0.1719,
                1.6232  1.4690 -0.0192  0.0979,
                0.3381  0.5307  0.0610  0.0374,
                -0.0383  0.2556  0.0370  0.0518,
                0.4493  0.3140  0.0177  0.1001,
                0.5896  0.0542  0.1991  0.2669,
                -0.2218  0.3772  0.1189  0.1234,
                1.1778 -0.0464 -0.1282  0.2171 };
```

```
//Compute: all 4 principal component vectors,
//          percent variance explained
//          matrix of factor loadings
{ p, v, a } = princomp(pcnt_return, 4);
```

After the code above:

```
p =  0.2662 -0.6077  0.0965 -0.2951      v = 0.8394      a
    = 2.4244  2.3264  0.1321  0.2227
```

print

```
0.0059 -0.1702 -0.3938 -0.1569 0.1436
0.9506 -0.9977 -0.1402 0.1566
0.5631 -0.0350 0.0953 0.6146 0.0144
0.0317 0.0153 -0.2757 -0.3420
-0.2170 0.0709 0.4012 -0.0219 0.0026
0.0208 -0.0188 0.1455 -0.1162
0.6491 0.0491 0.2359 -0.2269
0.1823 -0.1085 -0.0554 0.0444
0.0456 -0.1485 -0.1299 -0.1762
0.1624 0.0654 -0.1032 -0.1584
0.1445 0.2677 -0.6520 0.2598
0.0337 -0.3008 -0.3926 -0.2472
0.2447 0.6267 -0.0115 -0.5214
```

From the results above, we can see that approximately 83.9% of the variance in the *pcnt_return* is included in the first principal component vector and another 14.36% is included in the second principal component.

Remarks

Adapted from a program written by Mico Loretan.

The algorithm is based on Theil, Henri "Principles of Econometrics." Wiley, NY, 1971, 46-56.

print

Purpose

Prints matrices, arrays, strings and string arrays to the screen and/or auxiliary output.

Format

```
print list_of_expressions;
print [[/typ]] [[/fmted]] [[/mf]] [[/jnt]] list_of_expressions
```

print

```
[[,:]];
```

Input

<i>list_ of_ expres sions</i>	any GAUSS expressions that produce matrices, arrays, stings, or string arrays and/or names of variables to print, separated by spaces.
<i>/typ</i>	literal, symbol type flag. <i>/mat,</i> <i>/sa,</i> <i>/str</i>
	Indicate which symbol types you are setting the output format for: matrices and arrays (<i>/mat</i>), string arrays (<i>/sa</i>), and/or strings (<i>/str</i>). You can specify more than one <i>/ typ</i> flag; the format will be set for all types indicated. If no <i>/ typ</i> flag is listed, print assumes <i>/mat</i> .
<i>/fmted</i>	literal, enable formatting flag. <i>/on,</i> <i>/off</i>
	Enable/disable formatting. When formatting is disabled, the contents of a variable are dumped to the screen in a "raw" format. <i>/off</i> is currently supported only for strings. "Raw" format for strings means that the entire string is printed, starting at the current cursor position. When formatting is enabled for strings, they are handled the same as string arrays. This shouldn't be too surprising, since a string is actually a 1x1 string array.
<i>/mf</i>	literal, matrix format. It controls the way rows of a matrix are



separated from one another. The possibilities are:

<code>/m0</code>	no delimiters before or after rows when printing out matrices.
<code>/m1 or /mb1</code>	print 1 carriage return/line feed pair before each row of a matrix with more than 1 row.
<code>/m2 or /mb2</code>	print 2 carriage return/line feed pairs before each row of a matrix with more than 1 row.
<code>/m3 or /mb3</code>	print "Row 1", "Row 2"...before each row of a matrix with more than one row.
<code>/ma1</code>	print 1 carriage return/line feed pair after each row of a matrix with more than 1 row.
<code>/ma2</code>	print 2 carriage return/line feed pairs after each row of a matrix with more than 1 row.
<code>/a1</code>	print 1 carriage return/line feed pair after each row of a matrix.
<code>/a2</code>	print 2 carriage return/line feed pairs after each row of a matrix.
<code>/b1</code>	print 1 carriage return/line feed pair before each row of a matrix.
<code>/b2</code>	print 2 carriage return/line feed pairs before each row of a matrix.
<code>/b3</code>	print "Row 1", "Row 2"... before each row of a matrix.

`/jnt` literal, controls justification, notation, and the trailing

character.

Right-Justified

/rd

Signed decimal number in the form `[[-]]#####.####`, where `#####` is one or more decimal digits. The number of digits before the decimal point depends on the magnitude of the number, and the number of digits after the decimal point depends on the precision. If the precision is 0, no decimal point will be printed.

/re

Signed number in the form `[[-]]#.##E±###`, where `#` is one decimal digit, `##` is one or more decimal digits depending on the precision, and `###` is three decimal digits. If precision is 0, the form will be `[[-]]#E±###` with no decimal point printed.

/ro

This will give a format like */rd* or */re* depending on which is most compact for the number being printed. A format like */re* will be used only if the exponent value is less than -4 or greater than the precision. If a */re* format is used, a decimal point will always appear. The precision signifies the number of significant digits displayed.

/rz

This will give a format like */rd* or

/re depending on which is most compact for the number being printed. A format like */re* will be used only if the exponent value is less than -4 or greater than the precision. If a */re* format is used, trailing zeros will be suppressed and a decimal point will appear only if one or more digits follow it. The precision signifies the number of significant digits displayed.

Left-Justified

/ld

Signed decimal number in the form `[[-]] #####.####`, where `#####` is one or more decimal digits. The number of digits before the decimal point depends on the magnitude of the number, and the number of digits after the decimal point depends on the precision. If the precision is 0, no decimal point will be printed. If the number is positive, a space character will replace the leading minus sign.

/le

Signed number in the form `[[-]]#.##E±####`, where `#` is one decimal digit, `##` is one or more decimal digits depending on the precision, and `####` is three decimal digits. If precision is 0, the form will be `[[-]]#E±####` with no decimal point printed. If the number is positive, a space character will replace the

`/lo`

leading minus sign.
This will give a format like `/ld` or `/le` depending on which is most compact for the number being printed. A format like `/le` will be used only if the exponent value is less than -4 or greater than the precision. If a `/le` format is used, a decimal point will always appear. If the number is positive, a space character will replace the leading minus sign. The precision specifies the number of significant digits displayed.

`/lz`

This will give a format like `/ld` or `/le` depending on which is most compact for the number being printed. A format like `/le` will be used only if the exponent value is less than -4 or greater than the precision. If a `/le` format is used, trailing zeros will be suppressed and a decimal point will appear only if one or more digits follow it. If the number is positive, a space character will replace the leading minus sign. The precision specifies the number of significant digits displayed.

Trailing Character

The following characters can be added to the `/ jnt` parameters above to control the trailing character if any:

	format /rdn
1,3;	
s	The number will be followed immediately by a space character. This is the default.
c	The number will be followed immediately by a comma.
t	The number will be followed immediately by a tab character.
n	No trailing character.
	The default when GAUSS is first started is:
	format /ml /ro 16,8;
::	Double semicolons following a <code>print</code> statement will suppress the final carriage return/line feed.

Remarks

The list of expressions MUST be separated by spaces. In `print` statements, because a space is the delimiter between expressions, NO SPACES are allowed inside expressions unless they are within index brackets, quotes, or parentheses.

The printing of special characters is accomplished by the use of the backslash (\) within double quotes. The options are:

\b	backspace (ASCII 8)
\e	escape (ASCII 27)

print

<code>\f</code>	form feed (ASCII 12)
<code>\g</code>	beep (ASCII 7)
<code>\l</code>	line feed (ASCII 10)
<code>\r</code>	carriage return (ASCII 13)
<code>\t</code>	tab (ASCII 9)
<code>\####</code>	the character whose ASCII value is "####" (decimal).

Thus, `\13\10` is a carriage return/line feed sequence. The first three digits will be picked up here. So if the character to follow a special character is a digit, be sure to use three digits in the escape sequence. For example: `\0074` will be interpreted as 2 characters (ASCII 7, "4")

An expression with no assignment operator is an implicit `print` statement.

If **output on** has been specified, then all subsequent `print` statements will be directed to the auxiliary output as well as the window. (See `output`.) The `locate` statement has no effect on what will be sent to the auxiliary output, so all formatting must be accomplished using tab characters or some other form of serial output.

If the name of the symbol to be printed is prefixed with a `$`, it is assumed that the symbol is a matrix of characters.

```
print $x;
```

Note that **GAUSS** makes no distinction between matrices containing character data and those containing numeric data, so it is the responsibility of the user to use functions which operate on character matrices only on those matrices containing character data.

These matrices of character strings have a maximum of 8 characters per element. A precision of 8 or more should be set when printing out character matrices or the elements will be truncated.

Complex numbers are printed with the sign of the imaginary half separating them and an "i" appended to the imaginary half. Also, the current field width setting (see `format`) refers to the width of field for each half of the number, so a complex number printed with a field of 8 will actually take (at least) 20 spaces to print.

`print`'ing a sparse matrix results in a table of the non-zero values contained in the sparse matrix, followed by their corresponding row and column indices, respectively.

A `print` statement by itself will cause a blank line to be printed:

```
print;
```

print

Examples

Example 1: Print a matrix

```
x = { 1 2,  
      3 4 };  
print x;
```

returns:

```
1.0000000    2.0000000  
3.0000000    4.0000000
```

Example 2: Print an expression

```
x = 3;  
print (x + 2);
```

returns:

```
5.0000000
```

Notice the parentheses in the code above. Remember that `print` statements in **GAUSS** take a space separated list of items to print. The parentheses tell **GAUSS** to first evaluate the expression and then print the result. Without the parentheses (i.e. `print x + 2;`), the statement would tell **GAUSS** to print a list of three items (first print `x`, then print `+`, and finally print `2`). Since the second item in that list is an operator (the `+` sign), an error will occur.

Example 3

```
x = rndn(3,3);  
format /rd 16,8;  
print x;
```

returns:

```
0.14357994 -1.39272762 -0.91942414
0.51061645 -0.02332207 -0.02511298
-1.04675893 -1.04988540 0.07992059
```

Example 4: Scientific notation

```
format /re 12,2;
print x;
```

returns:

```
1.44E-001 -1.39E+000 -9.19E-001
5.11E-001 -2.33E-002 -2.51E-002
-1.55E+000 -1.05E+000 7.99E-002
```

Example 5: Append commas

```
x = rndn(3,3);
format /rd 16,8;
print x;
```

returns:

```
0.14357994, -1.39272762, -0.91942414,
0.51061645, -0.02332207, -0.02511298,
-1.04675893, -1.04988540, 0.07992059,
```

Example 6: Add row numbers

```
print /rd /m3 x;
```

returns: (NOTE: this example does not specify the precision and spacing, so you may see more decimal places printed if that is your default setting)

```
Row 1
    0.14    -1.39    -0.92
```

printdos

```
Row 2
      0.51      -0.02      -0.03
Row 3
     -1.55     -1.05      0.08
```

Example 7: Printing character data

Character data is text inside a **GAUSS** matrix. To print elements of a matrix as characters, you need to prepend the dollar sign (\$) to the name of the variable you want to print. In most cases, string arrays are recommended over character matrices..

```
let x = AGE PAY SEX;
format /m1 8,8;
print $x;
```

```
AGE
PAY
SEX
```

See Also

[printfm](#), [printdos](#)

printdos

Purpose

Prints a string to the standard output.

Format

```
printdos s;
```

Input

s string to be printed to the standard output.

Remarks

This function is useful for printing messages to the screen when **screen off** is in effect. The output of this function will not go to the auxiliary output.

This function was used in the past to send escape sequences to the `ansi.sys` device driver on DOS. It still works on some terminals.

Example

```
printfdos "\27[7m"; /* set for reverse video */
printfdos "\27[0m"; /* set for normal text */
```

See Also

[print](#), [printfm](#), [screen](#)

printfm

Purpose

Prints a matrix using a different format for each column of the matrix.

Format

```
y = printfm(x, mask, fmt);
```

Input

<i>x</i>	NxK matrix which is to be printed and which may contain both character and numeric data.
<i>mask</i>	LxM matrix, ExE conformable with <i>x</i> , containing ones and zeros, which is used to specify whether the particular row, column, or element is to be printed as a

p

printfm

	character (0) or numeric (1) value.
<i>fmt</i>	Kx3 or 1x3 matrix where each row specifies the format for the respective column of <i>x</i> .

Output

<i>y</i>	scalar, 1 if the function is successful and 0 if it fails.
----------	--

Remarks

The mask is applied to the matrix *x* following the rules of standard element-by-element operations. If the corresponding element of *mask* is 0, then that element of *x* is printed as a character string of up to 8 characters. If *mask* contains a 1, then that element of *x* is assumed to be a double precision floating point number.

The contents of *fmt* are as follows:

$[K, 1]$	format string,	a string 8 characters maximum.
$[K, 2]$	field width,	a number < 80.
$[K, 3]$	precision,	a number < 17.

The format strings correspond to the `format` slash commands as follows:

<code>/rdn</code>	<code>"*.*1f"</code>
<code>/ren</code>	<code>"*.*1E"</code>
<code>/ron</code>	<code>"#*.*1G"</code>
<code>/rzn</code>	<code>"*.*1G"</code>
<code>/ldn</code>	<code>"- *.*1f"</code>

```

/len      "- *.*lE"
/lon      "-# *.*lG"
/lzn      "- *.*lG"

```

Complex numbers are printed with the sign of the imaginary half separating them and an "i" appended to the imaginary half. The field width refers to the width of field for each half of the number, so a complex number printed with a field of 8 will actually take (at least) 20 spaces to print.

If the precision = 0, the decimal point will be suppressed.

The format string can be a maximum of 8 characters and is appended to a % sign and passed directly to the **fprintf** function in the standard C language I/O library. The *lf*, etc., are case sensitive. If you know C, you will easily be able to use this.

If you want special characters to be printed after *x*, then include them as the last characters of the format string. For example:

```

"*.*lf,"      right-justified decimal followed by a comma.
"-*.*s "      left-justified string followed by a space.
"*.*lf"       right-justified decimal followed by nothing.

```

If you want the beginning of the field padded with zeros, then put a "0" before the first "*" in the format string:

```

"0*.*lf"      right-justified decimal.

```

Example

Here is an example of **printfm** being used to print a mixed numeric and character matrix:

```

let x[4,3] = "AGE" 5.12345564 2.23456788

```

printfmt

```
"PAY" 1.23456677 1.23456789
"SEX" 1.14454345 3.44718234
"JOB" 4.11429432 8.55649341;

let mask[1,3] = 0 1 1;      /* character numeric numeric */
let fmt[3,3] = "-*. *s " 8 8 /* first column format */
"*. *lf," 10 3              /* second column format */
"*. *le " 12 4;             /* third column format */

d = printfm(x,mask,fmt);
```

The output looks like this:

```
AGE 5.123, 2.2346E+00
PAY 1.235, 1.2346E+00
SEX 1.145, 3.4471E+00
JOB 4.114, 8.5564E+00
```

When the column of *x* to be printed contains all character elements, use a format string of `"*. *s"` if you want it right-justified, or `"-*. *s"` if you want it left-justified. If the column is mixed character and numeric elements, then use the correct numeric format and **printfm** will substitute a default format string for those elements in the column that are character.

Remember, the mask value controls whether an element will be printed as a number or a character string.

See Also

[print](#), [printdos](#)

printfmt

Purpose

Prints character, numeric, or mixed matrix using a default format controlled by the functions **formatcv** and **formatnv**.

Format

```
y = printfmt(x, mask);
```

Input

x	NxK matrix which is to be printed.
$mask$	scalar, 1 if x is numeric or 0 if x is character. - or - 1xK vector of 1's and 0's. The corresponding column of x will be printed as numeric where $mask = 1$ and as character where $mask = 0$.

Output

y	scalar, 1 if the function is successful and 0 if it fails.
-----	--

Remarks

Default format for numeric data is: `''*.*lg '' 16 8`

Default format for character data is: `''*.*s '' 8 8`

Example

```
c1 = { "age", "height", "weight" };  
c2 = { 31, 70, 160 };  
  
//Horizontally concatenate c1 and c2  
c = c1~c2;  
  
//Print 'c' as numeric data
```

printfmt

```
print c;

//Print 'c' as character data
print $c;

//Print column 1 of 'c' as character data and column 2 as
//numeric data
//Note: call disregards the return value
mask = { 0 1 };
call printfmt(c, mask);
```

The output from the three different print statements will be:

+DEN	31.000000
+DEN	70.000000
+DEN	160.00000

age	
height	
weight	

age	31
height	70
weight	160

Only the final print statement from **printfmt** correctly prints both columns.

Source

gauss.src

Globals

__fmtcv, __fmtnv

See Also

[formatcv](#), [formatnv](#)

proc

Purpose

Begins the definition of a multi-line recursive procedure. Procedures are user-defined functions with local or global variables.

Format

```
proc nrets = name(arglist);  
proc name(arglist);
```

Input

<i>nrets</i>	constant, number of objects returned by the procedure. If <i>nrets</i> is not explicitly given, the default is 1. Legal values are 0 to 1023. The <code>ret</code> statement is used to return values from a procedure.
<i>name</i>	literal, name of the procedure. This name will be a global symbol.
<i>arglist</i>	a list of names, separated by commas, to be used inside the procedure to refer to the arguments that are passed to the procedure when the procedure is called. These will always be local to the procedure, and cannot be accessed from outside the procedure or from other procedures.

Remarks

A procedure definition begins with the `proc` statement and ends with the `endp` statement.

An example of a procedure definition is:

proc

```
proc dog(x,y,z); /* procedure declaration */
local a,b;      /* local variable declarations */
    a = x .* x;
    b = y .* y;
    a = a ./ x;
    b = b ./ y;
    z = z .* z;
    z = inv(z);
    retp(a'b*z); /* return with value of a'b*z */
endp;          /* end of procedure definition */
```

Procedures can be used just as if they were functions intrinsic to the language. Below are the possible variations depending on the number of items the procedure returns.

Returns 1 item:

```
y = dog(i,j,k);
```

Returns multiple items:

```
{ x,y,z } = cat(i,j,k);
```

Returns no items:

```
fish(i,j,k);
```

If the procedure does not return any items or you want to discard the returned items:

```
call
dog(i,j,k);
```

Procedure definitions may not be nested.

For more details on writing procedures, see **PROCEDURES AND KEYWORDS, CHAPTER 1**.

See Also

[keyword](#), [call](#), [endp](#), [local](#), [retp](#)

prodc

Purpose

Computes the products of all elements in each column of a matrix.

Format

```
y = prodc(x);
```

Input

x	$N \times K$ matrix.
-----	----------------------

Output

y	$K \times 1$ matrix containing the products of all elements in each column of x .
-----	---

Remarks

To find the products of the elements in each row of a matrix, transpose before applying **prodc**. If x is complex, use the bookkeeping transpose (`.'`).

To find the products of all of the elements in a matrix, use the **vecr** function before applying **prodc**.

psi

Example

```
x = { 1 2 3,
      4 5 6,
      7 8 9 };

y = prodc(x);
```

The code above assigns *y* to be equal to:

```
      28
y =   80
     162
```

See Also

[sumc](#), [meanc](#), [stdc](#)

psi

Purpose

Computes the Psi (or Digamma) function.

Format

$f = \mathbf{psi}(z);$

Input

z NxK matrix; *z* may be complex.

Output

f NxK matrix.

p

Remarks

This program uses the analytical derivative of the log of the Lanczos series approximation for the Gamma function.

References

1. C. Lanczos, SIAM JNA 1, 1964. pp. 86-96.
2. Y. Luke, "The Special ... approximations," 1969 pp. 29-31.
3. Y. Luke, "Algorithms ... functions," 1977.
4. J. Spouge, SIAM JNA 31, 1994. pp. 931.
5. W. Press, "Numerical Recipes."
6. S. Chang, "Computation of special functions," 1996.
7. Original code by Paul Godfrey

putarray

Purpose

Puts a contiguous subarray into an N-dimensional array and returns the resulting array.

Format

```
y = putarray(a, loc, src);
```

Input

<i>a</i>	N-dimensional array.
<i>loc</i>	Mx1 vector of indices into the array to locate the subarray of interest, where M is a value from 1 to N.

putarray

<i>src</i>	[N-M]-dimensional array, matrix, or scalar.
------------	---

Output

<i>y</i>	N-dimensional array.
----------	----------------------

Remarks

If *loc* is an Nx1 vector, then *src* must be a scalar. If *loc* is an [N-1]x1 vector, then *src* must be a 1-dimensional array or a 1xL vector, where L is the size of the fastest moving dimension of the array. If *loc* is an [N-2]x1 vector, then *src* must be a KxL matrix, or a KxL 2-dimensional array, where K is the size of the second fastest moving dimension.

Otherwise, if *loc* is an Mx1 vector, then *src* must be an [N-M]-dimensional array, whose dimensions are the same size as the corresponding dimensions of array *a*.

Example

```
//Create a 2x3x4x5x6 dimensional array with unspecified
//contents
a = arrayalloc(2|3|4|5|6,0);

//Create a 4x5x6 dimensional array with all elements equal
//to 5
src = arrayinit(4|5|6,5);

loc = { 2,1 };
a = putarray(a,loc,src);
```

This example sets the contiguous 4x5x6 subarray of *a* beginning at [2,1,1,1,1] to the array *src*, in which each element is set to the specified value 5.

See Also

[setarray](#)

putf

Purpose

Writes the contents of a string to a file.

Format

```
ret = putf(filename, str, start, len, mode, append);
```

Input

<i>filename</i>	string, name of output file.
<i>str</i>	string to be written to <i>filename</i> . All or part of <i>str</i> may be written out.
<i>start</i>	scalar, beginning position in <i>str</i> of output string.
<i>len</i>	scalar, length of output string.
<i>mode</i>	scalar, output mode, (0) ASCII or (1) binary.
<i>append</i>	scalar, file write mode, (0) overwrite or (1) append.

Output

<i>ret</i>	scalar, return code.
0	normal return
1	null file name
2	file open error
3	file write error
4	output string too long
5	null output string, or illegal <i>mode</i> value
6	illegal <i>append</i> value
16	(1) append specified but file did not exist; file

putvals

was created (warning only)

Remarks

If *mode* is set to (1) binary, a string of length *len* will be written to *filename*. If *mode* is set to (0) ASCII, the string will be output up to length *len* or until **putf** encounters a ^Z (ASCII 26) in *str*. The ^Z will not be written to *filename*.

If *append* is set to (0) overwrite, the current contents of *filename* will be destroyed. If *append* is set to (1) append, *filename* will be created if it does not already exist.

If an error occurs, **putf** will either return an error code or terminate the program with an error message, depending on the **trap** state. If bit 2 (the 4's bit) of the trap flag is 0, **putf** will terminate with an error message. If bit 2 of the trap flag is 1, **putf** will return an error code. The value of the trap flag can be tested with **trapchk**.

Source

putf.src

See Also

[getf](#)

putvals

Purpose

Inserts values into a matrix or N-dimensional array.

Format

```
y=putvals(x, inds, vals);
```

Input

<i>x</i>	MxK matrix or N-dimensional array.
<i>inds</i>	LxD matrix of indices, specifying where the new values are to be inserted, where D is the number of dimensions in <i>x</i> .
<i>vals</i>	Lx1 vector, new values to insert.

Output

<i>y</i>	MxK matrix or N-dimensional array, copy of <i>x</i> containing the new values in <i>vals</i> .
----------	--

Remarks

If *x* is a vector, *inds* should be an Lx1 vector. If *x* is a matrix, *inds* should be an Lx2 matrix. Otherwise if *x* is an N-dimensional array, *inds* should be an LxN matrix.

putvals allows you to insert multiple values into a matrix or N-dimensional array at one time. This could also be accomplished using indexing inside a [for](#) loop.

Example

```
x = { -0.8750  0.3616  0.6032 -0.3974,
      0.7644 -1.8509 -0.2703 -0.8190,
      0.7886  1.2678 -1.4998 -0.5876,
      0.6639 -0.7972  1.2713  0.1896,
      0.6303  0.7879 -0.7451 -0.5419 };
inds = { 1 1, 2 4, 3 2, 3 4, 5 3 };
v = seqa(1,1,5);
y = putvals(x,inds,v);
```

After the code above:

pvCreate

```
      1.000  0.362  0.603 -0.397      1.00
      0.764 -1.851 -0.270  2.000      2.00
y = 0.789  3.000 -1.500  4.000  v = 3.00
      0.664 -0.797  1.271  0.190      4.00
      0.630  0.788  5.000 -0.542      5.00
```

pvCreate

Purpose

Returns an initialized instance of structure of type **PV**.

Format

```
p1 = pvCreate();
```

Output

p1 an instance of structure of type **PV**

Example

```
//Declare 'p1' as an instance of a 'PV' structure
struct PV p1;

//Fill in 'p1' with default values
p1 = pvCreate();
```

Source

pv.src

pvGetIndex

Purpose

Gets row indices of a matrix in a parameter vector.

Format

```
id = pvGetIndex(p1, nm1);
```

Input

<i>p1</i>	an instance of structure of type PV .
<i>nm1</i>	name or row number of matrix.

Output

<i>id</i>	Kx1 vector, row indices of matrix described by <i>nm1</i> in parameter vector.
-----------	--

Source

pv.src

pvGetParNames

Purpose

Generates names for parameter vector stored in structure of type **PV**.

Include

pv.sdf

pvGetParNames

Format

```
s = pvGetParNames(p1);
```

Input

<i>p1</i>	an instance of structure of type PV .
-----------	--

Output

<i>s</i>	Kx1 string array, names of parameters.
----------	--

Remarks

If the vector in the structure of type **PV** was generated with matrix names, the parameter names will be concatenations of the matrix name with row and column numbers of the parameters in the matrix. Otherwise the names will have a generic prefix with concatenated row and column numbers.

Example

```
//Define PV structure
#include pv.sdf
//Declare 'p1' as an instance of a 'PV' structure
struct PV p1;

//Initialize 'p1' with default values
p1 = pvCreate;

//Data to pack into the 'PV' struct
x = { 1 2,
      3 4 };

//1's indicate an element to pack into the structure
```

```
//0's indicate elements to NOT pack into the structure
mask = { 1 0,
         0 1 };

//Pack values of 'x' selected by 'mask' into 'pi' and name
//this resulting vector, 'P'
p1 = pvPackm(p1,x,"P",mask);

print pvGetParNames(p1);
```

Since *mask* has ones in the [1,1] and [2,2] locations, the code above, produces:

```
P[1,1]
P[2,2]
```

Source

pv.src

pvGetParVector

Purpose

Retrieves parameter vector from structure of type **PV**.

Include

pv.sdf

Format

```
p = pvGetParVector(p1);
```

pvGetParVector

Input

p1 an instance of structure of type **PV**.

Output

p Kx1 vector, parameter vector.

Remarks

Matrices or portions of matrices (stored using a mask) are stored in the structure of type **PV** as a vector in the *p* member.

Example

```
//Define 'PV' structure
#include pv.sdf
//Declare 'p1' as an instance of a 'PV' structure
struct PV p1;

//Initialize 'p1' with default values
p1 = pvCreate;

x = { 1 2,
      3 4 };

//1's indicate elements to pack into 'p1' parameter vector
mask = { 1 1,
         0 0 };

p1 = pvPackm(p1,x,"X",mask);

print pvUnpack(p1,"X");
```

p

pvUnpack returns the entire value of x that was packed in. Therefore, the print statement above, produces:

```
1.000 2.000
3.000 4.000
```

```
print
pvGetParVector (p1) ;
```

pvGetParVector returns only those elements indicated by the *mask* variable and therefore the `print` statement above, returns:

```
1.000
2.000
```

Source

pv.src

pvLength

Purpose

Returns the length of a parameter vector.

Format

```
 $n$  = pvLength( $p1$ );
```

Input

$p1$ an instance of structure of type **PV**.

pvList

Output

n	scalar, length of parameter vector in $p1$.
-----	--

Source

`pv.src`

pvList

Purpose

Retrieves names of packed matrices in structure of type **PV**.

Format

```
 $n$  = pvList( $p1$ );
```

Input

$p1$	an instance of structure of type PV .
------	--

Output

n	Kx1 string vector, names of packed matrices.
-----	--

Source

`pv.src`

p

pvPack

Purpose

Packs general matrix into a structure of type **PV** with matrix name.

Include

pv.sdf

Format

```
p1 = pvPack(p1, x, nm);
```

Input

<i>p1</i>	an instance of structure of type PV .
<i>x</i>	MxN matrix or N-dimensional array.
<i>nm</i>	string, name of matrix/array.

Output

<i>p1</i>	an instance of structure of type PV .
-----------	--

Example

Example 1: Basic usage

```
//Create starting parameter vector
start_vals = { 0,
               1,
               1 };

//Declare 'p1' as an instance of a 'PV' structure
```

pvPacki

```
struct PV p1;

//Initialize 'p1' with default settings
p1 = pvCreate();

//Add a variable named 'b' in 'p1' containing the data from
'start_vals'
p1 = pvPack(p1, start_vals, "b");
```

The data can be extracted using the **pvUnpack** command:

```
b_out = pvUnpack(p1, "b");
print b_out;
```

The code above, should return the following output:

```
0
1
1
```

Source

pv.src

See Also

[pvPackm](#), [pvPacks](#), [pvUnpack](#)

pvPacki

Purpose

Packs general matrix or array into a **PV** instance with name and index.

Include

pv.sdf

Format

```
p1 = pvPacki(p1, x, nm, i);
```

Input

<i>p1</i>	an instance of structure of type PV .
<i>x</i>	MxN matrix or N-dimensional array.
<i>nm</i>	string, name of matrix or array, or null string.
<i>i</i>	scalar, index of matrix or array in lookup table.

Output

<i>p1</i>	an instance of structure of type PV .
-----------	--

Example

```
//Define the 'PV' structure
#include pv.sdf

y = rndn(100,1);
x = rndn(100,5);

//Declare 'p1' as an instance of a 'PV' structure
struct PV p1;

//Initialize 'p1' with default values
p1 = pvCreate;

//Pack the variables in with a variable name and an index
p1 = pvPacki(p1,y,"Y",1);
p1 = pvPacki(p1,x,"X",2);
```

These matrices can be extracted using the **pvUnpack** command, indicating the variable to unpack either by index or by variable name:

pvPackm

```
//Unpack variables by index
y = pvUnpack(p1,1);
x = pvUnpack(p1,2);

//Unpack variables by variable name
y = pvUnpack(p1,"Y");
x = pvUnpack(p1,"X");
```

See Also

[pvPack](#), [pvUnpack](#)

pvPackm

Purpose

Packs general matrix into a structure of type **PV** with a mask and matrix name.

Include

pv.sdf

Format

```
p1 = pvPackm(p1, x, nm, mask);
```

Input

p

<i>p1</i>	an instance of structure of type PV .
<i>x</i>	MxN matrix or N-dimensional array.
<i>nm</i>	string, name of matrix/array or N-dimensional array.
<i>mask</i>	MxN matrix, mask matrix of zeros and ones.

Output

p1 an instance of structure of type **PV**.

Remarks

The *mask* argument allows storing a selected portion of a matrix into the packed vector. The ones in *mask* indicate an element to be stored in the packed matrix. When the matrix is unpacked (using **pvUnpack**) the elements corresponding to the zeros are restored. Elements corresponding to the ones come from the packed vector which may have been changed.

If the mask is all zeros, the matrix or array is packed with the specified elements in the second argument but no elements of the matrix or array are entered into the parameter vector. When unpacked the matrix or array in the second argument is returned without modification.

Example

```
#include pv.sdf
struct PV p1;
p1 = pvCreate;

x = { 1 2,
      3 4 };

mask = { 1 0,
         0 1 };

p1 = pvPackm(p1,x,"X",mask);

print pvUnpack(p1,1);
```

pvPackmi

```
1.000 2.000
3.000 4.000
```

```
p1 = pvPutParVector(p1, 5 | 6);

print pvUnpack(p1, "X");
```

```
5.000 2.000
3.000 6.000
```

Source

pv.src

pvPackmi

Purpose

Packs general matrix or array into a **PV** instance with a mask, name, and index.

Include

pv.sdf

Format

```
p1 = pvPackmi(p1, x, nm, mask, i);
```

Input

<i>p1</i>	an instance of structure of type PV .
<i>x</i>	MxN matrix or N-dimensional array.

<i>nm</i>	string, matrix or array name.
<i>mask</i>	MxN matrix or N-dimensional array, <i>mask</i> of zeros and ones.
<i>i</i>	scalar, index of matrix or array in lookup table.

Output

<i>p1</i>	an instance of structure of type PV .
-----------	--

Remarks

The *mask* allows storing a selected portion of a matrix into the parameter vector. The ones in the *mask* matrix indicate an element to be stored in the parameter matrix. When the matrix is unpacked (using **pvUnpackm**) the elements corresponding to the zeros are restored. Elements corresponding to the ones come from the parameter vector.

If the mask is all zeros, the matrix or array is packed with the specified elements in the second argument but no elements of the matrix or array are entered into the parameter vector. When unpacked the matrix or array in the second argument is returned without modification.

Example

```
#include pv.sdf

struct PV p1;
p1 = pvCreate;

x = { 1 2,
      3 4 };

mask = { 1 0,
```

pvPacks

```
0 1 };  
  
p1 = pvPackmi (p1,x,"X",mask,1);  
  
print pvUnpack (p1,1);
```

```
1.000 2.000  
3.000 4.000
```

```
p1 = pvPutParVector (p1,5|6);  
  
print pvUnpack (p1,1);
```

```
5.000 2.000  
3.000 6.000
```

See Also

[pvPackm](#), [pvUnpack](#)

pvPacks

Purpose

Packs symmetric matrix into a structure of type **PV**.

Include

`pv.sdf`

Format

```
p1 = pvPacks(p1, x, nm);
```

Input

<i>p1</i>	an instance of structure of type PV .
<i>x</i>	MxM symmetric matrix.
<i>nm</i>	string, matrix name.

Output

<i>p1</i>	an instance of structure of type PV .
-----------	--

Remarks

pvPacks does not support the packing of arrays.

Example

```
#include pv.sdf

struct PV p1;
p1 = pvCreate;

x = { 1 2,
      2 1 };

p1 = pvPacks(p1, x, "A");
p1 = pvPacks(p1, eye(2), "I");
```

These matrices can be extracted using the **pvUnpack** command:

```
print pvUnpack(p1, "A");
```

```
1.000 2.000
2.000 1.000
```

```
print pvUnpack(p1, "I");
```

p

pvPacksi

```
1.000 0.000
0.000 1.000
```

Source

`pv.src`

See Also

[pvPacksm](#), [pvUnpack](#)

pvPacksi

Purpose

Packs symmetric matrix into a **PV** instance with matrix name and index.

Include

`pv.sdf`

Format

```
p1 = pvPacksi(p1, x, nm, i);
```

Input

P

<i>p1</i>	an instance of structure of type PV .
<i>x</i>	MxM symmetric matrix.
<i>nm</i>	string, matrix name.
<i>i</i>	scalar, index of matrix in lookup table.

Output

p1 an instance of structure of type **PV**.

Remarks

pvPacksi does not support the packing of arrays.

Example

```
#include pv.sdf

struct PV p1;
p1 = pvCreate;

x = { 1 2, 2 1 };

p1 = pvPacksi(p1,x, "A",1);
p1 = pvPacksi(p1, eye(2), "I",2);
```

These matrices can be extracted using the **pvUnpack** command.

```
print
pvUnpack(p1,1);
```

```
1.000 2.000
2.000 1.000
```

```
print
pvUnpack(p1,2);
```

```
1.000 0.000
0.000 1.000
```

pvPacksm

See Also

[pvPacks](#), [pvUnpack](#)

pvPacksm

Purpose

Packs symmetric matrix into a structure of type **PV** with a mask.

Include

`pv.sdf`

Format

```
p1 = pvPacksm(p1, x, nm, mask);
```

Input

<i>p1</i>	an instance of structure of type PV .
<i>x</i>	MxM symmetric matrix.
<i>nm</i>	string, matrix name.
<i>mask</i>	MxM matrix, mask matrix of zeros and ones.

Output

<i>p1</i>	an instance of structure of type PV .
-----------	--

Remarks

pvPacksm does not support the packing of arrays.

The mask allows storing a selected portion of a matrix into the packed vector. The ones in *mask* indicate an element to be stored in the packed matrix. When the matrix is unpacked (using **pvUnpack**) the elements corresponding to the zeros are restored. Elements corresponding to the ones come from the packed vector which may have been changed.

Only the lower left portion of the *mask* matrix is used, and only the lower left portion of the *x* matrix is stored in the packed vector.

If the mask is all zeros, the matrix is packed with the specified elements in the second argument but no elements of the matrix are entered into the parameter vector. When unpacked the matrix in the second argument is returned without modification.

Example

```
#include pv.sdf

struct PV p1;
p1 = pvCreate;

x = { 1 2 4,
      2 3 5,
      4 5 6};

mask = { 1 0 1,
         0 1 0,
         1 0 1 };

p1 = pvPacksm(p1, x, "A", mask);

print pvUnpack(p1, "A");
```

```
1.000 2.000 4.000
2.000 3.000 5.000
4.000 5.000 6.000
```

pvPacksmi

```
p2 = pvGetParVector (p1);
```

```
print p2;
```

```
1.000
```

```
3.000
```

```
4.000
```

```
6.000
```

```
p3 = { 10, 11, 12, 13 };
```

```
p1 = pvPutParVector (p1,p3);
```

```
print pvUnpack (p1, "A");
```

```
10.000  2.000 12.000
```

```
2.000 11.000  5.000
```

```
12.000  5.000 13.000
```

Source

pv.src

pvPacksmi

Purpose

Packs symmetric matrix into a **PV** instance with a mask, matrix name, and index.

Include

pv.sdf

Format

```
p1 = pvPacksmi(p1, x, nm, mask, i);
```

Input

<i>pl</i>	an instance of structure of type PV .
<i>x</i>	MxM symmetric matrix.
<i>nm</i>	string, matrix name.
<i>mask</i>	MxM matrix, symmetric mask matrix of zeros and ones.
<i>i</i>	scalar, index of matrix in lookup table.

Output

<i>pl</i>	an instance of structure of type PV .
-----------	--

Remarks

pvPacksmi does not support the packing of arrays.

The *mask* allows storing a selected portion of a matrix into the parameter vector. The ones in the *mask* matrix indicate an element to be stored in the parameter vector. When the matrix is unpacked (using **pvUnpackm**) the elements corresponding to the zeros are restored. Elements corresponding to the ones come from the parameter vector.

Only the lower left portion of the *mask* matrix is used, and only the lower left portion of the *x* matrix is stored in the packed vector.

If the mask is all zeros, the matrix is packed with the specified elements in the second argument but no elements of the matrix are entered into the parameter vector. When unpacked the matrix in the second argument is returned without modification.

Example

```
#include pv.sdf
```

pvPacksmi

```
struct PV p1;
p1 = pvCreate;

x = { 1 2 4,
      2 3 5,
      4 5 6};

mask = { 1 0 1,
         0 1 0,
         1 0 1 };

p1 = pvPacksmi(p1,x, "A",mask,1);

print pvUnpack(p1,1);
```

```
1.000 2.000 4.000
2.000 3.000 5.000
4.000 5.000 6.000
```

```
p2 = pvGetParVector(p1);

print p2;
```

```
1.000
3.000
4.000
6.000
```

```
p3 = { 10, 11, 12, 13 };
p1 = pvPutParVector(p1,p3);

print pvUnpack(p1,1);
```

p

```
10.000  2.000 12.000
 2.000 11.000  5.000
12.000  5.000 13.000
```

See Also

[pvPacksm](#), [pvUnpack](#)

pvPutParVector

Purpose

Inserts parameter vector into structure of type **PV**.

Include

pv.sdf

Format

$p1 = \text{pvPutParVector}(p1, \ p);$

Input

$p1$	an instance of structure of type PV .
p	Kx1 vector, parameter vector.

Output

$p1$	an instance of structure of type PV .
------	--

Remarks

Matrices or portions of matrices (stored using a *mask*) are stored in the structure of

pvPutParVector

type **PV** as a vector in the `p` member.

Example

```
#include pv.sdf

struct PV p1;
p1 = pvCreate;

x = { 1 2 4,
      2 3 5,
      4 5 6};

mask = { 1 0 1,
         0 1 0,
         1 0 1 };

//Packed as square matrix
p1 = pvPackm(p1,x,"A",mask);

print pvUnpack(p1,"A");
```

```
1.000 2.000 4.000
2.000 3.000 5.000
4.000 5.000 6.000
```

```
p3 = { 10, 11, 12, 13, 14 };
p1 = pvPutParVector(p1,p3);

print pvUnpack(p1,"A");
```

```
10.000  2.000 11.000
 2.000 12.000  5.000
13.000  5.000 14.000
```

p

Source

pv.src

pvTest

Purpose

Tests an instance of structure of type **PV** to determine if it is a proper structure of type **PV**.

Format

```
i = pvTest(p1);
```

Input

p1 an instance of structure of type **PV**.

Output

i scalar, if 0, *p1* is a proper structure of type **PV**, else if 1, an improper or uninitialized structure of type **PV**.

Source

pv.src

pvUnpack

Purpose

Unpacks matrices stored in a structure of type **PV**.

pvUnpack

Format

```
x = pvUnpack(p1, m);
```

Input

<i>p1</i>	an instance of structure of type PV .
<i>m</i>	string, name of matrix, or integer, index of matrix.

Output

<i>x</i>	MxN general matrix or MxM symmetric matrix or N-dimensional array.
----------	--

Source

pv.src



q

QNewton

Purpose

Optimizes a function using the BFGS descent algorithm.

Format

{ *x*, *f*, *g*, *ret* } = QNewton(&*fct*, *start*);

Input

<i>&fct</i>	pointer to a procedure that computes the function to be minimized. This procedure must have one input argument, a vector of parameter values, and one output argument, the value of the function evaluated at the input vector of parameter values.
<i>start</i>	Kx1 vector, start values.

Global Input

<i>_qn_</i> <i>RelGradTol</i>	scalar, convergence tolerance for relative gradient of estimated coefficients. Default = 1e-5.
<i>_qn_</i> <i>GradProc</i>	scalar, pointer to a procedure that computes the gradient of the function with respect to the parameters. This procedure must have a single input argument, a Kx1 vector of parameter values, and a single output argument, a Kx1 vector of gradients

QNewton

	of the function with respect to the parameters evaluated at the vector of parameter values. If <code>_qn_GradProc</code> is 0, QNewton uses gradp .
<code>_qn_MaxIters</code>	scalar, maximum number of iterations. Default = 1e+5. Termination can be forced by pressing C on the keyboard.
<code>_qn_PrintIters</code>	scalar, if 1, print iteration information. Default = 0. Can be toggled during iterations by pressing P on the keyboard.
<code>_qn_ParNames</code>	Kx1 vector, labels for parameters.
<code>_qn_PrintResults</code>	scalar, if 1, results are printed.

Output

<code>x</code>	Kx1 vector, coefficients at the minimum of the function.
<code>f</code>	scalar, value of function at minimum.
<code>g</code>	Kx1 vector, gradient at the minimum of the function.
<code>ret</code>	scalar, return code. 0 normal convergence 1 forced termination 2 max iterations exceeded 3 function calculation failed 4 gradient calculation failed 5 step length calculation failed 6 function cannot be evaluated at initial parameter values

Remarks

If you are running in terminal mode, **GAUSS** will not see any input until you press ENTER. Pressing C on the keyboard will terminate iterations, and pressing P will toggle iteration output.

To reset global variables for this function to their default values, call **QNewtonSet**.

Example

This example computes maximum likelihood coefficients and standard errors for a Tobit model:

```

/**qnewton.e - a Tobit model***/
//Get data
z = loadadd("tobit");
b0 = { 1, 1, 1, 1 };
{b,f,g,retcode} = qnewton(&lpr,b0);

//Covariance matrix of parameters
h = hessp(&lpr,b);
output file = qnewton.out reset;

print "Tobit Model";
print;
print "coefficients standard errors";
print b~sqrt(diag(invpd(h)));

output off;

//Log-likelihood proc
proc lpr(b);
    local s,m,u;
    s = b[4];
    if s <= 1e-4;

```

QNewtonmt

```
        retp(error(0));
    endif;
    m = z[:,2:4]*b[1:3,:];
    u = z[:,1] ./= 0;
    retp(-sumc(u.*lnpdfmvn(z[:,1]-m,s) + (1-u).*(ln(cdfnc
(m/sqrt(s))))));
endp;
```

produces:

```
Tobit Model
coefficients standard errors

    0.010417884  0.080220019
   -0.20805753  0.094551107
   -0.099749592 0.080006676
    0.65223067  0.099827309
```

Source

qnewton.src

QNewtonmt

Purpose

Minimize an arbitrary function.

Format

```
out = QNewtonmt(&fct, par);
out = QNewtonmt(&fct, par, ...);
out = QNewtonmt(&fct, par, c);
out = QNewtonmt(&fct, par, ..., c);
```

Input

<i>&fct</i>	pointer to a procedure that computes the function to be minimized. This procedure must have at least one input argument, an instance of a PV structure containing the parameters. And, one output argument, the value of the function evaluated at the input vector of parameter values.						
<i>par</i>	Optional, an instance of a PV structure. The <i>par</i> instance is passed to the user-provided procedure pointed to by <i>&fct</i> . <i>par</i> is constructed using the pvpPack functions.						
<i>...</i>	Optional extra arguments. These arguments are passed untouched to the user-provided objective function, by QNewtonmt .						
<i>c</i>	an instance of a QNewtonmtControl structure. Normally an instance is initialized by calling QNewtonmtControlCreate and members of this instance can be set to other values by the user. For an instance named <i>c</i> , the members are: <table> <tr> <td><i>c.CovType</i></td><td>scalar, if 1, ML covariance matrix, else if 2, QML covariance matrix is computed. Default is 0, no covariance matrix.</td></tr> <tr> <td><i>c.GradProc</i></td><td>scalar, pointer to a procedure that computes the gradient of the function with respect to the parameters. Default = ., i.e., no gradient procedure has been provided.</td></tr> <tr> <td><i>c.MaxIters</i></td><td>scalar, maximum number of iterations. Default = 1e+5.</td></tr> </table>	<i>c.CovType</i>	scalar, if 1, ML covariance matrix, else if 2, QML covariance matrix is computed. Default is 0, no covariance matrix.	<i>c.GradProc</i>	scalar, pointer to a procedure that computes the gradient of the function with respect to the parameters. Default = ., i.e., no gradient procedure has been provided.	<i>c.MaxIters</i>	scalar, maximum number of iterations. Default = 1e+5.
<i>c.CovType</i>	scalar, if 1, ML covariance matrix, else if 2, QML covariance matrix is computed. Default is 0, no covariance matrix.						
<i>c.GradProc</i>	scalar, pointer to a procedure that computes the gradient of the function with respect to the parameters. Default = ., i.e., no gradient procedure has been provided.						
<i>c.MaxIters</i>	scalar, maximum number of iterations. Default = 1e+5.						

QNewtonmt

<i>c.MaxTries</i>	scalar, maximum number of attempts in random search. Default = 100.
<i>c.relGradTol</i>	scalar, convergence tolerance for gradient of estimated coefficients. Default = 1e-5. When this criterion has been satisfied QNewtonmt exits the iterations.
<i>c.randRadius</i>	scalar, If zero, no random search is attempted. If nonzero, it is the radius of the random search. Default = .001.
<i>c.output</i>	scalar, if nonzero, results are printed. Default = 0.
<i>c.PrintIters</i>	scalar, if nonzero, prints iteration information. Default = 0.
<i>c.disableKey</i>	scalar, if nonzero, keyboard input disabled

Output

<i>out</i>	an instance of an QNewtonmtOut structure. For an instance named <i>out</i> , the members are:
<i>out.par</i>	instance of a PV structure containing the parameter estimates will be placed in the member matrix <i>out.par</i> .
<i>out.fct</i>	scalar, function evaluated at <i>x</i> .
<i>out.retcode</i>	scalar, return code: 0 normal convergence.

	1	forced exit.
	2	maximum number of iterations exceeded.
	3	function calculation failed.
	4	gradient calculation failed.
	5	Hessian calculation failed.
	6	line search failed.
	7	error with constraints.
	8	function complex.
<i>out.moment</i>		KxK matrix, covariance matrix of parameters, if <i>c.covType</i> > 0.
<i>out.hessian</i>		KxK matrix, matrix of second derivatives of objective function with respect to parameters.

Remarks

There is one required user-provided procedure, the one computing the objective function to be minimized, and another optional functions, the gradient of the objective function.

These functions have one input argument that is an instance of type struct **PV**. On input to the call to **QNewtonmt**, the first argument contains starting values for the parameters. The arguments following the **PV** structure (except for the optional control structure) contain any required data.

The **PV** structures are set up using the **PV** pack procedures, **pvpPack**, **pvpPackm**, **pvpPacks**, and **pvpPacksm**. These procedures allow for setting up a parameter vector in a variety of ways.

QNewtonmt

For example, we might have the following objective function for fitting a nonlinear curve to data:

```
proc (1) = Micherlitz(struct PV par1, y, x);  
    local p0,e,s2;  
    p0 = pvUnpack(par1, "parameters");  
    e = y - p0[1] - p0[2]*exp(-p0[3] * x);  
    retp(-lnpdfmvn(e,e'e/rows(e)));  
endp;
```

In this example the dependent and independent variables are passed to the procedure as the second and third arguments to the procedure.

If the objective function is the negative of a proper log-likelihood, and if *c.covType* is set to 1, the covariance matrix of the parameters is computed and returned in *out-moment*, and standard errors, t-statistics and probabilities are printed if *c.output* = 1.

If the objective function returns the negative of a vector of log-likelihoods, and if *c.covType* is set to 2, the quasi-maximum likelihood (QML) covariance matrix of the parameters is computed.

Example

```
//Define function to be minimized  
//The first input is a PV structure containing the para-  
meters  
//The following arguments contain data, other than the para-  
meters,  
//which is needed by the function  
proc (1) = Micherlitz(struct PV par1, y, x);  
    local p0,e,s2;  
    p0 = pvUnpack(par1, "parameters");
```

```

    e = y - p0[1] - p0[2]*exp(-p0[3] * x);
    retp(-lnpdfmvn(e,e'e/rows(e)));
endp;

//Create extra data needed by objective function
y = { 3.183,
      3.059,
      2.871,
      2.622,
      2.541,
      2.184,
      2.110,
      2.075,
      2.018,
      1.903,
      1.770,
      1.762,
      1.550 };

x = seqa(1,1,13);

//Declare 'par' to be a PV structure
struct PV par;

//Set PV defaults in 'par'
par = pvCreate();

//Add a variable named 'parameters' to par with a 3x1
//vector of starting values
par = pvPack(par, 1|1|0, "parameters");

//Declare 'out' to be a QNewtonmtOut structure
//to hold data returned by QNewtonmt
struct QNewtonmtout out;

//Minimize the 'Micherlitz' function

```

QNewtonmtControlCreate

```
out = QNewtonmt(&Micherlitz,par,y,x);

//Get returned parameters from the output structure
parms = pvGetParVector(out.par);

//Print returned parameters
print parms;
```

The code above should return the following output:

```
0.96312060
2.5189989
0.10305485
```

Source

qnewtonmt.src

See Also

[QNewtonmtControlCreate](#), [QNewtonmtOutCreate](#)

QNewtonmtControlCreate

Purpose

Creates default **QNewtonmtControl** structure.

Include

qnewtonmt.sdf

Format

```
c = QNewtonmtControlCreate();
```

Output

<code>c</code>	instance of QNewtonmtControl structure with members set to default values.
----------------	---

Source

`qnewtonmt.src`

See Also

[QNewtonmt](#)

QNewtonmtOutCreate

Purpose

Creates default **QNewtonmtOut** structure.

Format

```
c = QNewtonmtOutCreate();
```

Output

<code>c</code>	instance of QNewtonmtOut structure with members set to default values.
----------------	---

Source

`qnewtonmt.src`

See Also

[QNewtonmt](#)

QNewtonSet

QNewtonSet

Purpose

Resets global variables used by **QNewton** to default values.

Format

```
QNewtonSet;
```

Source

qnewton.src

QProg

Purpose

Solves the quadratic programming problem.

Format

```
{ x, u1, u2, u3, u4, u5 } = QProg(start, q, r, a, b, c, d, bnds);
```

Input

<i>start</i>	Kx1 vector, start values.
<i>q</i>	KxK matrix, symmetric model matrix.
<i>r</i>	Kx1 vector, model constant vector.
<i>a</i>	MxK matrix, equality constraint coefficient matrix, or scalar 0, no equality constraints.
<i>b</i>	Mx1 vector, equality constraint constant vector, or scalar 0, will be expanded to Mx1 vector of zeros.

<i>c</i>	NxK matrix, inequality constraint coefficient matrix, or scalar 0, no inequality constraints.
<i>d</i>	Nx1 vector, inequality constraint constant vector, or scalar 0, will be expanded to Nx1 vector of zeros.
<i>bnds</i>	Kx2 matrix, bounds on <i>x</i> , the first column contains the lower bounds on <i>x</i> , and the second column the upper bounds. If scalar 0, the bounds for all elements will default to $\pm 1e200$.

Global Input

<i>_qprog_ maxit</i>	scalar, maximum number of iterations. Default = 1000.
--------------------------	---

Output

<i>x</i>	Kx1 vector, coefficients at the minimum of the function.
<i>u1</i>	Mx1 vector, Lagrangian coefficients of equality constraints.
<i>u2</i>	Nx1 vector, Lagrangian coefficients of inequality constraints.
<i>u3</i>	Kx1 vector, Lagrangian coefficients of lower bounds.
<i>u4</i>	Kx1 vector, Lagrangian coefficients of upper bounds.
<i>ret</i>	scalar, return code. <div> <div>0</div> <div>successful termination</div> </div> <div> <div>1</div> <div>max iterations exceeded</div> </div> <div> <div>2</div> <div>machine accuracy is insufficient to maintain decreasing function values</div> </div> <div> <div>3</div> <div>model matrices not conformable</div> </div>

QProgmt

< 0 active constraints inconsistent

Remarks

QProg solves the standard quadratic programming problem:

$$\min \frac{1}{2}x'Qx - x'R$$

subject to constraints,

$$\begin{aligned} Ax &= B \\ Cx &\leq D \end{aligned}$$

and bounds,

$$x_{low} \leq x \leq x_{up}$$

Source

qprog.src

QProgmt

Purpose

Solves the quadratic programming problem.

Format

```
qOut = QProgmt(qIn );
```


Input

<i>qIn</i>	instance of a qprogMTIn structure containing the following members:
<i>qIn.start</i>	Kx1 vector, start values.
<i>qIn.q</i>	KxK matrix, symmetric model matrix.
<i>qIn.r</i>	Kx1 vector, model constant vector.
<i>qIn.a</i>	MxK matrix, equality constraint coefficient matrix, or scalar 0, no equality constraints.
<i>qIn.b</i>	Mx1 vector, equality constraint constant vector, or scalar 0, will be expanded to Mx1 vector of zeros.
<i>qIn.c</i>	NxK matrix, inequality constraint coefficient matrix, or scalar 0, no inequality constraints.
<i>qIn.d</i>	Nx1 vector, inequality constraint constant vector, or scalar 0, will be expanded to Nx1 vector of zeros.
<i>qIn.bounds</i>	Kx2 matrix, bounds on <i>qOut.x</i> , the first column contains the lower bounds on <i>qOut.x</i> , and the second column the upper bounds. If scalar 0, the bounds for all elements will default to $\pm 1e200$.
<i>qIn.maxit</i>	scalar, maximum number of iterations. Default = 1000.

Output

<i>qOut</i>	instance of a qprogMTOut structure containing the following members:
<i>qOut.x</i>	Kx1 vector, coefficients at the minimum of the function.
<i>qOut.lagrange</i>	instance of a qprogMTLagrange structure containing the following members:
<i>qOut.lagrange.lineq</i>	Mx1 vector, Lagrangian coefficients of equality constraints.
<i>qOut.lagrange.linineq</i>	Nx1 vector, Lagrangian coefficients of inequality constraints.
<i>qOut.lagrange.bounds</i>	Kx2 matrix, Lagrangian coefficients of bounds, the first column contains the lower bounds and the second the upper bounds.
<i>qOut.ret</i>	scalar, return code. 0 successful termination 1 max iterations exceeded 2 machine accuracy is insufficient to maintain decreasing function values 3 model matrices not conformable < 0 active constraints

inconsistent

Remarks

QProgmt solves the standard quadratic programming problem:

$$\min \frac{1}{2}x'Qx - x'R$$

subject to constraints,

$$Ax = Bx \leq D$$

and bounds,

$$x_{low} \leq x \leq x_{up}$$

Source

qprogmt.src

See Also

[QProgmtInCreate](#)

QProgmtInCreate

Purpose

Creates an instance of a structure of type **QProgmtInCreate** with the *maxit* member set to a default value.

Include

qprogmt.sdf

qqr

Format

```
s = QProgmtInCreate();
```

Output

s	instance of structure of type QProgmtIn.
---	--

Source

qprogmt.src

See Also

[QProgmt](#)

qqr

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix x , such that:
 $X = Q_I R$

Format

```
{ q1, r } = qqr(x);
```

Input

x	NxP matrix.
---	-------------

Output

q1	NxK unitary matrix, K = min (N,P).
r	KxP upper triangular matrix.

Remarks

Given X , there is an orthogonal matrix Q such that $Q'X$ is zero below its diagonal, i.e.,

$$Q'X = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 Q_2]$$

where Q_1 has P columns, then

$$X = Q_1 R$$

is the QR decomposition of X . If X has linearly independent columns, R is also the Cholesky factorization of the moment matrix of X , i.e., of $X'X$.

If you want only the R matrix, see the function **qr**. Not computing Q_1 can produce significant improvements in computing time and memory usage.

An unpivoted R matrix can also be generated using **cholup**:

```
r = cholup(zeros(cols(x), cols(x)), x);
```

For linear equation or least squares problems, which require Q_2 for computing residuals and residual sums of squares, see **olsqr** and **qtyr**.

For most problems an explicit copy of Q_1 or Q_2 is not required. Instead one of the following, $Q'Y$, QY , $Q_1'Y$, Q_1Y , $Q_2'Y$, or Q_2Y , for some Y , is required. These cases are all handled by **qtyr** and **gyr**. These functions are available because Q and Q_1 are typically very large matrices while their products with Y are more manageable.

If $N < P$, the factorization assumes the form:

qqre

$$Q'X = [R_1 \ R_2]$$

where R_1 is a $P \times P$ upper triangular matrix and R_2 is $P \times (N-P)$. Thus Q is a $P \times P$ matrix and R is a $P \times N$ matrix containing R_1 and R_2 . This type of factorization is useful for the solution of underdetermined systems. However, unless the linearly independent columns happen to be the initial rows, such an analysis also requires pivoting (see **qre** and **qrep**).

Source

qqr.src

See Also

[qre](#), [qrep](#), [qtyr](#), [qtyre](#), [qtyrep](#), [qyr](#), [qyre](#), [qyrep](#), [olsqr](#)

qqre

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix x , such that:
 $X[:,E] = Q_1 R$

Format

$\{ q1, r, \ e \} = \text{qqre}(x);$

Input

x $N \times P$ matrix.

Output

$q1$	$N \times K$ unitary matrix, $K = \mathbf{min}(N,P)$.
r	$K \times P$ upper triangular matrix.
e	$P \times 1$ permutation vector.

Remarks

Given $X[:,E]$, where E is a permutation vector that permutes the columns of X , there is an orthogonal matrix Q such that $Q'X[:,E]$ is zero below its diagonal, i.e.,

$$Q'R[:,E] = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X[:,E] = Q_1 R$$

is the QR decomposition of $X[:,E]$.

If you want only the R matrix, see **qqre**. Not computing Q_1 can produce significant improvements in computing time and memory usage.

If X has rank P , then the columns of X will not be permuted. If X has rank $M < P$, then the M linearly independent columns are permuted to the front of X by E . Partition the permuted X in the following way:

$$X[., E] = [X_1 \ X_2]$$

where X is $N \times M$ and X_2 is $N \times (P-M)$. Further partition R in the following way:

$$R = \begin{bmatrix} R_{11} & R_{12} \\ 0 & 0 \end{bmatrix}$$

where R_{11} is $M \times M$ and R_{12} is $M \times (P-M)$. Then

$$A = R^{-1}_{11} R_{12}$$

and

$$X_2 = X_1 A$$

that is, A is an $M \times (P-M)$ matrix defining the linear combinations of X_2 with respect to X_1 .

If $N < P$, the factorization assumes the form:

$$Q^T X = [R_1 \ R_2]$$

where R_1 is a $P \times P$ upper triangular matrix and R_2 is $P \times (N-P)$. Thus Q is a $P \times P$ matrix and R is a $P \times N$ matrix containing R_1 and R_2 . This type of factorization is useful for the solution of underdetermined systems. For the solution of

$$X[:, E] b = Y$$

it can be shown that

$$b = \text{qrsol}(Q'Y, R1) | \text{zeros}(N-P, 1);$$

The explicit formation here of Q , which can be a very large matrix, can be avoided by using the function **qtyre**.

For further discussion of QR factorizations see the remarks under **qqr**.

Source

qqr.src

See Also

qqr**qqr**, [qtyre](#), [olsqr](#)

qqrep

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix x , such that:
 $X[:, E] = Q_1 R$

Format

$\{ q1, r, e \} = \text{qqrep}(x, pvt);$

Input

x NxP matrix.

qqrep

pvt	<p>Px1 vector, controls the selection of the pivot columns:</p> <ul style="list-style-type: none">if $pvt[i] > 0$, $x[i]$ is an initial columnif $pvt[i] = 0$, $x[i]$ is a free columnif $pvt[i] < 0$, $x[i]$ is a final column <p>The initial columns are placed at the beginning of the matrix and the final columns are placed at the end. Only the free columns will be moved during the decomposition.</p>
-------	---

Output

$q1$	NxK unitary matrix, $K = \mathbf{min}(N,P)$.
r	KxP upper triangular matrix.
e	Px1 permutation vector.

Remarks

Given $X[.,E]$, where E is a permutation vector that permutes the columns of X , there is an orthogonal matrix Q such that $Q'X[.,E]$ is zero below its diagonal, i.e.,

$$Q'R[.,E] = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X[., E] = Q_1 R$$

is the QR decomposition of $X[., E]$.

qqrep allows you to control the pivoting. For example, suppose that x is a data set with a column of ones in the first column. If there are linear dependencies among the columns of x , the column of ones for the constant may get pivoted away. This column can be forced to be included among the linearly independent columns using `pvt`.

If you want only the R matrix, see **qrep**. Not computing Q_1 can produce significant improvements in computing time and memory usage.

Source

`qqr.src`

See Also

[qqr](#), [qre](#), [olsqr](#)

qr

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix x , such that:
 $X = Q_1 R$

Format

```
 $r = \mathbf{qr}(x);$ 
```

Input

x

NxP matrix.

qr

Output

r

KxP upper triangular matrix, $K = \min(N,P)$.

Remarks

qr is the same as **qqr** but doesn't return the Q_I matrix. If Q_I is not wanted, **qr** will save a significant amount of time and memory usage, especially for large problems.

Given X , there is an orthogonal matrix Q such that $Q'X$ is zero below its diagonal, i.e.,

$$Q'X = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X = Q_1 R$$

is the QR decomposition of X . If X has linearly independent columns, R is also the Cholesky factorization of the moment matrix of X , i.e., of $X'X$.

qr does not return the Q_I matrix because in most cases it is not required and can be very large. If you need the Q_I matrix, see the function **qqr**. If you need the entire Q matrix, call **qyr** with Y set to a conformable identity matrix.

For most problems $Q'Y$, $Q_I'Y$, or QY , Q_IY , for some Y , are required. For these cases see **qtyr** and **qyr**.

For linear equation or least squares problems, which require Q_2 for computing residuals and residual sums of squares, see **olsqr**.

If $N < P$, the factorization assumes the form:

$$Q'X = [R_1 R_2]$$

where R_1 is a $P \times P$ upper triangular matrix and R_2 is $P \times (N-P)$. Thus Q is a $P \times P$ matrix and R is a $P \times N$ matrix containing R_1 and R_2 . This type of factorization is useful for the solution of underdetermined systems. However, unless the linearly independent columns happen to be the initial rows, such an analysis also requires pivoting (see **qre** and **qrep**).

Source

`qr.src`

See Also

[qqr](#), [qrep](#), [qtypre](#)

qre

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix x , such that:
 $X[:,E] = Q_1 R$

Format

`{ r, e } = qre(x);`

gre

Input

x	$N \times P$ matrix.
-----	----------------------

Output

r	$K \times P$ upper triangular matrix, $K = \mathbf{min}(N,P)$.
e	$P \times 1$ permutation vector.

Remarks

gre is the same as **qgre** but doesn't return the Q_I matrix. If Q_I is not wanted, **gre** will save a significant amount of time and memory usage, especially for large problems.

Given $X[:,E]$, where E is a permutation vector that permutes the columns of x , there is an orthogonal matrix Q such that $Q'X[:,E]$ is zero below its diagonal, i.e.,

$$Q'X[:, E] = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_I has P columns, then

$$X[:, E] = Q_1 R$$

is the QR decomposition of $X[:,E]$.

qre does not return the Q_I matrix because in most cases it is not required and can be very large. If you need the Q_I matrix, see the function **qqre**. If you need the entire Q matrix, call **qyre** with Y set to a conformable identity matrix. For most problems $Q'Y$, $Q_I'Y$, or QY , Q_IY , for some y , are required. For these cases see **qtyre** and **qyre**.

If X has rank P , then the columns of X will not be permuted. If X has rank $M < P$, then the M linearly independent columns are permuted to the front of X by E . Partition the permuted X in the following way:

$$X[., E] = [X_1 \ X_2]$$

where X_1 is $N \times M$ and X_2 is $N \times (P-M)$. Further partition R in the following way:

where R_{11} is $M \times M$ and R_{12} is $M \times (P-M)$. Then

$$A = R_{11}^{-1} R_{12}$$

and

$$X_2 = X_1 A$$

that is, A is an $M \times (P-N)$ matrix defining the linear combinations of X_2 with respect to X_1

If $N < P$ the factorization assumes the form:

$$Q'X = [R_1 \ R_2]$$

qrep

where R_1 is a $P \times P$ upper triangular matrix and R_2 is $P \times (N-P)$. Thus Q is a $P \times P$ matrix and R is a $P \times N$ matrix containing R_1 and R_2 . This type of factorization is useful for the solution of underdetermined systems. For the solution of

$$X[:, E]b = Y$$

it can be shown that

```
b = qrsol(Q'Y, R1) | zeros(N-P,1);
```

The explicit formation here of Q , which can be a very large matrix, can be avoided by using the function **qtyre**.

For further discussion of QR factorizations see the remarks under **qqr**.

Source

qr.src

See Also

[qqr](#), [olsqr](#)

qrep

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix X , such that:

$$X[:, E] = Q_1 R$$

Format

```
{ r, e } = qrep(X, pvt);
```


Input

X	$N \times P$ matrix.
pvt	$P \times 1$ vector, controls the selection of the pivot columns: if $pvt[i] > 0$, $X[i]$ is an initial column. if $pvt[i] = 0$, $X[i]$ is a free column. if $pvt[i] < 0$, $X[i]$ is a final column. The initial columns are placed at the beginning of the matrix and the final columns are placed at the end. Only the free columns will be moved during the decomposition.

Output

r	$K \times P$ upper triangular matrix, $K = \min(N, P)$.
e	$P \times 1$ permutation vector.

Remarks

qrep is the same as **qqrep** but doesn't return the Q_I matrix. If Q_I is not wanted, **qrep** will save a significant amount of time and memory usage, especially for large problems.

Given $X[:, E]$, where E is a permutation vector that permutes the columns of X , there is an orthogonal matrix Q such that $Q'X[:, E]$ is zero below its diagonal, i.e.,

$$Q'X[:, E] = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

qrsol

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X[., E] = Q_1 R$$

is the QR decomposition of $X[., E]$.

qrep does not return the Q_1 matrix because in most cases it is not required and can be very large. If you need the Q_1 matrix, see the function **qqrep**. If you need the entire Q matrix, call **qyrep** with Y set to a conformable identity matrix. For most problems $Q'Y$, $Q_1'Y$, or QY , Q_1Y , for some Y , are required. For these cases see **qtyrep** and **qyrep**.

qrep allows you to control the pivoting. For example, suppose that X is a data set with a column of ones in the first column. If there are linear dependencies among the columns of X , the column of ones for the constant may get pivoted away. This column can be forced to be included among the linearly independent columns using **pvt**.

Source

`qr.src`

See Also

[qr](#), [gre](#), [qqrep](#)

qrsol

Purpose

Computes the solution of $Rx = b$ where R is an upper triangular matrix.

Format

```
 $x = \text{qrtsol}(b, R);$ 
```

Input

b	PxL matrix.
R	PxP upper triangular matrix.

Output

x	PxL matrix.
-----	-------------

Remarks

qrtsol applies a backsolve to $Rx = b$ to solve for x . Generally R will be the R matrix from a QR factorization. **qrtsol** may be used, however, in any situation where R is upper triangular.

Source

qrtsol.src

See Also

[qqr](#), [qr](#), [qtyr](#), [qrtsol](#)

qrtsol

Purpose

Computes the solution of $R'x = b$ where R is an upper triangular matrix.

qrtsol

Format

```
x = qrtsol(b, R);
```

Input

b	PxL matrix.
R	PxP upper triangular matrix.

Output

x	PxL matrix.
-----	-------------

Remarks

qrtsol applies a forward solve to $R'x = b$ to solve for x . Generally R will be the R matrix from a QR factorization. **qrtsol** may be used, however, in any situation where R is upper triangular. If R is lower triangular, transpose before calling **qrtsol**.

If R is not transposed, use **qrsol**.

Source

qrsol.src

See Also

qqr, **qr**, **qtyr**, **qrsol**

qtyr

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix X and returns $Q'Y$ and R .

Format

$\{ \text{qty}, r \} = \text{qtyr}(y, X);$

Input

y	NxL matrix.
X	NxP matrix.

Output

qty	NxL unitary matrix.
r	KxP upper triangular matrix, $K = \mathbf{min}(N,P)$.

Remarks

Given X , there is an orthogonal matrix Q such that $Q'X$ is zero below its diagonal, i.e.,

$$Q'X = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X = Q_1 R$$

is the QR decomposition of X . If X has linearly independent columns, R is also the Cholesky factorization of the moment matrix of X , i.e., of $X'X$. For most problems Q or Q_1 is not what is required. Rather, we require $Q'Y$ or $Q_1'Y$ where Y is an $N \times L$ matrix (if either QY or Q_1Y are required, see **qyr**). Since Q can be a very large matrix, **qtyr** has been provided for the calculation of $Q'Y$ which will be a much smaller matrix. $Q_1'Y$ will be a submatrix of $Q'Y$. In particular,

$$G = Q_1'Y = \text{qty}[1 : P, .]$$

and $Q_2'Y$ is the remaining submatrix:

$$H = Q_2'Y = \text{qty}[P + 1 : N, .]$$

Suppose that X is an $N \times K$ data set of independent variables, and Y is an $N \times 1$ vector of dependent variables. Then it can be shown that

$$b = R^{-1}G$$

and

$$s_j = \sum_{i=1}^{N-P} H_{i,j}, j = 1, 2, \dots, L$$

where b is a $P \times L$ matrix of least squares coefficients and s is a $1 \times L$ vector of residual sums of squares. Rather than invert R directly, however, it is better to apply **qrsol** to

$$Rb = Q_1'Y$$

For rank deficient least squares problems, see **qtyre** and **qtyrep**.

Example

The QR algorithm is the numerically superior method for the solution of least squares problems:

```
loadm x, y;
{ qty, r } = qtyr(y,x);
q1ty = qty[1:rows(r),.];
q2ty = qty[rows(r)+1:rows(qty),.];

//LS coefficients
b = qrsol(q1ty,r);

//Residual sums of squares
s2 = sumc(q2ty^2);
```

Source

qtyr.src

See Also

[qqr](#), [qtyre](#), [qtyrep](#), [olsqr](#)

qtyre

qtyre

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix X and returns $Q'Y$ and R .

Format

$\{ \text{qty}, r, e \} = \mathbf{qtyre}(y, x);$

Input

y	NxL matrix.
x	NxP matrix.

Output

qty	NxL unitary matrix.
r	KxP upper triangular matrix, $K = \mathbf{min}(N,P)$.
e	Px1 permutation vector.

Remarks

Given $X[.,E]$, where E is a permutation vector that permutes the columns of X , there is an orthogonal matrix Q such that $Q'X[.,E]$ is zero below its diagonal, i.e.,

$$Q'X \left[., E \right] = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X[., E] = Q_1 R$$

is the QR decomposition of $X[., E]$.

If X has rank P , then the columns of X will not be permuted. If X has rank $M < P$, then the M linearly independent columns are permuted to the front of X by E . Partition the permuted X in the following way:

$$X[., E] = [X_1 \ X_2]$$

where X_1 is $N \times M$ and X_2 is $N \times (P-M)$. Further partition R in the following way:

$$R = \begin{bmatrix} R_{11} & R_{12} \\ 0 & 0 \end{bmatrix}$$

where R_{11} is $M \times M$ and R_{12} is $M \times (P-M)$. Then

$$A = R_{11}^{-1} R_{12}$$

and

$$X_2 = X_1 A$$

qtyre

that is, A is an $M \times (P-N)$ matrix defining the linear combinations of X_2 with respect to X_1 .

For most problems Q or Q_1 is not it is required. Rather, we require $Q'Y$ or $Q_1'Y$ where Y is an $N \times L$ matrix. Since Q can be a very large matrix, **qtyre** has been provided for the calculation of $Q'Y$ which will be a much smaller matrix. $Q_1'Y$ will be a submatrix of $Q'Y$. In particular,

$$Q_1'Y = \text{qty}[1 : P, .]$$

and $Q_2'Y$ is the remaining submatrix:

$$Q_2'Y = \text{qty}[P + 1 : N, .]$$

Suppose that X is an $N \times K$ data set of independent variables and Y is an $N \times 1$ vector of dependent variables. Suppose further that X contains linearly dependent columns, i.e., X has rank $M < P$. Then define

$$C = Q_1'Y[1 : M, .]$$
$$A = R[1 : M, 1 : M]$$

and the vector (or matrix of $L > 1$) of least squares coefficients of the reduced, linearly independent problem is the solution of

$$Ab = C$$

To solve for b use **qrsol**:

```
b = qrsol (C, A) ;
```

If $N < P$, the factorization assumes the form:

$$Q'X [., E] = [R_1 R_2]$$

where R_1 is a $P \times P$ upper triangular matrix and R_2 is $P \times (N-P)$. Thus Q is a $P \times P$ matrix and R is a $P \times N$ matrix containing R_1 and R_2 . This type of factorization is useful for the solution of underdetermined systems. For the solution of

$$X[., E]b = Y$$

it can be shown that

```
b = qrsol (Q'Y, R1) | zeros (N-P, 1) ;
```

Source

qtyprep.src

See Also

[qqr](#), [qre](#), [qtyp](#)

qtyprep

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix X using a pivot vector and returns $Q'Y$ and R .

qtyrep

Format

$\{ \text{qty}, r, e \} = \mathbf{qtyrep}(y, x, \text{pvt});$

Input

y	NxL matrix.
x	NxP matrix.
pvt	Px1 vector, controls the selection of the pivot columns: if $\text{pvt}[i] > 0$, $x[i]$ is an initial column. if $\text{pvt}[i] = 0$, $x[i]$ is a free column. if $\text{pvt}[i] < 0$, $x[i]$ is a final column. The initial columns are placed at the beginning of the matrix and the final columns are placed at the end. Only the free columns will be moved during the decomposition.

Output

qty	NxL unitary matrix.
r	KxP upper triangular matrix, $K = \mathbf{min}(N,P)$.
e	Px1 permutation vector.

Remarks

Given $X[.,E]$, where E is a permutation vector that permutes the columns of X , there is an orthogonal matrix Q such that $Q'X[.,E]$ is zero below its diagonal, i.e.,

$$Q'X[:, E] = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X[:, E] = Q_1 R$$

is the QR decomposition of $X[:, E]$.

qtyrep allows you to control the pivoting. For example, suppose that X is a data set with a column of ones in the first column. If there are linear dependencies among the columns of X , the column of ones for the constant may get pivoted away. This column can be forced to be included among the linearly independent columns using *pvt*.

Source

`qtyr.src`

See Also

[qrep](#), [qtyre](#)

quantile

Purpose

Computes quantiles from data in a matrix, given specified probabilities.

quantile

Format

```
y=quantile(x, e);  
y=quantile(x, e, tp);
```

Input

x	NxK matrix of data.
e	Lx1 vector, quantile levels or probabilities.
tp	Scalar, 1, 2, ..., 9. Sample quantile type. Default is 4.

Output

y	LxK matrix, quantiles.
---	------------------------

Remarks

Let $\{x_{(1)}, \dots, x_{(n)}\}$ denote the order statistics, and let $\hat{Q}_i(p) = (1 - \gamma)x_{(j)} + \gamma x_{(j+1)}$ denotes the sample quantiles, where $\frac{j-m}{n} \leq p < \frac{j-m-1}{n}$, $m \in \mathbb{R}$, $0 \leq \gamma \leq 1$. The value of γ is a function of integer part $j = \text{floor}(pn + m)$ and fractional part $g = pn + m - j$. The m is a constant determined by sample quantile type.

Type	Definition
1	Discrete sample quantile type 1. Inverse of empirical distribution function.
2	Discrete sample quantile type 2. Similar to type 1 except that averaging at discontinuities.
3	Discrete sample quantile type 3. SAS definition, choose the nearest even order statistics.

- | | |
|---|---|
| 4 | Continuous sample quantile type 4. Interpolating the step function of definition 1. |
| 5 | Continuous sample quantile type 5. This is the value midway through each step of definition 1. |
| 6 | Continuous sample quantile type 6. The vertices divide the sample space into $n+1$ regions, each with probability $1/(n+1)$. |
| 7 | Continuous sample quantile type 7. The vertices divide the range into $n-1$ regions, and $100p\%$ of the intervals lie to the left and $100(1-p)\%$ lie to the right. |
| 8 | Continuous sample quantile type 8. The resulting sample quantile is median unbiased regardless the distribution. |
| 9 | Continuous sample quantile type 9. The resulting sample quantile is median unbiased if normal distribution. |

Example

```
//Set the rng seed for repeatable random numbers

rndseed 345567;

//Create a 1000x4 random normal matrix
x = rndn(1000,4);

//Quantile levels
e = { .025, .5, .975 };

y = quantile(x, e);

print "medians";
print y[2, .];
print;
```

quantiled

```
print "95 percentiles";  
print y[1,.];  
print y[3,.];
```

Produces the following output:

```
medians  
    -0.037801917    0.029923972   -0.010477829   -0.023937160  
  
95 percentiles  
    -2.0074122    -2.0798579    -1.9982702    -1.9605009  
     2.0437573     2.0271770     1.9025695     1.9228044
```

Source

quantile.src

quantiled

Purpose

Computes quantiles from data in a data set, given specified probabilities.

Format

```
y = quantiled(dataset, e, var);
```

Input

<i>dataset</i>	string, data set name, or NxM matrix of data.
<i>e</i>	Lx1 vector, quantile levels or probabilities.
<i>var</i>	Kx1 vector, scalar zero, string array, or formula string . If Kx1, character vector of labels selected for analysis, or numeric vector of column numbers in data set of variables

selected for analysis.

If *var* is scalar zero, all columns are selected.

If *dataset* is a matrix *var* cannot be a character vector.

If *dataset* includes variable names, then *var* could be a string array, e.g. "Height" \$| "Weight" or [formula string](#). e.g. "Height + Weight".

Output

y LxK matrix, quantiles.

Remarks

- **quantiled** will not succeed if $N * \text{minc}(e)$ is less than 1, or $N * \text{maxc}(e)$ is greater than $N - 1$. In other words, to produce a **quantile** for a level of .001, the input matrix must have more than 1000 rows.
- The supported data set types are [CSV](#), [XLS](#), [XLSX](#), [HDF5](#), [FMT](#), [DAT](#).
- For HDF5 file, the *dataset* must include [file schema](#) and both file name and data set name must be provided, e.g. **quantiled** ("h5://C:/gauss17/examples/testdata.h5/mydata", 0.5, 0).

Examples

Example 1 Use data set name

```
//Create file name with full path
file_name = getGAUSSHome() $+ "examples/fueleconomy.dat";
//Set up quantile levels
e = { .025, .5, .975 };
//Choose all variables in the data set
```

quantiled

```
var = 0;
//Compute quantiles
y = quantiled(file_name, e, var);

print "medians";
print y[2,.];
print;
print "95 percentiles";
print y[1,.];
print y[3,.];
```

produces:

```
medians
2.5000000      3.0000000

95 percentiles
1.5500000      1.4000000
4.0500000      6.2550000
```

Example 2 Use .csv file and variable index

```
//Create file name with full path
file_name = getGAUSSHome() $+ "examples/binary.csv";
//Set up quantile levels
e = { .025, .5, .975 };
//Set up variable index
var = 2|3;
//Compute quantiles
y = quantiled(file_name,e,var);

print "medians";
print y[2,.];
print;
```

```
print "95 percentiles";
print y[1,.];
print y[3,.];
```

After the above code:

```
medians
580.00000      3.3900000

95 percentiles
360.00000      2.6300000
               800.00000      4.0000000
```

Example 3 Use .xls file and [formula string](#)

```
//Create file name with full path
file_name = getGAUSSHome() $+ "examples/nba_ht_wt.xls";
//Set up quantile levels
e = { .025, .5, .975 };
//Set up formula string
var = "Height + Weight" ;
//Compute quantiles
y = quantiled(file_name,e,var);
print "Height" $~ "Weight";
print "medians";
print y[2,.];
print;
print "95 percentiles";
print y[1,.];
print y[3,.];
```

After the above code:

```
medians
220.00000      79.500000
```

qyr

```
95 percentiles
175.00000      72.000000
270.00000      84.000000
```

See also

[Formula String](#)

Source

quantile.src

qyr

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix X and returns QY and R .

Format

$\{ qY, r \} = \mathbf{qyr}(y, x);$

Input

y	NxL matrix.
X	NxP matrix.

Output

qY	NxL unitary matrix.
r	KxP upper triangular matrix, $K = \mathbf{min}(N,P)$.

Remarks

Given X , there is an orthogonal matrix Q such that $Q'X$ is zero below its diagonal, i.e.,

$$Q'X = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X = Q_1 R$$

is the QR decomposition of X . If X has linearly independent columns, R is also the Cholesky factorization of the moment matrix of X , i.e., of $X'X$.

For most problems Q or Q_1 is not what is required. Since Q can be a very large matrix, **qyr** has been provided for the calculation of QY , where Y is some $N \times L$ matrix, which will be a much smaller matrix.

If either $Q'Y$ or $Q_1'Y$ are required, see **qtyr**.

Source

`qyr.src`

See Also

[qqr](#), [qyre](#), [qyrep](#), [olsqr](#)

qyre

qyre

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix x and returns QY and R .

Format

$\{ qy, r, e \} = \text{qyre}(y, x);$

Input

y	NxL matrix.
x	NxP matrix.

Output

qy	NxL unitary matrix.
r	KxP upper triangular matrix, $K = \mathbf{min}(N,P)$.
e	Px1 permutation vector.

Remarks

Given $X[.,E]$, where E is a permutation vector that permutes the columns of X , there is an orthogonal matrix Q such that $Q'X[.,E]$ is zero below its diagonal, i.e.,

$$Q'X \left[., E \right] = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X[:, E] = Q_1 R$$

is the QR decomposition of $X[:, E]$.

For most problems Q or Q_1 is not what is required. Since Q can be a very large matrix, **qyre** has been provided for the calculation of QY , where Y is some NxL matrix, which will be a much smaller matrix.

If either $Q'Y$ or $Q_1'Y$ are required, see **qtyre**.

If $N < P$, the factorization assumes the form:

$$Q'X[:, E] = [R_1 \ R_2]$$

where R_1 is a PxP upper triangular matrix and R_2 is Px(N-P). Thus Q is a PxP matrix and R is a PxN matrix containing R_1 and R_2 .

Source

`qyr.src`

See Also

[qqr](#), [gre](#), [qyr](#)

qyrep

qyrep

Purpose

Computes the orthogonal-triangular (QR) decomposition of a matrix X using a pivot vector and returns QY and R .

Format

$\{ qY, r, e \} = \text{qyrep}(y, x, \text{pvt});$

Input

y	$N \times L$ matrix.
x	$N \times P$ matrix.
pvt	$P \times 1$ vector, controls the selection of the pivot columns: if $pvt[i] > 0$, $x[i]$ is an initial column. if $pvt[i] = 0$, $x[i]$ is a free column. if $pvt[i] < 0$, $x[i]$ is a final column. The initial columns are placed at the beginning of the matrix and the final columns are placed at the end. Only the free columns will be moved during the decomposition.

Output

qY	$N \times L$ unitary matrix.
r	$K \times P$ upper triangular matrix, $K = \text{min}(N,P)$.
e	$P \times 1$ permutation vector.

Remarks

Given $X[:, E]$, where E is a permutation vector that permutes the columns of X , there is an orthogonal matrix Q such that $Q'X[:, E]$ is zero below its diagonal, i.e.,

$$Q'X[:, E] = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

where R is upper triangular. If we partition

$$Q = [Q_1 \ Q_2]$$

where Q_1 has P columns, then

$$X[:, E] = Q_1 R$$

is the QR decomposition of $X[:, E]$.

qyrep allows you to control the pivoting. For example, suppose that X is a data set with a column of ones in the first column. If there are linear dependencies among the columns of X , the column of ones for the constant may get pivoted away. This column can be forced to be included among the linearly independent columns using `pvt`.

For most problems Q or Q_1 is not what is required. Since Q can be a very large matrix, **qyrep** has been provided for the calculation of QY , where Y is some $N \times L$ matrix, which will be a much smaller matrix.

If either $Q'Y$ or $Q_1'Y$ are required, see **qtyrep**.

If $N < P$, the factorization assumes the form:

qz

$$Q'X[.,E] = [R_1 R_2]$$

where R_1 is a PxP upper triangular matrix and R_2 is Px(N-P). Thus Q is a PxP matrix and R is a PxN matrix containing R_1 and R_2 .

Source

qyr.src

See Also

[qr](#), [qqrep](#), [qrep](#), [qtyrep](#)

qz

Purpose

Compute the complex QZ, or generalized Schur, form of a pair of real or complex general matrices with an option to sort the eigenvalues.

Format

```
{ S, T, Q, Z } = qz(A, B);  
{ S, T, Q, Z } = qz(A, B, sort_type);
```

Input

A	NxN matrix, real or complex general matrix
B	NxN matrix, real or complex general matrix

<i>sort_type</i>	Optional input, scalar or string specifying how to sort the eigenvalues. Options include:	
1	"udi"	Absolute value of the eigenvalue less than 1.0. (Unit disk inside)
2	"udo"	Absolute value of the eigenvalue greater than or equal to 1.0. (Unit disk outside)
3	"lhp"	Value of the real portion of the eigenvalue less than 0. (Left hand plane)
4	"rhp"	Value of the real portion of the eigenvalue greater than 0. (Right hand plane)
5	"ref"	Real eigenvalues first. (Complex portion less than <i>imagtol</i> see remarks section)
6	"cef"	Complex eigenvalues first. (Complex portion greater than <i>imagtol</i> see remarks section)

Output

<i>S</i>	NxN matrix, Schur form of <i>A</i>
<i>T</i>	NxN matrix, Schur form of <i>B</i>
<i>Q</i>	NxN matrix, left Schur vectors
<i>Z</i>	NxN matrix, right Schur vectors

Examples

Example 1: Basic usage

```
//For repeatable random numbers
rndseed 23434;

//Matrix dimensions
order = 4;

//Create 2 square, real matrices
A = rndn(order, order);
B = rndn(order, order);

//Perform 'QZ' decomposition
{ S, T, Q, Z } = qz(A,B);

//Calculate generalized eigenvalues
eig_vals = diag(S) ./ diag(T);

print "Generalized eigenvalues = ";
print eig_vals;

print "Absolute value of the generalized eigenvalues = ";
print abs(eig_vals);
```

The above code should return the following output:

```
Generalized eigenvalues =

    20.703871 -    1.9686543e-16i
    0.16170711 -    1.6939178e-17i
   -0.83402664 -    0.34681937i
   -0.83402664 +    0.34681937i
```

```

Absolute value of the generalized eigenvalues =

    20.703871
    0.16170711
    0.90326303
    0.90326303

```

Example 2: Ordering eigenvalues

You can order the eigenvalues, by passing in the optional third input, *sort_type*. The code below uses the same *A* and *B* variables made in the example above.

```

//Perform 'QZ' decomposition and
//reorder generalized eigenvalues, placing
//those with absolute value less than 1
//on the upper left
{ S, T, Q, Z } = qz(A, B, "udi");

//Calculate generalized eigenvalues
eig_vals = diag(S) ./ diag(T);

print "Generalized eigenvalues = ";
print (eig_vals);

print "Absolute value of the generalized eigenvalues = ";
print abs(eig_vals);

```

The code above should print out the sorted eigenvalues as we see below.

```

Generalized eigenvalues =

    0.16170711 -    1.6819697e-17i

```

```
-0.83402664 -      0.34681937i
-0.83402664 +      0.34681937i
 20.703871 -      2.1311282e-14i
```

Absolute value of the generalized eigenvalues =

```
0.16170711
0.90326303
0.90326303
20.703871
```

Remarks

- The pair of matrices S and T are in generalized complex Schur form if S and T are upper triangular and the diagonal of T contains positive real numbers.
- The real generalized eigenvalues can be computed by dividing the diagonal element of S by the corresponding diagonal element of T .
- The generalized Schur vectors Q and Z are orthogonal matrices ($Q'Q = I$ and $Z'Z = I$) that reduce A and B to Schur form:

```
S = Q'A*Z
T = Q'B*Z

A = Q*S*Z'
B = Q*T*Z'
```

- For the real generalized schur decomposition, call **lapgschur**.
- If only the generalized eigenvalues are needed, you can call **lapgeig**, or **lapgeigv**.
- By default **imagtol** is set to 2.23e-16. If your program requires *imagtol* to

be a different value, you may change it using **sysstate** case 21, like this:

```
//Set imagtol to 1e-15  
imagtol_org = sysstate(21, 1e-15);
```

Note that while the function **qz** IS threadsafe, setting *imagtol* is NOT threadsafe. Therefore, *imagtol* should not be changed inside of a [threadStat](#) or [threadBegin](#) block.

- This procedure calls the LAPACK routine ZGGES.

rank

r

rank

Purpose

Computes the rank of a matrix, using the singular value decomposition.

Format

```
 $k$  = rank( $x$ );
```

Input

x NxP matrix.

Global Input

`—svd_tol`

scalar, the tolerance used in determining if any of the singular values are effectively 0. The default value is $10e^{-13}$. This can be changed before calling the procedure.

Output

k an estimate of the rank of x . This equals the number of singular values of x that exceed a prespecified tolerance in absolute value.

Global Output

`_svderr` scalar, if not all of the singular values can be computed
`_svderr` will be nonzero.

Source

svd.src

rankindx

Purpose

Returns the vector of ranks of a vector.

Format

`y = rankindx(x, flag);`

Input

<code>x</code>	Nx1 vector.
<code>flag</code>	scalar, 1 for numeric data or 0 for character data.

Output

<code>y</code>	Nx1 vector containing the ranks of <code>x</code> . That is, the rank of the largest element is N and the rank of the smallest is 1. (To get ranks in descending order, subtract <code>y</code> from N+1).
----------------	--

readr

Remarks

rankindx assigns different ranks to elements that have equal values (ties). Missing values are assigned the lowest ranks.

Example

```
x = { 12, 4, 15, 7, 8 };  
r = rankindx(x,1);
```

After the code above, *r* is equal to:

```
      4  
      1  
r = 5  
      2  
      3
```

readr

Purpose

Reads a specified number of rows of data from a **GAUSS** data set (*.dat*) file or a **GAUSS** matrix (*.fmt*) file.

Format

```
y = readr(f1, r);
```

Input

<i>f1</i>	scalar, file handle of an open file.
<i>r</i>	scalar, number of rows to read.

Output

 y

NxK matrix, the data read from the file.

Remarks

The first time a **readr** statement is encountered, the first x rows will be read. The next time it is encountered, the next x rows will be read in, and so on. If the end of the data set is reached before x rows can be read, then only those rows remaining will be read.

After the last row has been read, the pointer is placed immediately after the end of the file. An attempt to read the file in these circumstances will cause an error message.

To move the pointer to a specific place in the file use **seekr**.

Example

```
open dt = dat1.dat;
m = 0;

do until eof(dt);
  x = readr(dt, 400);
  m = m + moment(x, 0);
endo;

dt = close(dt);
```

This code reads data from a data set 400 rows at a time. The moment matrix for each set of rows is computed and added to the sum of the previous moment matrices. The result is the moment matrix for the entire data set. **eof(dt)** returns 1 when the end of the data set is encountered.

See Also

[open](#), [create](#), [writer](#), [seekr](#), [eof](#)

real

real

Purpose

Returns the real part of x .

Format

```
 $zr = \text{real}(x);$ 
```

Input

x NxK matrix or N-dimensional array.

Output

zr NxK matrix or N-dimensional array, the real part of x .

Remarks

If x is not complex, zr will be equal to x .

Example

```
 $x = \{ \begin{matrix} 1 & 11+2i, \\ 7i & 3, \\ 2+1i & 1 \end{matrix} \};$   
 $zr = \text{real}(x);$ 
```

After the code above, x and zr are equal to:

```
 $x = \begin{matrix} 1+0i & 11+2i \\ 0+7i & 3+0i \\ 2+1i & 1+0i \end{matrix} \quad zr = \begin{matrix} 1 & 11 \\ 0 & 3 \\ 2 & 1 \end{matrix}$ 
```

See Also

[complex](#), [imag](#)

reclassify

Purpose

Replaces specified values of a matrix, array or string array

Format

```
x_new = reclassify(x, from, to);
```

Input

<i>x</i>	NxK matrix, string array or NxKxP array to be recoded (changed)
<i>from</i>	kx1 vector, or string array of values to change
<i>to</i>	kx1 vector, or string array containing the new values to be assigned to the recoded variable

Output

<i>x_new</i>	Matrix, multi-dimensional array or string array with the same dimensions as the input <i>x</i> , containing the recoded values of <i>x</i>
--------------	--

Examples

Example 1

Change instances of 1, 2 and 3 to 'low', 'medium' and 'high'.

reclassify

```
//Vector to be changed
x = { 2,
      3,
      2,
      1,
      2,
      3 };

from = { 1,
         2,
         3 };

//Create a 3x1 string array using
//string vertical concatenation operator
to = "low" $| "medium" $| "high";

x_new = reclassify(x, from, to);
print x_new ;
```

After the code above, `x_new` is equal to:

```
medium
high
medium
low
medium
high
```

Example 2

Change instances of tea types: 'black', 'green', 'oolong' to 9.95, 11.95 and 10.50, respectively.

```
string orders = { "green",
                  "green",
                  "oolong",
```

```
        "green",  
        "green",  
        "green",  
        "black" };  
  
string tea_types = { "black",  
                    "green",  
                    "oolong" };  
  
price = { 9.95, 11.95, 10.50 };  
  
order_prices = reclassify(orders, tea_types, price);  
print order_prices;
```

After the code above, *order_prices* is equal to:

```
11.95  
11.95  
10.50  
11.95  
11.95  
11.95  
9.95
```

Source

datatran.src

See Also

[code](#), [recode](#), [reclassifyCuts](#), [substute](#), [rescale](#), [dummy](#)

reclassifyCuts

reclassifyCuts

Purpose

Replaces values of a matrix or array within specified ranges

Format

```
x_new = reclassifyCuts(x, cut_pts);  
x_new = reclassifyCuts(x, cut_pts, close_right);
```

Input

<i>x</i>	NxK matrix, string array or NxKxP array to be recoded (changed)
<i>cut_pts</i>	kx1 vector, bounds of the specified ranges
<i>close_right</i>	Scalar, optional argument, 1 if the <i>cut_pts</i> should be the right end-point of the interval, or 0 if the values in <i>cut_pts</i> should start the next interval

Output

<i>x_new</i>	Matrix, multi-dimensional array or string array with the same dimensions as the input <i>x</i> , containing the recoded values of <i>x</i>
--------------	--

Examples

Example 1: Basic sequence


```
//Create column vector to place in categories
x = { 0,
      0.1,
      0.2,
      0.3,
      0.4,
      0.5,
      0.6,
      0.7 };

//Cut points for data in 'x'
cut_pts = { 0.2,
            0.5 };

//Class 0:      x <= 0.2
//Class 1: 0.2 < x <= 0.5
//Class 2: 0.5 < x
r_open = reclassifyCuts(x, cut_pts);

//Class 0:      x < 0.2
//Class 1: 0.2 < x < 0.5
//Class 2: 0.5 < x
r_closed = reclassifyCuts(x, cut_pts, 1);

print "x = " x;
print;
print "r_open = " r_open;
print;
print "r_closed = " r_closed;
print;
print "cut_pts = " cut_pts;
```

After the code above:

reclassifyCuts

```
x =  
0.00  
0.10  
0.20  
0.30  
0.40  
0.50  
0.60  
0.70  
  
r_open =  
0.00  
0.00  
0.00  
1.0  
1.0  
1.0  
2.0  
2.0  
  
r_closed =  
0.00  
0.00  
1.0  
1.0  
1.0  
2.0  
2.0  
2.0  
  
cut_pts =  
0.20  
0.50
```

Example 2: Classifying blood pressure data

```
//Create a column of blood pressure data
bp = { 87,
      154,
      127,
      112,
      159,
      90,
      151,
      109,
      125,
      107 };

//Assign cut points
cut_pts = { 120, 140 };

//Create categorical variable
bp_category = reclassifyCuts(bp, cut_pts);

print "bp = " bp;
print;
print "bp_category = " bp_category;
print;
print "cut_pts = " cut_pts;
```

After the code above:

```
bp =
87.00
154.0
127.0
112.0
159.0
90.00
151.0
```

reclassifyCuts

```
109.0
125.0
107.0

bp_category =
0.0000
2.000
1.000
0.0000
2.000
0.0000
2.000
0.0000
1.000
0.0000

cut_pts =
120.0
140.0
```

We can take the categorical data output from **reclassifyCuts** and use the **reclassify** function to change the numeric categories to string categories like this:

```
//Starting categories
from = { 0, 1, 2 };

//New categories
to = "normal" $| "prehypertension" $| "hypertension";

bp_category = reclassify(bp_category, from, to);
print "bp_category = " bp_category;
```

After the code above:

```
bp_category =  
normal  
hypertension  
prehypertension  
normal  
hypertension  
normal  
hypertension  
normal  
prehypertension  
normal
```

Source

datatran.src

See Also

[code](#), [recode](#), [reclassify](#), [substute](#), [rescale](#)

recode

Purpose

Changes the values of an existing vector from a vector of new values. Used in data transformations.

Format

```
y = recode(x, e, v);
```

Input

x Nx1 vector to be recoded (changed).

recode

e $N \times K$ matrix of 1's and 0's.
 v $K \times 1$ vector containing the new values to be assigned to the recoded variable.

Output

y $N \times 1$ vector containing the recoded values of x .

Remarks

There should be no more than a single 1 in any row of e .

For any given row N of x and e , if the K th column of e is 1, the K th element of v will replace the original element of x .

If every column of e contains a 0, the original value of x will be unchanged.

Example

```
x = { 20,  
      45,  
      32,  
      63,  
      29 };  
  
//Create 4 column vectors with a 1 where the statement  
//evaluates as 'true'  
e1 = (20 .lt x) .and (x .le 30);  
e2 = (30 .lt x) .and (x .le 40);  
e3 = (40 .lt x) .and (x .le 50);  
e4 = (50 .lt x) .and (x .le 60);  
  
//Horizontally concatenate the column vectors into a 5x4  
//matrix
```

```
e = e1~e2~e3~e4;

v = { 1.2,
      2.4,
      3.1,
      4.6 };

//Replace elements of 'x' with elements from 'v' based upon
//the 0's and 1's in 'e'
y = recode(x,e,v);
```

The above code assigns *e* and *y* as follows:

```

      0   0   0   0
      0   0   1   0
e = 0   1   0   0
      0   0   0   0
      1   0   0   0

//Since the third column of the second row of 'e' is equal
//to 1, the second row of 'y' is set equal to the third
//element of 'v', etc.
      20.000000
      3.1000000
y = 2.4000000
      63.000000
      1.2000000
```

Source

datatran.src

See Also

[code](#), [reclassifyCuts](#), [reclassify](#), [substute](#), [rescale](#), [dummy](#)

recode (dataloop)

recode (dataloop)

Purpose

Changes the value of a variable with different values based on a set of logical expressions.

Format

```
recode var with  
  or  
recode #var with  
  or  
recode $var with  
  val1 for expression_1,  
  val2 for expression_2,  
  .  
  .  
  .  
  valn for expression_n;
```

Input

<i>var</i>	literal, the new variable name.
<i>val</i>	scalar, value to be used if corresponding expression is TRUE.
<i>expression</i>	logical scalar-returning expression that returns nonzero TRUE or zero FALSE.

Remarks

If '\$' is specified, the variable will be considered a character variable. If '#' is

specified, the variable will be considered numeric. If neither is specified, the type of the variable will be left unchanged.

The logical expressions must be mutually exclusive, that is only one may return TRUE for a given row (observation).

If none of the expressions is TRUE for a given row (observation), its value will remain unchanged.

Any variables referenced must already exist, either as elements of the source data set, as [extern](#)'s, or as the result of a previous [make](#), [vector](#), or [code](#) statement.

Example

```
recode age with
  1 for age < 21,
  2 for age >= 21 and age < 35,
  3 for age >= 35 and age < 50,
  4 for age >= 50 and age < 65,
  5 for age >= 65;
```

```
recode $ sex with
  "MALE" for sex =\,= 1,
  "FEMALE" for sex =\,= 0;
```

```
recode # sex with
  1 for sex $=\,= "MALE",
  0 for sex $=\,= "FEMALE";
```

See Also

[code \(dataloop\)](#)

L

recserar

recserar

Purpose

Computes a vector of autoregressive recursive series.

Format

$y = \text{recserar}(x, y0, a);$

Input

x	$N \times K$ matrix
$y0$	$P \times K$ matrix.
a	$P \times K$ matrix.

Output

y	$N \times K$ matrix containing the series.
-----	--

Remarks

recserar is particularly useful in dealing with time series.

Typically, the result would be thought of as K vectors of length N .

$y0$ contains the first P values of each of these vectors (thus, these are prespecified). The remaining elements are constructed by computing a P th order "autoregressive" recursion, with weights given by a , and then by adding the result to the corresponding elements of x . That is, the t th row of y is given by:

$$y[t, :] = x[t, :] + a[1, :] * y[t-1, :] + \dots + a[P, :] * y[t-p, :], \quad t = P + 1, \dots, N$$

and

$$y[t, :] = y0[t, :], \quad t = 1, \dots, P$$

Note that the first P rows of x are not used.

Example

```
n = 10;
fn multnorm(n, sigma) = rndn(n, rows(sigma))*chol(sigma);
let sig[2,2] = { 1 -.3, -.3 1 };
rho = 0.5~0.3;
y0 = 0~0;
e = multnorm(n, sig);
x = ones(n,1)~rndn(n,3);
b = 1|2|3|4;
y = recserar(x*b+e, y0, rho);
```

In this example, two autoregressive series are formed using simulated data. The general form of the series can be written:

$$\begin{aligned} y[1, t] &= \rho[1, 1] * y[1, t-1] + x[t, :] * b + e[1, t] \\ y[2, t] &= \rho[2, 1] * y[2, t-1] + x[t, :] * b + e[2, t] \end{aligned}$$

The error terms ($e[1, t]$ and $e[2, t]$) are not individually serially correlated, but they are contemporaneously correlated with each other. The variance-covariance matrix is sig .

See Also

[recsercp](#), [recserre](#)

L

recsercp

recsercp

Purpose

Computes a recursive series involving products. Can be used to compute cumulative products, to evaluate polynomials using Horner's rule, and to convert from base *b* representations of numbers to decimal representations among other things.

Format

$y = \text{recsercp}(x, z);$

Input

<i>x</i>	NxK or 1xK matrix
<i>z</i>	NxK or 1xK matrix.

Output

<i>y</i>	NxK matrix in which each column is a series generated by a recursion of the form: <div>$y(1) = x(1) + z(1)$$y(t) = y(t-1) * x(t) + z(t), \quad t=2, \dots, N$</div>
----------	---

Remarks

The following **GAUSS** code could be used to emulate **recsercp** when the number of rows in *x* and *z* is the same:

```

/* assume here that rows(z) is also n */
n = rows(x);
y = zeros(n, 1);
y[1,.] = x[1,.] + z[1,.];

i = 2;
do until i > n;
    y[i,.] = y[i-1,.] .* x[i,.] + z[i,.];
    i = i + 1;
endo;

```

Note that K series can be computed simultaneously, since x and z can have K columns (they must both have the same number of columns).

recsercp allows either x or z to have only 1 row.

recsercp(x , 0) will produce the cumulative products of the elements in x .

Example

```

c1 = c[1,.];
n = rows(c) - 1;
y = recsercp(x, trimr(c ./ c1,1,0));
p = c1 .* y[n,.];

```

If x is a scalar and c is an $(N+1) \times 1$ vector, the result p will contain the value of the polynomial whose coefficients are given in c . That is:

$$p = c[1,.] \cdot x^n + c[2,.] \cdot x^{(n-1)} + \dots + c[n+1,.]$$

Note that both x and c could contain more than 1 column, and then this code would evaluate the entire set of polynomials at the same time. Note also that if $x = 2$, and if c contains the digits of the binary representation of a number, then p will be the decimal representation of that number.

recserrc

See Also

[recserar](#), [recserrc](#)

recserrc

Purpose

Computes a recursive series involving division.

Format

$y = \text{recserrc}(x, z);$

Input

x	1xK or Kx1 vector.
z	NxK matrix.

Output

y NxK matrix in which each column is a series generated by a recursion of the form:

$$\begin{aligned} y[1] &= x \bmod z[1], & x &= \text{trunc}(x/z[1]) \\ y[2] &= x \bmod z[2], & x &= \text{trunc}(x/z[2]) \\ y[3] &= x \bmod z[3], & x &= \text{trunc}(x/z[3]) \\ &\vdots \\ &\vdots \end{aligned}$$

$$y[n] = x \bmod z[n]$$

Remarks

Can be used to convert from decimal to other number systems (radix conversion).

Example

```
x = 2|8|10;
b = 2;
n = maxc(log(x) ./ log(b)) + 1;
z = reshape(b, n, rows(x));
y = rev(recserrc(x, z))';
```

The result, y , will contain in its rows (note that it is transposed in the last step) the digits representing the decimal numbers 2, 8, and 10 in base 2:

```
0 0 1 0
1 0 0 0
1 0 1 0
```

Source

recserrc.src

See Also

[recserar](#), [recsercp](#)

renamefile

renamefile

Purpose

Changes file name.

Format

```
ret = renamefile("oldname","newname");
```

Input

<i>oldname</i>	string, existing file name.
<i>newname</i>	string, new file name.

Output

<i>ret</i>	scalar, 0 if successful.
------------	--------------------------

Example

```
ret = renamefile("myfile.gss","mynewfile.gss");
```

In this example, a file in the current working directory with the name "myfile.gss" will be renamed "mynewfile.gss" in the same directory. Full path information may also be included:

```
//On Windows
ret = renamefile("c:\\gauss17\\myfile.gss",
    "c:\\gauss17\\mynewfile.gss");

//On Linux/Mac
```



```
ret = renamefile("/home/user/gauss17/myfile.gss",  
    "/home/user/gauss17/mynewfile.gss");
```

rerun

Purpose

Displays the most recently created graphics file.

Library

pgraph

Format

```
rerun;
```

Remarks

`rerun` is used by the **endwind** function.

Source

pcart.src

Globals

_pcmdlin, _pnotify, _psilent, _ptek, _pzoom

rescale

Purpose

Scales the columns of a matrix

rescale

Format

```
{ x_s, location, scale_factor } = rescale(x, method);
x_s = rescale(x, location, scale_factor);
```

Input

<i>x</i>	NxK matrix or NxK array to be rescaled		
<i>location</i>	1xK vector, used for column centering		
<i>scale_factor</i>	1xK vector, used for column scaling		
<i>method</i>	String, name of scaling and centering method:		
	<i>Method</i>	<i>Location</i>	<i>scale_factor</i>
	"euclidean"	0	Euclidean length: $\sqrt{\sum_{i=1}^n x_i^2}$
	"mad"	Median	Absolute deviation from the median
	"maxabs"	0	Maximum absolute value
	"midrange"	$(Max + Min)/2$	$Range/2$
	"range"	Minimum	Range
	"standardize"	Mean	Standard deviation

"sum"	0	Sum
"ustd"	0	Standard deviation about the origin

Output

<i>x_s</i>	Matrix or multi-dimensional array, containing the scaled columns of <i>x</i>
<i>location</i>	1xK vector, containing the values used to center the columns of the input matrix <i>x</i>
<i>scale_factor</i>	1xK vector, containing the values used to scale the columns of the input matrix <i>x</i>

Examples

Example 1: Specifying a scaling method

```
//Create a column vector
x = { 12.5,
      18.2,
      10.8,
      8.3,
      15.4,
      21.5,
      14.6,
      16.7 };

//Standardize 'x' and return the location and scaling
factors
{ x_s, location, scale_factor} = rescale(x, "standardize");

print "x_s = " x_s;
```

rescale

```
print "location = " location;
print "scale_factor = " scale_factor;
```

After the code above:

```
x_s =
    -0.53463295
     0.81977052
    -0.93857785
    -1.5326145
     0.15444952
     1.6038989
    -0.035642197
     0.46334856

location =      14.750000
scale_factor =    4.2084948
```

Example 2: Specifying a scaling method for multiple columns

```
//Create a matrix with 2 columns
x = {   12.5 1088.5,
       18.2  879.3,
       10.8 1232.0,
        8.3 1189.8,
       15.4  932.1,
       21.5 1009.2,
       14.6  656.7,
       16.7 1251.5 };

//Standardize 'x' and return the location and scaling
factors
{ x_s, location, scale_factor } = rescale(x,
```

```

"standardize");

print"x_s = " x_s;
print"location = " location;
print"scale_factor = " scale_factor;

```

After the code above:

```

x_s =
-0.53463295      0.28751716
 0.81977052     -0.73869039
-0.93857785      0.99144060
-1.5326145       0.78443315
 0.15444952     -0.47968581
 1.6038989      -0.10148025
-0.035642197    -1.8306302
 0.46334856      1.0870957

location =      14.750000      1029.8875
scale_factor =   4.2084948     203.85740

```

Example 3: Applying previously created location and scaling factors

Continuing with the variables used in example 2, we can apply the returned location and scaling factors to standardize additional observations of our two variables.

```

//Additional observations
x_new = {
    9.3  964.1,
    10.9 1173.7,
    11.1 1232.0,
    9.1  1051.2,
    14.6 1124.1,
    18.4  815.3,

```

rescale

```
20.2 1292.6,  
18.5 833.1 };  
  
//Standardize 'x' using the location and scaling factors  
print"location = " location;  
print"scale_factor = " scale_factor;  
  
//returned in example 2  
x_new_s = rescale(x_new, location, scale_factor);  
  
print"x_new_s = " x_new_s;  
print"x_new = " x_new;
```

After the code above:

```
location =          14.750000          1029.8875  
scale_factor =      4.2084948          203.85740  
  
x_new_s =  
    -1.2949998      -0.32271333  
    -0.91481638      0.70545637  
    -0.86729345      0.99144060  
    -1.3425227      0.10454612  
    -0.035642197     0.46214904  
     0.86729345     -1.0526353  
     1.2949998      1.2887072  
     0.89105492     -0.96531940  
  
x_new =  
     9.3000000      964.10000  
    10.900000      1173.7000  
    11.100000      1232.0000  
     9.100000      1051.2000  
    14.600000      1124.1000  
    18.400000      815.30000
```

20.200000	1292.6000
18.500000	833.10000

See Also

[code](#), [recode](#), [reclassifyCuts](#), [reclassify](#), [rescale](#), [substute](#)

reshape

Purpose

Reshapes a matrix.

Format

```
y = reshape(x, r, c);
```

Input

x	NxK matrix.
r	scalar, new row dimension.
c	scalar, new column dimension.

Output

y	$r \times c$ matrix created from the elements of x .
-----	--

Remarks

Matrices are stored in row major order.

reshape

The first c elements are put into the first row of y , the second in the second row, and so on. If there are more elements in x than in y , the remaining elements are discarded. If there are not enough elements in x to fill y , then when **reshape** runs out of elements, it goes back to the first element of x and starts getting additional elements from there.

Example

```
y = reshape(x, 2, 6);
```

```
      1  2  3  4
if x = 5  6  7  8 then y = 1  2  3  4  5  6
      9 10 11 12           7  8  9 10 11 12
```

```
      1  2  3
if x = 4  5  6 then y = 1  2  3  4  5  6
      7  8  9           7  8  9  1  2  3
```

```
      1  2  3  4  5
if x = 6  7  8  9 10 then y = 1  2  3  4  5  6
     11 12 13 14 15           7  8  9 10 11 12
```

```
if x = 1  2 then y = 1 2 3 4 1 2
      3  4           3 4 1 2 3 4
```

```
if x = 1 then y = 1 1 1 1 1 1
                1 1 1 1 1 1
```

See Also

[submat](#), [vec](#)

retp

Purpose

Returns from a procedure or keyword.

Format

```
retp;  
retp(x, y,...);
```

Remarks

For more details, see **PROCEDURES AND KEYWORDS**, CHAPTER 1.

In a `retp` statement 0-1023 items may be returned. The items may be expressions. Items are separated by commas.

It is legal to return with no arguments, as long as the procedure is defined to return 0 arguments.

See Also

[proc](#), [keyword](#), [endp](#)

return

Purpose

Returns from a subroutine.

Format

```
return;  
return(x, y,...);
```

rev

Remarks

The number of items that may be returned from a subroutine in a `return` statement is limited only by stack space. The items may be expressions. Items are separated by commas.

It is legal to return with no arguments and therefore return nothing.

See Also

[gosub](#), [pop](#)

rev

Purpose

Reverses the order of the rows in a matrix.

Format

```
 $y = \mathbf{rev}(x);$ 
```

Input

x	NxK matrix.
-----	-------------

Output

y	NxK matrix containing the reversed rows of x .
-----	--

Remarks

The first row of y will be where the last row of x was and the last row will be where

the first was and so on. This can be used to put a sorted matrix in descending order.

Example

```
//Set the rng seed for repeatable results
rndseed 345345;

//Set print formatting to print 4 spaces for each column
//and 0 numbers after the decimal
format /rd 4,0

//Create some random integers
x = round(rndn(5,3)*10);

//Reverse the order of the columns
y = rev(x);

print "x = " x;
print "y = " y;
```

The code above produces the following output:

```
x =
  10  -14   -7
   3   -1   -5
  -7    4    2
   1    1    1
   7   -7    2
y =
   7   -7    2
   1    1    1
  -7    4    2
   3   -1   -5
  10  -14   -7
```

rfft

See Also

[sortc](#)

rfft

Purpose

Computes a real 1- or 2-D Fast Fourier transform.

Format

```
 $y = \mathbf{rfft}(x);$ 
```

Input

x	NxK real matrix.
-----	------------------

Output

y	LxM matrix, where L and M are the smallest powers of 2 greater than or equal to N and K, respectively.
-----	--

Remarks

Computes the RFFT of x , scaled by $1/(L*M)$.

This uses a Temperton Fast Fourier algorithm.

If N or K is not a power of 2, x will be padded out with zeros before computing the transform.

See Also

[rffti](#), [fft](#), [ffti](#), [fftm](#), [fftmf](#)

rffti

Purpose

Computes inverse real 1- or 2-D Fast Fourier transform.

Format

```
y = rffti(x);
```

Input

x	NxK matrix.
---	-------------

Output

y	LxM real matrix, where L and M are the smallest prime factor products greater than or equal to N and K.
---	---

Remarks

It is up to the user to guarantee that the input will return a real result. If in doubt, use `ffti`.

See Also

[rfft](#), [fft](#), [ffti](#), [fftm](#), [fftmf](#)

rfftip

Purpose

Computes an inverse real 1- or 2-D FFT. Takes a packed format FFT as input.

rfftip

Format

```
y = rfftip(x);
```

Input

x	$N \times K$ matrix or K -length vector.
-----	--

Output

y	$L \times M$ real matrix or M -length vector.
-----	---

Remarks

rfftip assumes that its input is of the same form as that output by **rfftp** and **rfftnp**.

rfftip uses the Temperton prime factor FFT algorithm. This algorithm can compute the inverse FFT of any vector or matrix whose dimensions can be expressed as the product of selected prime number factors. **GAUSS** implements the Temperton algorithm for any integer power of 2, 3, and 5, and one factor of 7. Thus, **rfftip** can handle any matrix whose dimensions can be expressed as:

$$2^p \times 3^q \times 5^r \times 7^s$$

$$p, q, r \geq 0$$

$$s = 0 \text{ or } 1$$

If a dimension of x does not meet this requirement, it will be padded with zeros to the next allowable size before the inverse FFT is computed. Note that **rfftip** assumes the length (for vectors) or column dimension (for matrices) of x is $K-1$ rather than K , since the last element or column does not hold FFT information, but the Nyquist frequencies.

The sizes of x and y are related as follows: L will be the smallest prime factor product greater than or equal to N , and M will be twice the smallest prime factor product greater than or equal to $K-1$. This takes into account the fact that x contains both positive and negative frequencies in the row dimension (matrices only), but only positive frequencies, and those only in the first $K-1$ elements or columns, in the length or column dimension.

It is up to the user to guarantee that the input will return a real result. If in doubt, use **ffti**. Note, however, that **ffti** expects a full FFT, including negative frequency information, for input.

Do not pass **rffftip** the output from **rffft** or **rfftn**-it will return incorrect results. Use **rffti** with those routines.

See Also

[fft](#), [ffti](#), [fftn](#), [fftni](#), [fftn](#), [rfft](#), [rffti](#), [rfftn](#), [rfftnp](#), [rfftp](#)

rfftn

Purpose

Computes a real 1- or 2-D FFT.

Format

$y = \text{rfftn}(x);$

Input

x $N \times K$ real matrix.

Output

y $L \times M$ matrix, where L and M are the smallest prime

rfftn

factor products greater than or equal to N and K, respectively.

Remarks

rfftn uses the Temperton prime factor FFT algorithm. This algorithm can compute the FFT of any vector or matrix whose dimensions can be expressed as the product of selected prime number factors. **GAUSS** implements the Temperton algorithm for any power of 2, 3, and 5, and one factor of 7. Thus, **rfftn** can handle any matrix whose dimensions can be expressed as:

```
2p x 3q x 5r x 7s

p, q, r ≥ 0      -- for rows of matrix
p > 0, q, r ≥ 0 -- for columns of matrix
p > 0, q, r ≥ 0 -- for length of a vector
s = 0 or 1       -- for all dimensions
```

If a dimension of *x* does not meet these requirements, it will be padded with zeros to the next allowable size before the FFT is computed.

rfftn pads matrices to the next allowable size; however, it generally runs faster for matrices whose dimensions are highly composite numbers, i.e., products of several factors (to various powers), rather than powers of a single factor. For example, even though it is bigger, a 33600x1 vector can compute as much as 20 percent faster than a 32768x1 vector, because 33600 is a highly composite number, $2^6 \times 3 \times 5^2 \times 7$, whereas 32768 is a simple power of 2, 2^{15} . For this reason, you may want to hand-pad matrices to optimum dimensions before passing them to **rfftn**. The **Run-Time Library** includes two routines, **optn** and **optnevn**, for determining optimum dimensions. Use **optn** to determine optimum rows for matrices, and **optnevn** to determine optimum columns for matrices and optimum lengths for vectors.

The **Run-Time Library** also includes the **nextn** and **nextnevn** routines, for determining allowable dimensions for matrices and vectors. (You can use these to see the dimensions to which **rfftn** would pad a matrix or vector.)

rfftn scales the computed FFT by $1/(L*M)$.

See Also

[fft](#), [ffti](#), [fftm](#), [fftimi](#), [fftn](#), [rfft](#), [rffti](#), [rfftip](#), [rfftnp](#), [rfftp](#)

rfftnp

Purpose

Computes a real 1- or 2-D FFT. Returns the results in a packed format.

Format

```
y = rfftnp(x);
```

Input

x	$N \times K$ real matrix or K -length real vector.
-----	--

Output

y	$L \times (M/2+1)$ matrix or $(M/2+1)$ -length vector, where L and M are the smallest prime factor products greater than or equal to N and K , respectively.
-----	--

Remarks

For 1-D FFT's, **rfftnp** returns the positive frequencies in ascending order in the first

rfftnp

M/2 elements, and the Nyquist frequency in the last element. For 2-D FFT's, **rfftnp** returns the positive and negative frequencies for the row dimension, and for the column dimension, it returns the positive frequencies in ascending order in the first M/2 columns, and the Nyquist frequencies in the last column. Usually the FFT of a real function is calculated to find the power density spectrum or to perform filtering on the waveform. In both these cases only the positive frequencies are required. (See also **rfft** and **rfftn** for routines that return the negative frequencies as well.)

rfftnp uses the Temperton prime factor FFT algorithm. This algorithm can compute the FFT of any vector or matrix whose dimensions can be expressed as the product of selected prime number factors. **GAUSS** implements the Temperton algorithm for any power of 2, 3, and 5, and one factor of 7. Thus, **rfftnp** can handle any matrix whose dimensions can be expressed as:

```
2p x 3q x 5r x 7s

p, q, r ≥ 0      -- for rows of matrix

p > 0, q, r ≥ 0 -- for columns of matrix

p > 0, q, r ≥ 0 -- for length of a vector

s = 0 or 1      -- for all dimensions
```

If a dimension of *x* does not meet these requirements, it will be padded with zeros to the next allowable size before the FFT is computed.

rfftnp pads matrices to the next allowable size; however, it generally runs faster for matrices whose dimensions are highly composite numbers, i.e., products of several factors (to various powers), rather than powers of a single factor. For example, even though it is bigger, a 33600x1 vector can compute as much as 20 percent faster than a 32768x1 vector, because 33600 is a highly composite number, 2⁶x3x5²x7, whereas 32768 is a simple power of 2, 2¹⁵. For this reason, you may want to hand-pad matrices to optimum dimensions before passing them to **rfftnp**. The **Run-Time Library** includes two routines, **optn** and **optnevn**, for determining optimum dimensions. Use

rfftnp scales the computed FFT by $1/(L*M)$.

[fft](#), [ffti](#), [fftm](#), [fftm_i](#), [fftn](#), [rfft](#), [rffti](#), [rfftip](#), [rfftn](#), [rfftp](#)

Computes a real 1- or 2-D FFT. Returns the results in a packed format.

```
y = rfftp(x);
```

\mathbf{x} NxK real matrix or K-length real vector.

y $L \times (M/2+1)$ matrix or $(M/2+1)$ -length vector, where L and M are the smallest powers of 2 greater than or equal to N and K , respectively.

rndBernoulli

Remarks

If a dimension of x is not a power of 2, it will be padded with zeros to the next allowable size before the FFT is computed.

For 1-D FFT's, **rffftp** returns the positive frequencies in ascending order in the first $M/2$ elements, and the Nyquist frequency in the last element. For 2-D FFT's, **rffftp** returns the positive and negative frequencies for the row dimension, and for the column dimension, it returns the positive frequencies in ascending order in the first $M/2$ columns, and the Nyquist frequencies in the last column. Usually the FFT of a real function is calculated to find the power density spectrum or to perform filtering on the waveform. In both these cases only the positive frequencies are required. (See also **rfft** and **rfftn** for routines that return the negative frequencies as well.)

rffftp scales the computed FFT by $1/(L*M)$.

rffftp uses the Temperton FFT algorithm.

See Also

[fft](#), [ffti](#), [fftm](#), [fftn](#), [fftni](#), [fftnm](#), [fftnr](#), [fftnr](#), [fftnr](#), [fftnr](#), [fftnr](#)

rndBernoulli

Purpose

Computes Bernoulli distributed random numbers.

Format

```
{ r, newstate } = rndBernoulli(r, c, prob, state);  
r = rndBernoulli(r, c, prob);
```

Input

<i>r</i>	Scalar, number of rows of the output matrix.
<i>c</i>	Scalar, number of columns of the output matrix.
<i>prob</i>	Scalar, probability parameter.
<i>state</i>	Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number functions.

Output

<i>r</i>	<i>r</i> x <i>c</i> matrix, Bernoulli random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$E(X) = \text{prob}$$

$$\text{Var}(X) = \text{prob} * (1 - \text{prob})$$

Example

```
//Bernoulli random numbers can be used to model qualitative
//binary data (i.e., yes/no, true/false), such as marital
```

rndBeta

```
//status.  
  
//Set the random seed for repeatable numbers.  
  
rndseed 723940439;  
  
//The percentage of married people in the population we  
//would like to model.  
prob = 0.7;  
  
//Create 10,000 Bernoulli random numbers  
r = rndBernoulli(10000, 1, prob);  
  
//The mean of 'r' should approximately equal 'prob'  
mu = mean(r);  
print mu;
```

```
0.70270000
```

See Also

[rndMVn](#), [rndCreateState](#)

rndBeta

Purpose

Computes beta pseudo-random numbers with a choice of underlying random number generator.

Format

```
{ x, newstate } = rndBeta(r, c, a, b, state);  
x = rndBeta(r, c, a, b);
```

Input

<i>r</i>	Scalar, number of rows of resulting matrix.
<i>c</i>	Scalar, number of columns of resulting matrix.
<i>a</i>	<i>r</i> x <i>c</i> matrix, or <i>r</i> x 1 vector, or 1 x <i>c</i> vector, or scalar, first shape argument for beta distribution.
<i>b</i>	<i>r</i> x <i>c</i> matrix, or <i>r</i> x 1 vector, or 1 x <i>c</i> vector, or scalar, second shape argument for beta distribution.
<i>state</i>	Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, beta distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$E(x) = a/(a+b)$$

$$Var(x) = a*b/((a+b+1) * (a+b)^2)$$

rndBeta

$$\begin{aligned}0 < x < 1 \\ a > 0 \\ b > 0\end{aligned}$$

r and c will be truncated to integers if necessary.

Example

Example 1

This example illustrates basic usage of **rndBeta**, leaving the management of the random number state to GAUSS to handle internally.

```
num_rows = 100;
num_cols = 5;
a = 3;
b = 2;
x = rndBeta(num_rows, num_cols, a, b);
```

Example 2

```
//Starting seed for random number generator
seed = 235235;

//If a 'seed' or 'state' vector is passed in,
//then a state vector will be returned
{ x, newstate } = rndBeta(100, 5, 3, 2, seed);
```

Technical Notes

The default generator for **rndBeta** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndCauchy

Purpose

Computes Cauchy random numbers with a choice of underlying random number generator.

Format

```
{ r, newstate } = rndCauchy(rows, cols, location, scale,  
state);  
r = rndCauchy(rows, cols, location, scale);
```

Input

<i>rows</i>	Scalar, number of rows of resulting matrix.
<i>cols</i>	Scalar, number of columns of resulting matrix.
<i>location</i>	Scalar or ExE conformable matrix with <i>rows</i> and <i>cols</i> .
<i>scale</i>	Scalar or ExE conformable matrix with <i>rows</i> and <i>cols</i> .
<i>state</i>	Optional argument - scalar or opaque vector.
Scalar case:	
<i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock.	
Opaque vector case:	
<i>state</i> = the state vector returned from a previous call to one of the standard random number functions.	

rndChiSquare

Output

<i>r</i>	<i>rows</i> x <i>cols</i> matrix, Cauchy distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

```
E(x) = undefined  
Var(x) = undefined  
Median(x) = location
```

r and *c* will be truncated to integers if necessary.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndChiSquare

Purpose

Creates pseudo-random numbers with a chi-squared distribution, with an optional non-centrality parameter and a choice of underlying random number generator.

Format

```
{ x, newstate } = rndChiSquare(r, c, df, s_ncp, state);  
x = rndChiSquare(r, c, df, s_ncp);  
x = rndChiSquare(r, c, df);
```

Input

<i>r</i>	Scalar, number of rows of resulting matrix.
<i>c</i>	Scalar, number of columns of resulting matrix.
<i>df</i>	Scalar, degrees of freedom.
<i>s_ncp</i>	Optional argument - scalar, non-centrality parameter. NOTE: This is the <u>square root</u> of the noncentrality parameter that sometimes goes under the symbol lambda.
<i>state</i>	Optional argument - scalar or opaque vector. Scalar case: <i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock. Opaque vector case: <i>state</i> = the state vector returned from a previous call to one of the rnd random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, chi-square distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$E(x) = k + \lambda$$

$$\sigma^2(x) = 2*k + 4*\lambda$$

`rndcon`, `rndmult`, `rndseed`

where:

```
k = df
λ = s_ncp^2
```

Technical Notes

The default generator for **`rndChiSquare`** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **`rndCreateState`**.

See Also

[`rndCreateState`](#), [`rndStateSkip`](#)

`rndcon`, `rndmult`, `rndseed`

Purpose

Resets the parameters of the linear congruential random number generator that is the basis for **`rndu`**, **`rndi`** and **`rndn`**.

Format

```
rndcon c;  
rndmult a;  
rndseed seed;
```

Input

<i>c</i>	scalar, constant for the random number generator.
<i>a</i>	scalar, multiplier for the random number generator.
<i>seed</i>	scalar, initial seed for the random number generator.

Parameter default values and ranges:

<code>seed</code>	<code>time(0)</code>	$0 < seed < 2^{32}$	<code>a</code>	1664525	0
$< a < 2^{32}$	<code>c</code>	1013904223	$0 < a < 2^{32}$		

Remarks

A linear congruential uniform random number generator is used by **rndu**, and is also called by **rndn**. These statements allow the parameters of this generator to be changed.

The procedure used to generate the uniform random numbers is as follows. First, the current "seed" is used to generate a new seed:

$$new_seed = ((a * seed) \% 2^{32}) + c) \% 2^{32}$$

(where $\%$ is the mod operator). Then a number between 0 and 1 is created by dividing the new seed by 2^{32} :

$$x = new_seed / 2^{32}$$

rndcon resets `c`.

rndmult resets `a`.

rndseed resets `seed`. This is the initial seed for the generator. The default is that **GAUSS** uses the clock to generate an initial seed when **GAUSS** is invoked.

GAUSS goes to the clock to seed the generator only when it is first started up. Therefore, if **GAUSS** is allowed to run for a long time, and if large numbers of random numbers are generated, there is a possibility of recycling (that is, the sequence of "random numbers" will repeat itself). However, the generator used has an extremely long cycle, so that should not usually be a problem.

The parameters set by these commands remain in effect until new commands are encountered, or until **GAUSS** is restarted.

L

rndCreateState

See Also

[rndu](#), [rndn](#), [rndi](#), [rndLCi](#), [rndKMi](#)

rndCreateState

Purpose

Creates a new random number stream for a specified generator type from a seed value.

Format

```
state = rndCreateState(brng, seed);
```

Input

<i>brng</i>	String, generator name. Options include: "mrg32k3a" L'Ecuyer's MRG32K3A "mt19937" Mersenne-Twister 19937 "sfmt19937" optimized Mersenne-Twister 19937 "mt2203-01" Mersenne-Twister 2203 "niederreiter" Niederreiter quasi-random numbers "sobol" Sobol quasi-random numbers "wh-01" Wichmann-Hill
<i>seed</i>	Scalar, starting seed value. if -1, GAUSS computes the starting seed based on the system clock. NOTE: For the quasi-random number generators, "sobol" and "niederreiter", this second input is the dimension rather than a starting seed. For "sobol", 1 ≤ dimension ≤ 40. For "niederreiter", 1 ≤ dimension ≤ 318. See examples below.

Output

state Opaque vector, the newly created state.

Examples

Example 1: Basic usage

```
//Starting seed value
seed = 123456;

//Create state for generator 'mrg32k3a'
state = rndCreateState("mrg32k3a", seed);

//Create a 5x1 vector of random normal numbers with
//the state created above
{ r, newstate } = rndn(5, 1, state);
```

After the code above, *r* will equal:

```
0.51489262
0.14053340
r = 1.2128406
0.17112172
-0.18788202
```

Example 2: creating a state from a numbered stream

Most random number generators have one single stream in which you can think of the starting state as a bookmark. The "mt2203" and "wh" (or Wichmann-Hill) each have multiple separate streams. The example below shows how to use these random number streams.

```
seed = 123456;
```

rndCreateState

```
//Create a state from the 1028th substream of the
//Mersenne-Twister 2203 RNG
state_mt = rndCreateState("mt2203-1028", seed);

//Create a state from the 112th substream of the
//Wichmann-Hill RNG
state_wh = rndCreateState("wh-112", seed);

//Generate numbers using the states
{ r1, state_mt } = rndu(4, 1, state_mt);
{ r2, state_wh } = rndu(4, 1, state_wh);
```

After the code above, *r1* and *r2* should equal:

r1 = 0.14291687	r2 = 0.0073824407
0.99670199	0.93756896
0.59512065	0.071140446
1.5776604e-06	0.021328991

Example 3: initializing the Sobol quasi-random number generator

```
//Initialize random seed with a dimension of 2
state = rndCreateState("sobol", 2);

//Create some random numbers using this state
{ r, state } = rndu(10, 2, state);
```

After the code above, *r*, should be equal to:

r = 0.5000	0.5000
0.7500	0.2500
0.2500	0.7500
0.3750	0.3750
0.8750	0.8750
0.6250	0.1250

0.1250	0.6250
0.1875	0.3125
0.6875	0.8125
0.9375	0.0625

Remarks

The states returned from this function may NOT be used with **rndMTu** or any of the **rndKM** or **rndLC** functions.

See Also

[rndStateSkip](#), [rndn](#), [rndu](#), [rndBeta](#)

rndExp

Purpose

Computes exponentially distributed random numbers with a choice of underlying random number generator.

Format

```
{ r, newstate } = rndExp(rows, cols, scale, state);  
r = rndExp(rows, cols, scale);
```

Input

<i>rows</i>	Scalar, number of rows of resulting matrix.
<i>cols</i>	Scalar, number of columns of resulting matrix.
<i>scale</i>	Scalar or a matrix that is ExE conformable with the dimensions of the output. The scale parameter sometimes called β

L

`rndExp`

state

Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number functions.

Output

r

rows x *cols* matrix, exponentially distributed random numbers.

newstate

Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are specified in terms of the scale parameter sometimes called β . This is the reciprocal of the rate parameter which is sometimes called λ :

$$\begin{aligned} E(x) &= \text{scale} = \beta = 1/\text{rate} = 1/\lambda \\ \text{Var}(x) &= \text{scale}^2 = \beta^2 = 1/\text{rate}^2 = 1/\lambda^2 \end{aligned}$$

See Also

[rndCreateState](#), [rndStateSkip](#)

rndgam

Purpose

Computes pseudo-random numbers with gamma distribution. NOTE: **rndgam** is deprecated and should be replaced with **rndGamma**.

Format

```
x = rndgam(r, c, alpha);
```

Input

<i>r</i>	scalar, number of rows of resulting matrix.
<i>c</i>	scalar, number of columns of resulting matrix.
<i>alpha</i>	MxN matrix, ExE conformable with <i>r</i> x <i>c</i> resulting matrix, shape parameters for gamma distribution.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, gamma distributed pseudo-random numbers.
----------	--

Remarks

The properties of the pseudo-random numbers in *x* are:

$$E(x) = \alpha \quad Var(x) = \alpha \quad \alpha > 0 \quad \alpha > 0$$

Source

random.src

L

rndGamma

See Also

[rndGamma](#)

rndGamma

Purpose

Computes gamma pseudo-random numbers with a choice of underlying random number generator.

Format

```
{ x, newstate } = rndGamma(r, c, shape, scale, state);
x = rndGamma(r, c, shape, scale);
```

Input

<i>r</i>	Scalar, number of rows of resulting matrix.
<i>c</i>	Scalar, number of columns of resulting matrix.
<i>shape</i>	<i>r</i> x <i>c</i> matrix, or <i>r</i> x 1 vector, or 1 x <i>c</i> vector, or scalar, shape argument for gamma distribution.
<i>scale</i>	<i>r</i> x <i>c</i> matrix, or <i>r</i> x 1 vector, or 1 x <i>c</i> vector, or scalar, scale argument for gamma distribution.
<i>state</i>	Optional argument - scalar or opaque vector. Scalar case: <i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock. Opaque vector case: <i>state</i> = the state vector returned from a previous call to one of the rnd random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, gamma distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$\begin{aligned} E(x) &= \text{shape} * \text{scale} \\ \text{Var}(x) &= \text{shape} * \text{scale}^2 \\ x &> 0 \\ \text{shape} &> 0 \\ \text{scale} &> 0 \end{aligned}$$

Example

Example 1

```
num_rows = 5;
num_cols = 1;
shape = 3;
scale = 2;

x = rndGamma(num_rows, num_cols, shape, scale);
```

Example 2

The gamma distribution is sometimes described in terms of a shape parameter and an inverse scale parameter, called the rate parameter. The rate parameter is the reciprocal of the scale parameter. With this parameterization, the random numbers will have the following properties:

$$\begin{aligned} E(x) &= \text{shape} / \text{rate} \\ \text{Var}(x) &= \text{shape} / (\text{rate}^2) \end{aligned}$$

rndGeo

If you prefer to think about the gamma distribution in these terms, then pass in the reciprocal of the rate parameter as the fourth argument to **rndGamma**.

```
shape = 3;
rate = 2;

x = rndGamma(5, 1, shape, 1/rate);
```

Technical Notes

The default generator for **rndGamma** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndGeo

Purpose

Computes geometric pseudo-random numbers with a choice of underlying random number generator.

Format

```
{ y, newstate } = rndGeo(r, c, prob, state);
y = rndGeo(r, c, prob);
```

Input

<i>r</i>	Scalar, row dimension.
<i>c</i>	Scalar, column dimension.
<i>prob</i>	Scalar or matrix: ExE conformable with <i>r</i> and <i>c</i>

<i>state</i>	<p>columns.</p> <p>Optional argument - scalar or opaque vector.</p> <p>Scalar case:</p> <p><i>state</i> = starting seed value. If -1, GAUSS computes the starting seed based on the system clock.</p> <p>Opaque vector case:</p> <p><i>state</i> = the state vector returned from a previous call to one of the rnd random number generators.</p>
--------------	---

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of geometrically distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

<p>The properties of the pseudo-random numbers in <i>y</i> are:</p> <div>$E(y) = (1 - \text{prob}) / \text{prob};$$Var(y) = (1 - \text{prob}) / \text{prob}^2$</div> <p><i>r</i> and <i>c</i> will be truncated to integers if necessary.</p>

See Also

[rndCreateState](#), [rndStateSkip](#)

L

`rndGumbel`

Technical Notes

The default generator for **rndGeo** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

`rndGumbel`

Purpose

Computes Gumbel distributed random numbers with a choice of underlying random number generator.

Format

```
{ r, newstate } = rndGumbel(rows, cols, location, scale, state);  
r = rndGumbel(rows, cols, scale);
```

Input

<i>rows</i>	Scalar, number of rows of resulting matrix.
<i>cols</i>	Scalar, number of columns of resulting matrix.
<i>location</i>	Scalar or ExE conformable matrix with <i>rows</i> and <i>cols</i> .
<i>scale</i>	Scalar or ExE conformable matrix with <i>rows</i> and <i>cols</i> .
<i>state</i>	Optional argument - scalar or opaque vector. Scalar case: <i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number functions.

Output

<i>r</i>	<i>rows</i> x <i>cols</i> matrix, Gumbel distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

This function uses the definition of the Gumbel distribution corresponding to the minimum extreme. The properties of the pseudo-random numbers in *y* are:

$$E(y) = \text{location} - \gamma \cdot \text{scale} \approx \text{location} - 0.5772 \cdot \text{scale}$$
$$\gamma = \text{Euler-Mascheroni constant}$$
$$\text{Var}(y) = (\pi^2 \cdot \text{scale}^2) / 6$$

r and *c* will be truncated to integers if necessary.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndHyperGeo

Purpose

Computes the random numbers for the hypergeometric distribution

L

`rndHyperGeo`

Format

```
x = rndHyperGeo(r, c, m, k, n);  
{ x, new_state } = rndHyperGeo(r, c, m, k, n, state);
```

Input

<i>r</i>	Scalar, row dimension of the return matrix <i>x</i>
<i>c</i>	Scalar, column dimension of the return matrix <i>x</i>
<i>m</i>	The size of the population from which draws will be made. ExE conformable with the row and column dimensions of the return matrix, <i>r</i> and <i>c</i>
<i>k</i>	The number of items in the population which possess a specified trait. ExE conformable with row and column dimensions of the return matrix, <i>r</i> , and <i>c</i>
<i>n</i>	The number of items drawn from the population. ExE conformable with the dimensions of the return matrix, <i>r</i> and <i>c</i>
<i>state</i>	Optional argument - scalar or opaque vector Scalar case: <i>state</i> = starting seed value. If -1, GAUSS computes the starting seed based on the system clock. Opaque vector case: <i>state</i> = the state vector returned from a previous call to one of the rndn random number generators

Output

<i>x</i>	The probability of drawing <i>x</i> items which possess a specified trait. NxK matrix, Nx1 vector or scalar
----------	---

new_state

Opaque vector, the updated state

Examples

Example 1: Basic Example

```
//Population size
m = 100;

//Number of marked items
k = 25;

//Number of items drawn
n = 40;

//Compute 1 random number
x = rndHyperGeo(1, 1, m, k, n);
```

Example 2

The example below shows how to create a random matrix in which each column has different parameters.

```
//Population size
m = 100;

//Number of marked items
k = 25;

//1x2 row vector, number of items drawn
n = { 40 50 };

//Compute a 10 x 2 matrix of random number
x = rndHyperGeo(10, 2, m, k, n);
```

randi

Both columns of the variable x created in the code above use the same values for m and k . However, the first column of x will be calculated using the first element of n , 40. The second column of x will be calculated using the second element of n , 50.

Example 3: Passing in a state vector

```
//Starting seed value
seed = 23424;

//Population size
m = 100;

//Number of marked items
k = 25;

//Number of items drawn
n = 40;

//Compute 1000x1 vector of random numbers
{ x, state } = randHyperGeo(1000, 1, m, k, n, seed);
```

See Also

[cdfHyperGeo](#), [pdfHyperGeo](#)

randi

Purpose

Returns a matrix of random integers from a user defined range.

Format

```
y = randi(r, c);
y = randi(r, c, range);
{ y, newstate } = randi(r, c, range, state);
```

Input

<i>r</i>	scalar, row dimension.
<i>c</i>	scalar, column dimension.
<i>range</i>	Optional argument. 2x1 matrix, the requested range of the random integers. The first element is the range minimum and the second element is the range maximum. If <i>range</i> is not supplied, the default range is $0 \leq y < 2^{32}$.
<i>state</i>	Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number generators.

Output

<i>y</i>	$r \times c$ matrix of random integers in the specified range.
<i>newstate</i>	Opaque vector, the updated state.

Examples

Example 1: Basic example

```
//Create a 10x5 vector of random
//integers between 0 and 2^32 - 1
r_int = randi(10, 5);
```

Example 2: Basic range

```
//Create a 10x1 vector of random
//integers between 1 and 100
range_start = 1;
range_end = 100;
idx = randi(10, 1, range_start | range_end);
```

Example 3: Using 'randi' to sample with replacement from a dataset

```
//Load data from the 'fueleconomy' dataset
//in the GAUSS examples directory
file_name = getGAUSSHome() $+ "examples/fueleconomy.dat";
fueleconomy = loadadd(file_name);

//Create a 100x1 vector of random
//integers between 1 and 100
range_start = 1;
range_end = rows(fueleconomy);
idx = randi(100, 1, range_start | range_end);

//Draw a 100 observation sample from 'fueleconomy'
fuel_sample = fueleconomy[idx, .];
```

Example 4: Using a state-vector

```
//Create a 1050x1 vector of random
//integers between 20 and 150
```

```
seed_start = 5423432;
range = { 20, 150 };
{ idx, state } = rndi(1050, 1, range, seed_start);
```

Remarks

r and c will be truncated to integers if necessary.

This generator is automatically seeded using the system clock when **GAUSS** first starts. However, that can be overridden using the [rndseed](#) statement, or passing in a seed or state as the last input to **rndi**.

See Also

[rndu](#), [rndn](#), [rndseed](#), [rndCreateState](#)

rndKMbeta

Purpose

Computes beta pseudo-random numbers.

Format

```
{ x, newstate } = rndKMbeta(r, c, a, b, state);
```

Input

r	scalar, number of rows of resulting matrix.
c	scalar, number of columns of resulting matrix.
a	$r \times c$ matrix, or $rx1$ vector, or $1 \times c$ vector, or scalar, first shape argument for beta distribution.
b	$r \times c$ matrix, or $rx1$ vector, or $1 \times c$ vector, or scalar, second shape argument for beta distribution.

L

rndKMbeta

<i>state</i>	scalar or 500x1 vector. Scalar case: <i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock. 500x1 vector case: <i>state</i> = the state vector returned from a previous call to one of the rndKM random number functions.
--------------	--

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, beta distributed random numbers.
<i>newstate</i>	500x1 vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$\begin{aligned} E(x) &= a/(a+b) \\ Var(x) &= a*b/((a+b+1) * (a+b^2)) \\ 0 < x < 1 &a > 0 b > 0 \end{aligned}$$

r and *c* will be truncated to integers if necessary.

Source

randkm.src

Technical Notes

rndKMbeta uses the recur-with-carry KISS+Monster algorithm described in the **rndKMi** Technical Notes.

rndKMgam

Purpose

Computes Gamma pseudo-random numbers.

Format

```
{ x, newstate } = rndKMgam(r, c, alpha, state);
```

Input

<i>r</i>	scalar, number of rows of resulting matrix.
<i>c</i>	scalar, number of columns of resulting matrix.
<i>alpha</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, shape argument for gamma distribution.
<i>state</i>	scalar or 500x1 vector.

Scalar case:

state = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

500x1 vector case:

state = the state vector returned from a previous call to one of the **rndKM** random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, gamma distributed random numbers.
<i>newstate</i>	500x1 vector, the updated state.

rndKMi

Remarks

The properties of the pseudo-random numbers in x are:

$$E(x) = \alpha \text{Var}(x) = \alpha x > 0 \alpha > 0$$

To generate **gamma**(α , θ) pseudo-random numbers where θ is a scale parameter, multiply the result of **rndKMgam** by θ .

Thus

$$z = \theta * \text{rndgam}(1, 1, \alpha);$$

has the properties

$$E(z) = \alpha * \theta \text{Var}(z) = \alpha * \theta^2 z > 0 \alpha > 0 \theta > 0$$

r and c will be truncated to integers if necessary.

Source

randkm.src

Technical Notes

rndKMgam uses the recur-with-carry KISS+Monster algorithm described in the **rndKMi** Technical Notes.

rndKMi

Purpose

Returns a matrix of random integers, $0 \leq y < 2^{32}$, and the state of the random number generator.

Format

```
{ y, newstate } = rndKMi(r, c, state);
```

Input

<i>r</i>	scalar, row dimension.
<i>c</i>	scalar, column dimension.
<i>state</i>	scalar or 500x1 vector.
Scalar case:	
<i>state</i> = starting seed value. If -1, GAUSS computes the starting seed based on the system clock.	
500x1 vector case:	
<i>state</i> = the state vector returned from a previous call to one of the rndKM random number generators.	

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of random integers between 0 and 2 ³² - 1, inclusive.
<i>newstate</i>	500x1 vector, the updated state.

Remarks

r and *c* will be truncated to integers if necessary.

Example

This example generates two thousand vectors of random integers, each with one million elements. The state of the random number generator after each iteration is used as

rndKMn

an input to the next generation of random numbers.

```
state = 13;
n = 2000;
k = 1000000;
c = 0;
min = 2^32+1;
max = -1;

do while c < n;
    { y,state } = rndKMi(k,1,state);
    min = minc(min | minc(y));
    max = maxc(max | maxc(y));
    c = c + k;
endo;

print "min " min;
print "max " max;
```

See Also

[rndKMn](#), [rndKMu](#)

Technical Notes

rndKMi generates random integers using a KISS+Monster algorithm developed by George Marsaglia. KISS initializes the sequence used in the recur-with-carry Monster random number generator. For more information on this generator see <http://www.Aptech.com/random>.

rndKMn

Purpose

Returns a matrix of standard normal (pseudo) random variables and the state of the random number generator.

Format

```
{ y, newstate } = rndKMn(r, c, state);
```

Input

<i>r</i>	scalar, row dimension.
<i>c</i>	scalar, column dimension.
<i>state</i>	scalar or 500x1 vector.

Scalar case:

state = starting seed value. If -1, **GAUSS** computes the starting seed based on the system clock.

500x1 vector case:

state = the state vector returned from a previous call to one of the **rndKM** random number generators.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of standard normal random numbers.
<i>newstate</i>	500x1 vector, the updated state.

Remarks

r and *c* will be truncated to integers if necessary.

Example

This example generates two thousand vectors of standard normal random numbers, each with one million elements. The state of the random number generator after each iteration is used as an input to the next generation of random numbers.

rndKMnb

```
state = 13;
n = 2000;
k = 1000000;
c = 0;
submean = {};

do while c < n;
    { y, state } = rndKMn(k, 1, state);
    submean = submean | meanc(y);
    c = c + k;
endo;

mean = meanc(submean);
print mean;
```

See Also

[rndKMu](#), [rndKMi](#)

Technical Notes

rndKMn calls the uniform random number generator that is the basis for **rndKMu** multiple times for each normal random number generated. This is the recur-with-carry KISS+Monster algorithm described in the **rndKMi** Technical Notes. Potential normal random numbers are filtered using the fast acceptance-rejection algorithm proposed by Kinderman, A.J. and J.G. Ramage, "Computer Generation of Normal Random Numbers," *Journal of the American Statistical Association*, December 1976, Volume 71, Number 356, pp. 893-896. It employs the error correction from Tirler et al. (2004), "An error in the Kinderman-Ramage method and how to fix it," *Computational and Data Analysis*, Vol. 47, 433-40.

rndKMnb

Purpose

Computes negative binomial pseudo-random numbers.

Format

```
{ x, newstate } = rndKMnb(r, c, k, p, state);
```

Input

<i>r</i>	scalar, number of rows of resulting matrix.
<i>c</i>	scalar, number of columns of resulting matrix.
<i>k</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, "event" argument for negative binomial distribution.
<i>p</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, "probability" argument for negative binomial distribution.
<i>state</i>	scalar or 500x1 vector. Scalar case: <i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock. 500x1 vector case: <i>state</i> = the state vector returned from a previous call to one of the rndKM random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, negative binomial distributed random numbers.
<i>newstate</i>	500x1 vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

L

rndKmp

$$E(x) = (k * p) / (1 - p)$$
$$Var(x) = (k * p) / (1 - p)^2 \quad x = 0, 1, \dots, k > 0, 0 < p < 1$$

r and *c* will be truncated to integers if necessary.

Source

randkm.src

Technical Notes

rndKMnb uses the recur-with-carry KISS+Monster algorithm described in the **rndKMi** Technical Notes.

rndKmp

Purpose

Computes Poisson pseudo-random numbers.

Format

{ *x*, *newstate* } = **rndKmp**(*r*, *c*, *lambda*, *state*);

Input

<i>r</i>	scalar, number of rows of resulting matrix.
<i>c</i>	scalar, number of columns of resulting matrix.
<i>lambda</i>	<i>r</i> x <i>c</i> matrix, or <i>r</i> x 1 vector, or 1 x <i>c</i> vector, or scalar, shape argument for Poisson distribution.
<i>state</i>	scalar or 500x1 vector.
Scalar case:	
<i>state</i> = starting seed value only. If -1, GAUSS computes	

the starting seed based on the system clock.

500x1 vector case:

state = the state vector returned from a previous call to one of the **rndKM** random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, Poisson distributed random numbers.
<i>newstate</i>	500x1 vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$E(x) = \lambda, \text{Var}(x) = \lambda, x = 0, 1, \dots, \lambda > 0$$

r and *c* will be truncated to integers if necessary.

Source

randkm.src

Technical Notes

rndKMp uses the recur-with-carry KISS+Monster algorithm described in the **rndKMi** Technical Notes.

rndKMu

Purpose

Returns a matrix of uniform (pseudo) random variables and the state of the random number generator.

L

`rndKMu`

Format

```
{ y, newstate } = rndKMu(r, c, state);
```

Input

<i>r</i>	scalar, row dimension.
<i>c</i>	scalar, column dimension.
<i>state</i>	scalar, 2x1 vector, or 500x1 vector.
Scalar case:	
<i>state</i> = starting seed value. If -1, GAUSS computes the starting seed based on the system clock.	
2x1 vector case:	
<i>[1]</i> the starting seed, uses the system clock if -1	
<i>[2]</i> 0 for $0 \leq y < 1$	
1 for $0 \leq y \leq 1$	
500x1 vector case:	
<i>state</i> = the state vector returned from a previous call to one of the rndKM random number generators.	

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of uniform random numbers, $0 \leq y < 1$.
<i>newstate</i>	500x1 vector, the updated state.

Remarks

r and *c* will be truncated to integers if necessary.

Example

This example generates two thousand vectors of uniform random numbers, each with one million elements. The state of the random number generator after each iteration is used as an input to the next generation of random numbers.

```
state = 13;
n = 2000;
k = 1000000;
c = 0;
submean = {};

do while c < n;
    { y, state } = rndKMu(k,1,state);
    submean = submean | meanc(y);
    c = c + k;
endo;

mean = meanc(submean);
print 0.5-mean;
```

See Also

[rndKMn](#), [rndKMi](#)

Technical Notes

rndKMu uses the recur-with-carry KISS-Monster algorithm described in the **rndKMi** Technical Notes. Random integer seeds from 0 to $2^{32}-1$ are generated. Each integer is divided by 2^{32} or $2^{32}-1$.

rndKMvm

Purpose

Computes von Mises pseudo-random numbers.

L

`rndKMvm`

Format

```
{ x, newstate } = rndKMvm(r, c, m, k, state);
```

Input

<i>r</i>	scalar, number of rows of resulting matrix.
<i>c</i>	scalar, number of columns of resulting matrix.
<i>m</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, means for vm distribution.
<i>k</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, shape argument for vm distribution.
<i>state</i>	scalar or 500x1 vector.
Scalar case:	
<i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock.	
500x1 vector case:	
<i>state</i> = the state vector returned from a previous call to one of the rndKM random number functions.	

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, von Mises distributed random numbers.
<i>newstate</i>	500x1 vector, the updated state.

Remarks

r and *c* will be truncated to integers if necessary.

Source

`randkm.src`

Technical Notes

rndKMvm uses the recur-with-carry KISS+Monster algorithm described in the **rndKMi** Technical Notes.

rndLaplace

Purpose

Computes Laplacian pseudo-random numbers with the choice of underlying random number generator.

Format

```
{ x, newstate } = rndLaplace(r, c, loc, scale, state);
x = rndLaplace(r, c, loc, scale);
```

Input

<i>r</i>	Scalar, number of rows of resulting matrix.
<i>c</i>	Scalar, number of columns of resulting matrix.
<i>loc</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, location parameter.
<i>scale</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, scale parameter.
<i>state</i>	Optional argument - scalar or opaque vector.
Scalar case:	
<i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock.	

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`rndLaplace`

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, Laplacian distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$\begin{aligned} E(x) &= \text{location} \\ Var(x) &= 2 * \text{scale}^2 \end{aligned}$$

r and *c* will be truncated to integers if necessary.

Technical Notes

The default generator for **rndLaplace** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndLCbeta

Purpose

Computes beta pseudo-random numbers. NOTE: This function is deprecated--use **rndBeta**--but remains for backward compatibility.

Format

```
{ x, newstate } = rndLCbeta(r, c, a, b, state);
```

Input

<i>r</i>	scalar, number of rows of resulting matrix.
<i>c</i>	scalar, number of columns of resulting matrix.
<i>a</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, first shape argument for beta distribution.
<i>b</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, second shape argument for beta distribution.
<i>state</i>	scalar, or 3x1 vector, or 4x1 vector.

Scalar case:

state = starting seed value only. System default values are used for the additive and multiplicative constants.

The defaults are 1013904223, and 1664525, respectively. These may be changed with [rndcon](#) and [rndmult](#).

If *state* = -1, **GAUSS** computes the starting seed based on the system clock.

3x1 vector case:

- [1] the starting seed, uses the system clock if -1
- [2] the multiplicative constant

L

rndLCbeta

[3] the additive constant

4x1 vector case:

state = the state vector returned from a previous call to one of the **rndLC** random number generators.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, beta distributed random numbers.
<i>newstate</i>	4x1 vector:
	[1] the updated seed
	[2] the multiplicative constant
	[3] the additive constant
	[4] the original initialization seed

Source

randlc.src

Technical Notes

This function uses a linear congruential method, discussed in Kennedy, W.J. Jr., and J.E. Gentle, *Statistical Computing*, Marcel Dekker, Inc. 1980, pp. 136-147. Each seed is generated from the preceding seed using the formula

$$new_seed = ((a * seed) \% 2^{32}) + c) \% 2^{32}$$

where $\%$ is the mod operator and where *a* is the multiplicative constant and *c* is the additive constant.

rndLCgam

Purpose

Computes Gamma pseudo-random numbers. NOTE: This function is deprecated--use `rndGamma`--but remains for backward compatibility.

Format

```
{ x, newstate } = rndLCgam(r, c, alpha, state);
```

Input

<i>r</i>	scalar, number of rows of resulting matrix.
<i>c</i>	scalar, number of columns of resulting matrix.
<i>alpha</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, shape argument for gamma distribution.
<i>state</i>	scalar, or 3x1 vector, or 4x1 vector.

Scalar case:

state = starting seed value only. System default values are used for the additive and multiplicative constants.

The defaults are 1013904223, and 1664525, respectively. These may be changed with `rndcon` and `rndmult`.

If *state* = -1, **GAUSS** computes the starting seed based on the system clock.

3x1 vector case:

- [1] the starting seed, uses the system clock if -1
- [2] the multiplicative constant
- [3] the additive constant

4x1 vector case:

rndLCi

state = the state vector returned from a previous call to one of the **rndLC** random number generators.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, gamma distributed random numbers.
<i>newstate</i>	4x1 vector: <ul style="list-style-type: none"> [1] the updated seed [2] the multiplicative constant [3] the additive constant [4] the original initialization seed

Source

randlc.src

Technical Notes

This function uses a linear congruential method, discussed in Kennedy, W.J. Jr., and J.E. Gentle, *Statistical Computing*, Marcel Dekker, Inc. 1980, pp. 136-147. Each seed is generated from the preceding seed using the formula

$$new_seed = (((a * seed) \% 2^{32}) + c) \% 2^{32}$$

where $\%$ is the mod operator and where *a* is the multiplicative constant and *c* is the additive constant.

rndLCi

Purpose

Returns a matrix of random integers, $0 \leq y < 2^{32}$, and the state of the random number generator. NOTE: This function is deprecated but remains for backward

compatibility.

Format

```
{ y, newstate } = rndLCi(r, c, state);
```

Input

<i>r</i>	scalar, row dimension.
<i>c</i>	scalar, column dimension.
<i>state</i>	scalar, or 3x1 vector, or 4x1 vector.

Scalar case:

state = starting seed value only. System default values are used for the additive and multiplicative constants.

The defaults are 1013904223, and 1664525, respectively. These may be changed with [rndcon](#) and [rndmult](#).

If *state* < 0, **GAUSS** computes the starting seed based on the system clock.

3x1 vector case:

[1] the starting seed, uses the system clock if < 0

[2] the multiplicative constant

[3] the additive constant

4x1 vector case:

state = the state vector returned from a previous call to one of the **rndLC** random number generators.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of random integers between 0 and $2^{32} - 1$,
----------	--

rndLCi

newstate inclusive.
4x1 vector:
[1] the updated seed
[2] the multiplicative constant
[3] the additive constant
[4] the original initialization seed

Remarks

r and c will be truncated to integers if necessary.

Each seed is generated from the preceding seed, using the formula

$$new_seed = ((a * seed) \% 2^{32}) + c) \% 2^{32}$$

where $\%$ is the mod operator and where a is the multiplicative constant and c is the additive constant. The new seeds are the values returned.

Example

```
state = 13;
n = 2000000000;
k = 1000000;
c = 0;
min = 2^32+1;
max = -1;

do while c < n;
    { y, state } = rndLCi(k, 1, state);
    min = minc(min | minc(y));
    max = maxc(max | maxc(y));
    c = c + k;
endo;
```

```
print "min " min;
print "max " max;
```

See Also

[rndLCn](#), [rndLCu](#), [rndcon](#), [rndmult](#)

rndLCn

Purpose

Returns a matrix of standard normal (pseudo) random variables and the state of the random number generator. NOTE: This function is deprecated--use **`rndn`**--but remains for backward compatibility.

Format

```
{ y, newstate } = rndLCn(r, c, state);
```

Input

<i>r</i>	scalar, row dimension.
<i>c</i>	scalar, column dimension.
<i>state</i>	scalar, or 3x1 vector, or 4x1 vector.

Scalar case:

state = starting seed value only. System default values are used for the additive and multiplicative constants.

The defaults are 1013904223, and 1664525, respectively. These may be changed with [rndcon](#) and [rndmult](#).

3x1 vector case:

[1] the starting seed, uses the system clock if < 0

rndLCn

If $state < 0$, **GAUSS** computes the starting seed based on the system clock.

[2] the multiplicative constant

[3] the additive constant

4x1 vector case:

$state$ = the state vector returned from a previous call to one of the **rndLC** random number generators.

Output

y	$r \times c$ matrix of standard normal random numbers.
$newstate$	4x1 vector:
	[1] the updated seed
	[2] the multiplicative constant
	[3] the additive constant
	[4] the original initialization seed

Remarks

r and c will be truncated to integers if necessary.

Example

```
state = 13;
n = 2000000000;
k = 1000000;
c = 0;
submean = {};

do while c < n;
    { y, state } = rndLCn(k, 1, state);
```

```

        submean = submean | meanc(y);
        c = c + k;
    endo;

    mean = meanc(submean);
    print mean;

```

See Also

[rndLCu](#), [rndLCi](#), [rndcon](#), [rndmult](#)

Technical Notes

The normal random number generator is based on the uniform random number generator, using the fast acceptance-rejection algorithm proposed by Kinderman, A.J. and J.G. Ramagge, "Computer Generation of Normal Random Numbers," *Journal of the American Statistical Association*, December 1976, Volume 71, Number 356, pp. 893-896. This algorithm calls the linear congruential uniform random number generator multiple times for each normal random number generated. See [rndLCu](#) for a description of the uniform random number generator algorithm.

rndLCnb

Purpose

Computes negative binomial pseudo-random numbers. NOTE: This function is deprecated--use **rndNegBinomial**--but remains for backward compatibility.

Format

```
{ x, newstate } = rndLCnb(r, c, k, p, state);
```

rndLCnb

Input

r	scalar, number of rows of resulting matrix.
c	scalar, number of columns of resulting matrix.
k	$r \times c$ matrix, or $r \times 1$ vector, or $1 \times c$ vector, or scalar, "event" argument for negative binomial distribution.
p	$r \times c$ matrix, or $r \times 1$ vector, or $1 \times c$ vector, or scalar, "probability" argument for negative binomial distribution.
$state$	scalar, or 3×1 vector, or 4×1 vector.

Scalar case:

$state$ = starting seed value only. System default values are used for the additive and multiplicative constants.

The defaults are 1013904223, and 1664525, respectively. These may be changed with `rndcon` and `rndmult`.

If $state = -1$, **GAUSS** computes the starting seed based on the system clock.

3x1 vector case:

[1] the starting seed, uses the system clock if -1

[2] the multiplicative constant

[3] the additive constant

4x1 vector case:

$state$ = the state vector returned from a previous call to one of the **rndLC** random number generators.

Output

x	$r \times c$ matrix, negative binomial distributed random
-----	---

newstate numbers.
 4x1 vector:
 [1] the updated seed
 [2] the multiplicative constant
 [3] the additive constant
 [4] the original initialization seed

Source

randlc.src

Technical Notes

This function uses a linear congruential method, discussed in Kennedy, W.J. Jr., and J.E. Gentle, *Statistical Computing*, Marcel Dekker, Inc. 1980, pp. 136-147. Each seed is generated from the preceding seed using the formula

$$new_seed = ((a * seed) \% 2^{32}) + c) \% 2^{32}$$

where $\%$ is the mod operator and where a is the multiplicative constant and c is the additive constant.

rndLCp

Purpose

Computes Poisson pseudo-random numbers. NOTE: This function is deprecated--use **rndPoisson**--but remains for backward compatibility.

Format

$\{ x, newstate \} = \mathbf{rndLCp}(r, c, lambda, state);$

rndLCp

Input

<i>r</i>	scalar, row dimension.
<i>c</i>	scalar, column dimension.
<i>lambda</i>	scalar, mean parameter.
<i>state</i>	scalar, or 3x1 vector, or 4x1 vector.

Scalar case:

state = starting seed value only. System default values are used for the additive and multiplicative constants.

The defaults are 1013904223, and 1664525, respectively. These may be changed with [rndcon](#) and [rndmult](#).

3x1 vector case:

[1] the starting seed, uses the system clock if < 0

If *state* < 0, **GAUSS** computes the starting seed based on the system clock.

[2] the multiplicative constant

[3] the additive constant

4x1 vector case:

state = the state vector returned from a previous call to one of the **rndLC** random number generators.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix of Poisson distributed random numbers.
<i>newstate</i>	4x1 vector: [1] the updated seed [2] the multiplicative constant [3] the additive constant

[4] the original initialization seed

Source

randlc.src

Technical Notes

This function uses a linear congruential method, discussed in Kennedy, W.J. Jr., and J.E. Gentle, *Statistical Computing*, Marcel Dekker, Inc. 1980, pp. 136-147. Each seed is generated from the preceding seed using the formula

$$new_seed = ((a * seed) \% 2^{32}) + c) \% 2^{32}$$

where $\%$ is the mod operator and where a is the multiplicative constant and c is the additive constant.

rndLCu

Purpose

Returns a matrix of uniform (pseudo) random variables and the state of the random number generator. NOTE: This function is deprecated but remains for backward compatibility.

Format

$\{ y, newstate \} = \text{rndLCu}(r, c, state);$

Input

r	scalar, row dimension.
c	scalar, column dimension.
$state$	scalar, or 3x1 vector, or 4x1 vector.

L

rndLCu

Scalar case:

state = starting seed value only. System default values are used for the additive and multiplicative constants.

The defaults are 1013904223, and 1664525, respectively. These may be changed with `rndcon` and `rndmult`.

3x1 vector case:

[1] the starting seed, uses the system clock if < 0
If *state* < 0, **GAUSS** computes the starting seed based on the system clock.

[2] the multiplicative constant

[3] the additive constant

4x1 vector case:

state = the state vector returned from a previous call to one of the **rndLC** random number generators.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of uniform ($0 < x < 1$) random numbers.
<i>newstate</i>	4x1 vector: [1] the updated seed [2] the multiplicative constant [3] the additive constant [4] the original initialization seed

Remarks

r and *c* will be truncated to integers if necessary.
Each seed is generated from the preceding seed, using the formula

$$new_seed = ((a * seed) \% 2^{32}) + c) \% 2^{32}$$

where $\%$ is the mod operator and where a is the multiplicative constant and c is the additive constant. A number between 0 and 1 is created by dividing new_seed by 2^{32} .

Example

```
state = 13;
n = 2000000000;
k = 1000000;
c = 0;
submean = {};

do while c < n;
    { y, state } = rndLCu(k,1,state);
    submean = submean | meanc(y);
    c = c + k;
endo;

mean = meanc(submean);
print 0.5-mean;
```

See Also

[rndLCn](#), [rndLCi](#), [rndcon](#), [rndmult](#)

Technical Notes

This function uses a linear congruential method, discussed in Kennedy, W. J. Jr., and J. E. Gentle, *Statistical Computing*, Marcel Dekker, Inc., 1980, pp. 136-147.

rndLCvm

rndLCvm

Purpose

Computes von Mises pseudo-random numbers. NOTE: This function is deprecated but remains for backward compatibility.

Format

```
{ x, newstate } = rndLCvm(r, c, m, k, state);
```

Input

<i>r</i>	scalar, number of rows of resulting matrix.
<i>c</i>	scalar, number of columns of resulting matrix.
<i>m</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, means for vm distribution.
<i>k</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, shape argument for vm distribution.
<i>state</i>	scalar, or 3x1 vector, or 4x1 vector.

Scalar case:

state = starting seed value only. System default values are used for the additive and multiplicative constants.

The defaults are 1013904223, and 1664525, respectively. These may be changed with [rndcon](#) and [rndmult](#).

If *state* = -1, **GAUSS** computes the starting seed based on the system clock.

3x1 vector case:

- [1] the starting seed, uses the system clock if -1
- [2] the multiplicative constant

[3] the additive constant

4x1 vector case:

state = the state vector returned from a previous call to one of the **rndLC** random number generators.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, von Mises distributed random numbers.
<i>newstate</i>	4x1 vector:
	[1] the updated seed
	[2] the multiplicative constant
	[3] the additive constant
	[4] the original initialization seed

Remarks

r and *c* will be truncated to integers if necessary.

Source

randlc.src

Technical Notes

This function uses a linear congruential method, discussed in Kennedy, W.J. Jr., and J.E. Gentle, *Statistical Computing*, Marcel Dekker, Inc. 1980, pp. 136-147. Each seed is generated from the preceding seed using the formula

$$new_seed = ((a * seed) \% 2^{32}) + c) \% 2^{32}$$

where $\%$ is the mod operator and where *a* is the multiplicative constant and *c* is the additive constant.

rndLogNorm

rndLogNorm

Purpose

Computes lognormal pseudo-random numbers with the choice of underlying random number generator.

Format

```
{ x, newstate } = rndLogNorm(r, c, mu, sigma, state);  
x = rndLogNorm(r, c, mu, sigma);
```

Input

<i>r</i>	Scalar, number of rows of resulting matrix.
<i>c</i>	Scalar, number of columns of resulting matrix.
<i>mu</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, mean.
<i>sigma</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, standard deviation.
<i>state</i>	Optional argument - scalar or opaque vector. Scalar case: <i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock. Opaque vector case: <i>state</i> = the state vector returned from a previous call to one of the rnd random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, lognormal distributed random numbers.
----------	---

newstate Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in x are:

$$E(x) = \exp(\mu - 0.5 \cdot \sigma^2)$$

$$Var(x) = (\exp(\sigma^2) - 1) * \exp(2 \cdot \mu + \sigma^2)$$

r and c will be truncated to integers if necessary.

Technical Notes

The default generator for **rndLogNorm** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndMVn

Purpose

Computes multivariate normal random numbers given a covariance matrix.

Format

```
{ r, newstate } = rndMVn(num, mu, cov, state);
r = rndMVn(num, mu, cov);
```

rndMVn

Input

<i>num</i>	Scalar, number of random vectors to create.
<i>mu</i>	Nx1 matrix, mean vector.
<i>cov</i>	NxN covariance matrix.
<i>state</i>	Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number functions.

Output

<i>r</i>	<i>num</i> xN matrix, multivariate normal random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$\begin{aligned}E(x) &= \mu \\Var(x) &= cov\end{aligned}$$

Example

```
//covariance matrix
cov = { 1 0.3,
```

```

        0.3    1  };

//mean for each column of 'cov'
mu = { 0, 0 };

x = rndMVn(100, mu, cov);

```

See Also

[rndCreateState](#), [rndStateSkip](#)

rndMVt

Purpose

Computes multivariate Student-t distributed random numbers given a covariance matrix.

Format

```

{ r, newstate } = rndMVt(num, cov, df, state);
r = rndMVt(num, cov, df);

```

Input

<i>num</i>	Scalar, number of random vectors to create.
<i>cov</i>	NxN covariance matrix.
<i>df</i>	Scalar, degrees of freedom.
<i>state</i>	Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

rndMvt

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number functions.

Output

<i>x</i>	num x N matrix, multivariate student-t distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

$$E(x) = 0$$
$$Var(x) = (df / (df - 2)) * sigma$$

Example

```
//degrees of freedom
df = 8;

//covariance matrix
sigma = { 1 0.3,
          0.3 1 };

x = rndMvt(100, sigma, df);
```

See Also

[rndMVn](#), [rndCreateState](#)

rndn

Purpose

Computes normally distributed pseudo-random numbers with a choice of underlying random number generator.

Format

```
{ y, newstate } = rndn(r, c, state);  
y = rndn(r, c);
```

Input

<i>r</i>	Scalar, row dimension.
<i>c</i>	Scalar, column dimension.
<i>state</i>	Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rndn** random number generators.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of standard normal random numbers.
<i>newstate</i>	Opaque vector, the updated state.

rndn

Remarks

r and c will be truncated to integers if necessary.

Example

Example 1

```
//Create a 100 by 1 vector of standard normal numbers  
my_var = rndn(100, 1);
```

Example 2

This example simulates the linear model: $y = \alpha + \beta_1 X + \varepsilon$

```
num_obs = 100;  
alpha = 2.5;  
beta_1 = 0.8;  
  
//Simulate error term  
err = rndn(num_obs, 1);  
  
//Simulate 'x' variable  
x = rndn(num_obs, 1);  
  
//Simulate data generating process  
y = alpha + beta_1*x + err;
```

Example 3

This example generates two thousand vectors of standard normal random numbers, each with one million elements. The state of the random number generator after each iteration is used as an input to the next generation of random numbers.

```
state = 13;  
n = 2000;  
k = 1000000;  
  
//Create vector to hold 'n' submeans
```

```

submean = zeros(n, 1);

for i(1, n, 1);
    //Create a kx1 vector of random normal numbers,
    //using the optional 'state' input
    { y,state } = rndn(k,1,state);

    submean[i] = mean(y);
endfor;

mean = mean(submean);
print mean;

```

Technical Notes

The default generator for **`rndn`** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **`rndCreateState`**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndnb

Purpose

Computes pseudo-random numbers with negative binomial distribution.

Format

$x = \text{rndnb}(r, c, k, p);$

Input

r scalar, number of rows of resulting matrix.

rndnb

c	scalar, number of columns of resulting matrix.
k	MxN matrix, ExE conformable with $r \times c$ resulting matrix, "event" parameters for negative binomial distribution.
p	KxL matrix, ExE conformable with $r \times c$ resulting matrix, "probability" parameters for negative binomial distribution.

Output

x	$r \times c$ matrix, negative binomial distributed pseudo-random numbers.
-----	---

Remarks

The properties of the pseudo-random numbers in x are:

$$\begin{aligned}E(x) &= k * p / (1 - p) \\Var(x) &= k * p / (1 - p)^2 \\x &= 0, 1, 2, \dots, k \\k &> 0 \\p &> 0 \\p &< 1\end{aligned}$$

Source

random.src
pseudo-random numbers with negative binomial distribution

rndNegBinomial

Purpose

Computes negative binomial pseudo-random numbers with a choice of underlying random number generator.

Format

```
{ x, newstate } = rndNegBinomial(r, c, ns, prob, state);  
x = rndNegBinomial(r, c, ns, prob);
```

Input

<i>r</i>	Scalar, number of rows of resulting matrix.
<i>c</i>	Scalar, number of columns of resulting matrix.
<i>ns</i>	<i>r</i> × <i>c</i> matrix, or <i>r</i> × 1 vector, or 1 × <i>c</i> vector, or scalar, "event" argument for negative binomial distribution.
<i>prob</i>	<i>r</i> × <i>c</i> matrix, or <i>r</i> × 1 vector, or 1 × <i>c</i> vector, or scalar, "probability" argument for negative binomial distribution.
<i>state</i>	Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the state returning random number functions.

rndNegBinomial

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, negative binomial distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *x* are:

```

$$E(x) = \text{num\_s} * (1 - \text{prob}) / \text{prob}$$

$$\text{Var}(x) = \text{num\_s} * (1 - \text{prob}) / \text{prob}^2$$

$$\text{num\_s} > 0$$

$$0 < \text{prob} < 1$$

```

rndNegBinomial has a different parameterization than the deprecated **rndnbn**. To convert a call to **rndnbn** to an equivalent call to **rndNegBinomial**, pass in $1 - \text{prob}$ in place of *prob*. For example, the following two calls are equivalent.

```
x_1 = rndnbn(1e6, 1, 15, 0.3);  
x_2 = rndNegBinomial(1e6, 1, 15, 0.7);
```

r and *c* will be truncated to integers if necessary.

Example

Example 1

Simulate the number of failures before 30 successes where each trial has a 70% probability of success.

```
num_obs = 100;  
  
num_s = 30;
```

```
prob = 0.70;  
  
num_f = rndNegBinomial(num_obs, 1, num_s, prob);
```

Example 2

An alternative parameterization specifies the negative binomial distribution in terms of a dispersion parameter (*dp*) and a mean parameter (*mu*). If you would prefer to think of it in those terms, you may do so by passing in the dispersion parameter *dp*, in place of *num_s* and passing in $dp / (dp + mu)$ in place of *prob*.

```
//dispersion parameter  
dp = 12;  
  
//mean parameter  
mu = 3;  
  
x = rndNegBinomial(100, 1, dp, dp./(dp + mu));
```

Technical Notes

The default generator for **rndNegBinomial** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndp

Purpose

Computes pseudo-random numbers with Poisson distribution.

rndp

Format

```
 $x = \text{rndp}(r, c, \text{lambda});$ 
```

Input

r	scalar, number of rows of resulting matrix.
c	scalar, number of columns of resulting matrix.
lambda	MxN matrix, ExE conformable with $r \times c$ resulting matrix, shape parameters for Poisson distribution.

Output

x	$r \times c$ matrix, Poisson distributed pseudo-random numbers.
-----	---

Remarks

The properties of the pseudo-random numbers in x are:

$E(x)$	=	lambda
$\text{Var}(x)$	=	lambda
x	=	0,1,2,...
lambda	>	0

Source

random.src

rndPoisson

Purpose

Computes Poisson pseudo-random numbers with a choice of underlying random number generator.

Format

```
{ x, newstate } = rndPoisson(r, c, lambda, state);  
x = rndPoisson(r, c, lambda);
```

Input

<i>r</i>	Scalar, number of rows of resulting matrix.
<i>c</i>	Scalar, number of columns of resulting matrix.
<i>lambda</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, mean parameter for Poisson distribution.
<i>state</i>	Optional argument, scalar or opaque vector. Scalar case: <i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock. Opaque vector case: <i>state</i> = the state vector returned from a previous call to one of the rndMT random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, Poisson distributed random numbers.
<i>newstate</i>	Opaque vector, the updated state.

rndStateSkip

Remarks

The properties of the pseudo-random numbers in x are:

$$\begin{aligned}E(x) &= \text{lambda} \\ \text{Var}(x) &= \text{lambda}\end{aligned}$$

r and c will be truncated to integers if necessary.

Example

The example below simulates 100 observations of a Poisson process with a mean of 17.

```
lambda = 17;  
  
x = rndPoisson(100, 1, lambda);
```

Technical Notes

The default generator for **rndPoisson** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndStateSkip

Purpose

To advance a state vector by a specified number of values.

Format

```
newState = rndStateSkip(numSkip, state);
```

Input

<i>numSkip</i>	Scalar, the number of values to skip.
<i>state</i>	Opaque state vector.

Output

<i>newState</i>	Opaque vector, the advanced state.
-----------------	------------------------------------

Example

```
seed = 9192834;

//Create a state from the 118th substream of the
//Wichmann-Hill RNG
state = rndCreateState(wh-118", seed);

//Create a new state that is advanced by 2 numbers.
newState = rndStateSkip(2, state);

//Create and compare numbers from the two state vectors
{ r, state } = rndu(4, 1, state );
{ r2, newState } = rndu(2, 1, newState);
```

```
0.54973563
r = 0.81642451
0.68583300
0.09105558
```

rndu

```
r2 = 0.68583300
      0.09105558
```

Technical Notes

This function applies ONLY to the MRG32K3A and Wichmann-Hill random number generators.

See Also

[rndCreateState](#), [rndn](#), [rndu](#), [rndBeta](#), [rndGamma](#)

rndu

Purpose

Computes uniform random numbers with a choice of underlying random number generator.

Format

```
{ y, newstate } = rndu(r, c, state);
y = rndu(r, c);
```

Input

<i>r</i>	Scalar, row dimension.
<i>c</i>	Scalar, column dimension.
<i>state</i>	Optional argument - scalar, or opaque vector.

Scalar case:

state = starting seed value. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rand** random number generators.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of uniform random numbers, $0 \leq y < 1$.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

r and *c* will be truncated to integers if necessary.

Example

Example 1

Basic usage. If a state or seed is not passed in, then only the random numbers are returned.

```
//Create a 100x1 vector of uniform random numbers
y = randu(100, 1);
```

Example 2

randu can be used to create a vector of random integers in a specified range. The example below, creates 30 random integers in the range [1, 1000].

```
//Largest number in integer range
size = 1000;

//Number of integers to calculate
```

rndu

```
num_indices = 30;

idx = ceil(size .* rndu(num_indices, 1));
```

Example 3

This example generates two thousand vectors of uniform random numbers, each with one million elements. The state of the random number generator after each iteration is used as an input to the next generation of random numbers.

```
//starting seed
state = 13;

//Number of submeans to calculate
n_iters = 2000;

//Number of random numbers to generate
//on each iteration
k = 1000000;

//Pre-allocate 'submean' vector
submean = zeros(n_iters, 1);

for i(1, n_iters, 1);
    { y, state } = rndu(k, 1, state);
    submean[i] = meanc(y);
endfor;

mean = meanc(submean);
print 0.5-mean;
```

Technical Notes

The default generator for **rndu** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndvm

Purpose

Computes von Mises pseudo-random numbers.

Format

```
 $x = \text{rndvm}(r, c, m, k);$ 
```

Input

r	scalar, number of rows of resulting matrix.
c	scalar, number of columns of resulting matrix.
m	$N \times K$ matrix, ExE conformable with $r \times c$, means for von Mises distribution.
k	$L \times M$ matrix, ExE conformable with $r \times c$, shape argument for von Mises distribution.

Output

x	$r \times c$ matrix, von Mises distributed random numbers.
-----	--

Source

random.src

L

rndWeibull

rndWeibull

Purpose

Computes Weibull pseudo-random numbers with the choice of underlying random number generator.

Format

```
{ x, newstate } = rndWeibull(r, c, shape, scale, state);  
x = rndWeibull(r, c, shape, scale);
```

Input

<i>r</i>	Scalar, number of rows of resulting matrix.
<i>c</i>	Scalar, number of columns of resulting matrix.
<i>shape</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, shape parameter.
<i>scale</i>	<i>r</i> x <i>c</i> matrix, or <i>rx1</i> vector, or <i>1xc</i> vector, or scalar, scale parameter.
<i>state</i>	Optional argument - scalar or opaque vector. Scalar case: <i>state</i> = starting seed value only. If -1, GAUSS computes the starting seed based on the system clock. Opaque vector case: <i>state</i> = the state vector returned from a previous call to one of the rnd random number functions.

Output

<i>x</i>	<i>r</i> x <i>c</i> matrix, Weibull distributed random numbers.
----------	---

newstate Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in x are:

$$E(x) = \text{scale} * \text{gamma}(1 + 1/\text{shape})$$

$$\text{Var}(x) = \text{scale}^2 * (\text{gamma}(1 + 2/\text{shape}) - (\text{gamma}(1 + 1/\text{shape}))^2)$$

r and c will be truncated to integers if necessary.

Technical Notes

The default generator for **rndWeibull** is the SFMT Mersenne-Twister 19937. You can specify a different underlying random number generator with the function **rndCreateState**.

See Also

[rndCreateState](#), [rndStateSkip](#)

rndWishart

Purpose

Computes Wishart distributed random numbers given a covariance matrix.

Format

```
{ r, newstate } = rndWishart(numMats, cov, df, state);
r = rndWishart(numMats, cov, df);
```

rndWishart

Input

<i>numMats</i>	Scalar, number of Wishart random matrices to create.
<i>cov</i>	NxM covariance matrix.
<i>df</i>	Scalar, degrees of freedom.
<i>state</i>	Optional argument - scalar or opaque vector.

Scalar case:

state = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

Opaque vector case:

state = the state vector returned from a previous call to one of the **rnd** random number functions.

Output

<i>r</i>	<i>numMats</i> * rows (<i>cov</i>) x N matrix, wishart random matrices.
<i>newstate</i>	Opaque vector, the updated state.

Remarks

The properties of the pseudo-random numbers in *X* are:

$$\begin{aligned}E(X) &= df * cov \\Var(X_{ij}) &= df * (cov^2_{ij} + cov_{ii} * cov_{jj})\end{aligned}$$

Example

```
//covariance matrix
```

```

cov = { 1  0.5,
        0.5 1 };

//degrees of freedom
df = 7;

X = rndWishart(1, cov, df);

X = 7.6019339 4.7744799
     4.7744799 7.7341260

```

See Also

[rndWishartInv](#), [rndMVn](#), [rndCreateState](#)

rndWishartInv

Purpose

Computes the inverse Wishart distributed random numbers given a covariance matrix.

Format

$y = \text{rndWishartInv}(cov, df);$

Input

<i>cov</i>	p x p positive definite covariance matrix.
<i>df</i>	Scalar, degrees of freedom.

rotater

Output

y	Matrix, a random matrix from inverse Wishart distribution.
-----	--

Example

```
rndseed 223;
cov = {1 .5,
       .5 1};
df = 10;

// A random matrix from inverse Wishart distribution
y = rndWishartInv(cov, df);

print y;
```

After above code,

0.081211791	0.036818644
0.036818644	0.097064472

See also

[rndWishart](#), [rndMVn](#), [rndCreateState](#)

rotater

Purpose

Rotates the rows of a matrix.

Format

```
 $y$  = rotater( $x$ ,  $r$ );
```


Input

x	$N \times K$ matrix to be rotated.
r	$N \times 1$ or 1×1 matrix specifying the amount of rotation.

Output

y	$N \times K$ rotated matrix.
-----	------------------------------

Remarks

The rotation is performed horizontally within each row of the matrix. A positive rotation value will cause the elements to move to the right. A negative rotation value will cause the elements to move to the left. In either case, the elements that are pushed off the end of the row will wrap around to the opposite end of the same row.

If the rotation value is greater than or equal to the number of columns in x , then the rotation value will be calculated using $(r \% \text{cols}(x))$.

Example

```
y = rotater(x,r);
```

If $x =$	1 2 3	and $r =$	1	Then $y =$	3 1 2
	4 5 6		-1		5 6 4

	1 2 3		0		1 2 3
If $x =$	4 5 6	and $r =$	1	Then $y =$	6 4 5
	7 8 9		2		8 9 7
	10 11 12		3		10 11 12

round

See Also

[shiftr](#)

round

Purpose

Round to the nearest integer.

Format

```
y = round(x);
```

Input

x	NxK matrix or N-dimensional array.
---	------------------------------------

Output

y	NxK matrix or N-dimensional array containing the rounded elements of x.
---	---

Example

```
let x = { 77.68 -14.10,  
          4.73 -158.88 };  
y = round(x);  
print y;
```

```
78.00 -14.00
5.00 -159.00
```

See Also

[trunc](#), [floor](#), [ceil](#)

rows

Purpose

Returns the number of rows in a matrix.

Format

```
y = rows(x);
```

Input

x NxK matrix or sparse matrix.

Output

y scalar, number of rows in the specified matrix.

Remarks

Use **getorders** to return both the number of rows and columns in one call. If **x** is an empty matrix, **rows(x)** and **cols(x)** return 0.

Example

```
x = ones(3,5);
```

rowsf

```
y = rows(x);  
print x;
```

```
1.00 1.00 1.00  
1.00 1.00 1.00  
1.00 1.00 1.00
```

```
print y;
```

```
3.00
```

See Also

[cols](#), [getorders](#), [show](#)

rowsf

Purpose

Returns the number of rows in a **GAUSS** data set (.dat) file or **GAUSS** matrix (.fmt) file.

Format

```
y = rowsf(f);
```

Input

f	file handle of an open file.
-----	------------------------------

Output

y	scalar, number of rows in the specified file.
-----	---

Example

```
open fp = wilshire.dat;  
r = rowsf(fp);  
c = colsf(fp);  
print r;
```

```
324.00
```

```
print c;
```

```
7.00
```

See Also

[colsf](#), [open](#), [typedef](#)

rref

Purpose

Computes the reduced row echelon form of a matrix.

Format

```
y = rref(x);
```

Input

x	MxN matrix.
---	-------------

Output

y	MxN matrix containing reduced row echelon form of
---	---

rref

$$X.$$
Remarks

The tolerance used for zeroing elements is computed inside the procedure using:

```
tol = maxc(m|n) * eps * maxc(abs(sumc(x')));
```

where $eps = 2.24e-16$.

This procedure can be used to find the rank of a matrix. It is not as stable numerically as the singular value decomposition (which is used in the **rank** function), but it is faster for large matrices.

There is some speed advantage in having the number of rows be greater than the number of columns, so you may want to transpose if all you care about is the rank.

The following code can be used to compute the rank of a matrix:

```
r = sumc(sumc(abs(y')) .> tol);
```

where y is the output from **rref**, and tol is the tolerance used. This finds the number of rows with any nonzero elements, which gives the rank of the matrix, disregarding numeric problems.

Example

```
// Since (row 2) = 2*(row 1), we do not expect this
// matrix to have full rank
x[3,3] = 1 2 3
        2 4 6
        3 5 2;
y = rref(x);

// compute rank of x
```

run

```
r = sumc(sumc(abs(rref(x)))) .> 1e-15);  
print "The rank of x = " r;
```

```
The rank of x = 2.000
```

Source

rref.src

run

Purpose

Runs a source code or compiled code program.

Format

```
run filename;  
run -r filename;
```

Input

<i>filename</i>	literal or ^string, name of file to run.
<i>-r</i>	flag, returns control to the calling program.

Remarks

The filename can be any legal file name. Filename extensions can be whatever you want, except for the compiled file extension, .gcg. Pathnames are okay. If the name is to be taken from a string variable, then the name of the string variable must be preceded by the ^ (caret) operator.

run

The `run` statement can be used both from the command line and within a program. If used in a program, once control is given to another program through the `run` statement, there is no return to the original program unless the flag `-r` is used.

If you specify a filename without an extension, **GAUSS** will first look for a compiled code program (i.e., a `.gcg` file) by that name, then a source code program by that name. For example, if you enter

```
run dog;
```

GAUSS will first look for the compiled code file `dog.gcg`, and run that if it finds it. If **GAUSS** cannot find `dog.gcg`, it will then look for the source code file `dog` with no extension.

If a path is specified for the file, then no additional searching will be attempted if the file is not found.

If a path is not specified, the current directory will be searched first, then each directory listed in `src_path`. The first instance found is run. `src_path` is defined in `gauss.cfg`.

```
run  
/gauss/myprog.prg;
```

No additional
search will be
made if the file
is not found.

```
run myprog.prg;
```

The directories
listed in `src_`
`path` will be
searched for
`myprog.prg`
if the file is not
found in the
current
directory.

Programs can also be run by typing the filename on the OS command line when starting **GAUSS**.

Example

Example 1

```
run myprog.prg;
```

Example 2

```
name = "myprog.prg";  
run ^name;
```

Example 3

```
x = rndn(3,3);  
run -r myprog.prg;  
y = inv(x);  
e = x*y;
```

In this case, **GAUSS** will execute the lines after the `run` command. If the `-r` is omitted, the lines following the `run` command will not be executed within a program.

See Also

[#include](#)

sampleData

s

sampleData

Purpose

Returns a sample of the rows of a matrix, chosen with or without replacement

Format

```
s = sampleData(x, size);  
s = sampleData(x, size, replace);
```

Input

<i>x</i>	matrix, population from which to take a sample
<i>size</i>	scalar, the requested sample size
<i>replace</i>	Optional argument. Scalar, if <i>replace</i> is 0, the sample is drawn without replacement. If <i>replace</i> is 1, the sample is drawn with replacement. Default is 0.

Output

<i>s</i>	<i>size</i> x cols (<i>x</i>) matrix, containing the sample taken from <i>x</i> .
----------	--

Examples

Example 1: Basic example without replacement

```
//Set seed for repeatable random draws
rndseed 23423;

//Create a 7x1 vector
x = { 1,
      2,
      3,
      4,
      5,
      6,
      7 };

//Take a sample of 3 elements without replacement
s = sampleData(x, 3);
```

After running the code above, s is equal to:

```
5
3
7
```

Example 2: Basic example with replacement

```
//Set seed for repeatable random draws
rndseed 23423;

//Create a 7x2 vector
x = { 1.2 1.8,
      2.7 2.1,
      3.0 3.3,
      4.8 4.1,
      5.1 5.4,
```

satostrC

```
        6.0 2.8,  
        7.2 3.9 };  
  
replace = 1;  
  
//Take a sample of 5 rows of 'x' with replacement  
sample = sampleData(x, 5, replace);
```

After running the code above, *sample* is equal to:

```
5.1    5.4  
3.0    3.3  
6.0    2.8  
4.8    4.1  
3.0    3.3
```

Remarks

Indices for taking a random sample can be created with **GAUSS** function **rndi**.

The random number generator used in **sampleData** to choose the samples is automatically seeded using the system clock when **GAUSS** first starts. However, that can be overridden using the **rndseed** statement.

See Also

[rndi](#), [rndn](#), [rndseed](#)

satostrC

Purpose

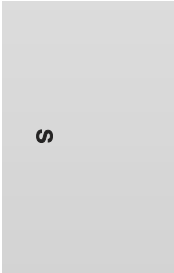
Copies from one string array to another using a C language format specifier string for each element.

Format

```
y = satostrC(sa, fmt);
```

Input

sa	NxM string array.
fmt	1x1, 1xM, or Mx1 format specifier for each element copy.



Output

y	NxM formatted string array.
---	-----------------------------

Examples

Example 1: Basic example

```
//Create a 3x1 column vector
length = { 12, 25, 18 };

//Convert numeric data to a string array
length = ntos(length);

//Add '(cm)' after each number
fmt = "%s (cm)";

length_fmt = satostrc(length, fmt);
```

After the code above, *length_fmt* will equal:

```
"12 (cm) "
"25 (cm) "
"18 (cm) "
```

Example 2: Different formats for each column

```
//Create numeric matrices
year = { 2012, 2013, 2014 };
beef = { 187.9, 183.6, 224.1 };
fish = { 4.8, 6.8, 6.6 };

//Create a 3x3 matrix using horizontal concatenation
commodity_prices = year ~ beef ~ fish;

//%s indicates the location of the contents of the original
string
fmt = "Year %s" $~ "%s cts/lb" $~ "%s $/kg";

//Convert the numeric matrix to a string
```

```
commodity_prices = ntos(commodity_prices);

//Apply formatting
commodity_prices_fmt = satostrC(commodity_prices, fmt);
```

After the code above, *commodity_prices_fmt* should be equal to:

"Year 2012"	"187.9 cts/lb"	"4.8 \$/kg"
"Year 2013"	"183.6 cts/lb"	"6.8 \$/kg"
"Year 2014"	"224.1 cts/lb"	"6.6 \$/kg"

Source

strfns.src

See Also

[strcombine](#)

save

Purpose

Saves matrices, strings, or procedures to a disk file.

Format

```
save vflag path=path x, lpath=y;
save path=path x;
save x;
```

Input

<i>vflag</i>	version flag.
--------------	---------------

save

S

- v89 not supported
- v92 supported on UNIX, Windows
- v96 supported on all platforms

See also **FOREIGN LANGUAGE INTERFACE**, CHAPTER 1, for details on the various versions. The default format can be specified in `gauss.cfg` by setting the `dat_fmt_version` configuration variable. If `dat_fmt_version` is not set, the default is `v96`.

<i>path</i>	literal or ^string, a default path to use for this and subsequent <code>save</code> 's.
<i>x</i>	a symbol name, the name of the file the symbol will be saved in is the same as this with the proper extension added for the type of the symbol.
<i>lpath</i>	literal or ^string, a local path and filename to be used for a particular symbol. This path will override the path previously set and the filename will override the name of the symbol being saved. The extension cannot be overridden.
<i>y</i>	the symbol to be saved to <i>lpath</i> .

Remarks

`save` can be used to save matrices, strings, procedures, and functions. Procedures and functions must be compiled and resident in memory before they can be `save`'d.

The following extensions will be given to files that are `save`'d:

matrix	.fmt
--------	------

string	.fst
procedure	.fcg
function	.fcg
keyword	.fcg

If the **path=** subcommand is used with **save**, the path string will be remembered until changed in a subsequent command. This path will be used whenever none is specified. The **save** path can be overridden in any particular **save** by specifying an explicit path and filename.

Example

```
spath = "/gauss";
save path = ^spath x,y,z;
```

Save *x*, *y*, and *z* using /gauss as the path. This path will be used for the next **save** if none is specified.

```
svp = "/gauss/data";
save path = ^svp n, k, /gauss/quad1=quad;
```

n and *k* will be saved using /gauss/data as the **save** path, *quad* will be saved in /gauss with the name *quad1.fmt*. On platforms that use the backslash as the path separator, the double backslash is required inside double quotes to produce a backslash because it is the escape character in quoted strings. It is not required when specifying literals.

```
save path=/procs;
```

Change **save** path to /procs.

```
save path = /miscdata;
save /data/mydata1 = x, y, hisdata = z;
```

saveall

In the above program:

x would be saved in `/data/mydata1.fmt`

y would be saved in `/miscdata/y.fmt`

z would be saved in `/miscdata/hisdata.fmt`

See Also

[datasave](#), [load](#), [saveall](#), [saved](#)

saveall

Purpose

Saves the current state of the machine to a compiled file. All procedures, global matrices and strings will be saved.

Format

```
saveall fname;
```

Input

<i>fname</i>	literal or ^string, the path and filename of the compiled file to be created.
--------------	---

Remarks

The file extension will be `.gcg`.

A file will be created containing all your matrices, strings, and procedures. No main code segment will be saved. This just means it will be a `.gcg` file with no main program code (see [compile](#)). The rest of the contents of memory will be saved,

including all global matrices, strings, functions and procedures. Local variables are not saved. This can be used inside a program to take a snapshot of the state of your global variables and procedures. To reload the compiled image, use `run` or `use`.

```
library pgraph;  
external proc xy, logx, logy, loglog, hist;  
saveall pgraph;
```

This would create a file called `pgraph.gcg`, containing all the procedures, strings and matrices needed to run **Publication Quality Graphics** programs. Other programs could be compiled very quickly with the following statement at the top of each:

```
use pgraph;
```

See Also

[compile](#), [run](#), [use](#)

saved

Purpose

Writes a matrix in memory to a **GAUSS** data set on disk.

Format

```
y = saved(x, dataset, vnames);
```

Input

<i>x</i>	NxK matrix to save in .dat file.
<i>dataset</i>	string, name of data set.
<i>vnames</i>	string or Kx1 character vector, names for the columns

saved

of the data set.

Output

y scalar, 1 if successful, otherwise 0.

Remarks

If *dataset* is null or 0, the data set name will be `temp.dat`.

If *vnames* is a null or 0, the variable names will begin with "X" and be numbered 1-K.

If *vnames* is a string or has fewer elements than *x* has columns, it will be expanded as explained under [create](#).

The output data type is double precision.

Example

```
x = rndn(100,3);
dataset = "mydata";
vnames = { height, weight, age };

if not saved(x,dataset,vnames);
    errorlog "Write error";
end;
endif;
```

Source

saveload.src

See Also

[load](#), [writer](#), [create](#)

savestruct

Purpose

Saves a matrix of structures to a file on the disk.

Format

```
retcode = saveStruct(instance, file_name);
```

Input

<i>instance</i>	MxN matrix, instances of a structure.
<i>file_name</i>	string, name of file on disk to contain matrix of structures.

Output

<i>retcode</i>	scalar, 0 if successful, otherwise it will be non-zero.
----------------	---

Remarks

The file on the disk will be given a `.fsr` extension

Example

```
struct DS p0;  
p0 = reshape(dsCreate(), 2, 3);  
retc = saveStruct(p2, "p2");
```

savewind

savewind

Purpose

Save the current graphic panel configuration to a file. Note: This function is for use with the deprecated PQG graphics.

Library

pgraph

Format

```
err = savewind(filename);
```

Input

<i>filename</i>	string, name of file.
-----------------	-----------------------

Output

<i>err</i>	scalar, 0 if successful, 1 if graphic panel matrix is invalid. Note that the file is written in either case.
------------	--

Remarks

See the discussion on using graphics panels in **Tiled Graphic Panels**, Section 1.0.1.

Source

pwindow.src

See Also

[loadwind](#)

scale

Purpose

Fixes the scaling for subsequent graphs. The axes endpoints and increments are computed as a best guess based on the data passed to it. Note: This function is for use with the deprecated PQG graphics.

Library

pgraph

Format

```
scale(x, y);
```

Input

x	matrix, the X axis data.
y	matrix, the Y axis data.

Remarks

x and y must each have at least 2 elements. Only the minimum and maximum values are necessary.

This routine fixes the scaling for all subsequent graphs until **graphset** is called. This also clears **xtics** and **yticks** whenever it is called.

If either of the arguments is a scalar missing, the main graphics function will set the scaling for that axis using the actual data.

If an argument has 2 elements, the first will be used for the minimum and the last will be used for the maximum.

scale3d

If an argument has 2 elements, and contains a missing value, that end of the axis will be scaled from the data by the main graphics function.

If you want direct control over the axes endpoints and tick marks, use **xtics** or **ytics**. If **xtics** or **ytics** have been called after **scale**, they will override **scale**.

Source

pscale.src

See Also

[xtics](#), [ytics](#), [ztics](#), [scale3d](#)

scale3d

Purpose

Fixes the scaling for subsequent graphs. The axes endpoints and increments are computed as a best guess based on the data passed to it. Note: This function is for use with the deprecated PQG graphics.

Library

pgraph

Format

```
scale3d(x, y, z);
```

Input

<i>x</i>	matrix, the X axis data.
<i>y</i>	matrix, the Y axis data.
<i>z</i>	matrix, the Z axis data.

Remarks

x , y and z must each have at least 2 elements. Only the minimum and maximum values are necessary.

This routine fixes the scaling for all subsequent graphs until **graphset** is called. This also clears **xtics**, **ytics** and **ztics** whenever it is called.

If any of the arguments is a scalar missing, the main graphics function will set the scaling for that axis using the actual data.

If an argument has 2 elements, the first will be used for the minimum and the last will be used for the maximum.

If an argument has 2 elements, and contains a missing value, that end of the axis will be scaled from the data by the main graphics function.

If you want direct control over the axes endpoints and tick marks, use **xtics**, **ytics**, or **ztics**. If one of these functions have been called, they will override **scale3d**.

Source

pscale.src

See Also

[scale](#), [xtics](#), [ytics](#), [ztics](#)

scalerr

Purpose

Tests for a scalar error code.

Format

```
 $y$  = scalerr( $c$ );
```

scalerr

Input

c	$N \times K$ matrix or sparse matrix or N -dimensional array, generally the return argument of a function or procedure call.
-----	--

Output

y	scalar or $[N-2]$ -dimensional array, 0 if the argument is not a scalar error code, or the value of the error code as an integer if the argument is an error code.
-----	--

Remarks

Error codes in **GAUSS** are NaN's (Not A Number). These are not just scalar integer values. They are special floating point encodings that the math chip recognizes as not representing a valid number. See also **error**.

scalerr can be used to test for either those error codes that are predefined in **GAUSS** or an error code that the user has defined using **error**.

If c is an N -dimensional array, y will be an $[N-2]$ -dimensional array, where each element corresponds to a 2-dimensional array described by the last two dimensions of c . For each 2-dimensional array in c that does not contain a scalar error code, its corresponding element in y will be set to zero. For each 2-dimensional array in c that does contain a scalar error code, its corresponding element in y will be set to the value of that error code as an integer. In other words, if c is a $5 \times 5 \times 10 \times 10$ array, y will be a 5×5 array, in which each element corresponds to a 10×10 array in c and contains either a zero or the integer value of a scalar error code.

If c is an empty matrix, **scalerr** will return 65535.

Certain functions will either return an error code or terminate a program with an error message, depending on the trap state. The **trap** command is used to set the trap state.

The error code that will be returned will appear to most commands as a missing value code, but the **scalerr** function can distinguish between missing values and error codes and will return the value of the error code.

Following are some of the functions that are affected by the trap state:

	trap 1	trap 0
function	error code	error message
chol	10	Matrix not positive definite
invpd	20	Matrix not positive definite
solpd	30	Matrix not positive definite
/	40	Matrix not positive definite (second argument not square)
	41	Matrix singular (second argument is square)
inv	50	Matrix singular

Example

```
trap 1;
cm = invpd(x);
trap 0;

if scalerr(cm);
    cm = inv(x);
endif;
```

In this example **invpd** will return a scalar error code if the matrix **x** is not positive definite. If **scalerr** returns with a nonzero value, the program will use the **inv** function, which is slower, to compute the inverse. Since the trap state has been turned off, if **inv** fails, the program will terminate with a Matrix singular error message.

scalinfnanmiss

See Also

[error](#), [trap](#), [trapchk](#)

scalinfnanmiss

Purpose

Returns true if the argument is a scalar infinity, NaN, or missing value.

Format

```
y = scalinfnanmiss(x);
```

Input

x	NxK matrix.
---	-------------

Output

y	scalar, 1 if x is a scalar, infinity, NaN, or missing value, else 0.
---	--

Example

```
//Create an infinity
x = 1/0;

if scalInfNanMiss(x);
    print "x = " x;
else;
    print "x is Not: a Nan, Infinity, or Missing";
endif;
```

See Also

[isinfnanmiss](#), [ismiss](#), [scalmiss](#)

scalmiss

Purpose

Tests to see if its argument is a scalar missing value.

Format

```
y = scalmiss(x);
```

Input

x	NxK matrix.
-----	-------------

Output

y	scalar, 1 if argument is a scalar missing value, 0 if not.
-----	--

Remarks

scalmiss first tests to see if the argument is a scalar. If it is not scalar, **scalmiss** returns a 0 without testing any of the elements.

To test whether any element of a matrix is a missing value, use **ismiss**. **scalmiss** will execute much faster if the argument is a large matrix, since it will not test each element of the matrix but will simply return a 0.

An element of x is considered to be a missing if and only if it contains a missing value in the real part. Thus, **scalmiss** and **scalmiss** would return a 1 for complex $x = . + 1i$, and a 0 for $x = 1 + .i$.

schtoc

Example

```
clear s;
do until eof(fp);
    y = readr(fp,nr);
    y = packr(y);
    if scalmiss(y);
        continue;
    endif;
    s = s+sumc(y);
endo;
```

In this example the **packr** function will return a scalar missing if every row of its argument contains missing values, otherwise it will return a matrix that contains no missing values. **scalmiss** is used here to test for a scalar missing returned from **packr**. If the test returns true, then the sum step will be skipped for that iteration of the read loop because there were no rows left after the rows containing missings were packed out.

schtoc

Purpose

Reduces any 2x2 blocks on the diagonal of the real Schur matrix returned from **schur**. The transformation matrix is also updated.

Format

```
{ schc, transc } = schtoc(sch, trans);
```

Input

<i>sch</i>	real NxN matrix in Real Schur form, i.e., upper triangular except for possibly 2x2 blocks on the
------------	--

<i>trans</i>	diagonal. real NxN matrix, the associated transformation matrix.
--------------	---

Output

<i>schc</i>	NxN matrix, possibly complex, strictly upper triangular. The diagonal entries are the eigenvalues.
<i>transc</i>	NxN matrix, possibly complex, the associated transformation matrix.

Remarks

Other than checking that the inputs are strictly real matrices, no other checks are made. If the input matrix *sch* is already upper triangular, it is not changed. Small off-diagonal elements are considered to be zero. See the source code for the test used.

Example

```
{ schc, transc } = schtoc(schur(a));
```

This example calculates the complex Schur form for a real matrix *a*.

Source

schtoc.src

See Also

[schur](#)

schur

schur

Purpose

Computes the real or complex Schur form of a square matrix.

Format

```
{ S, Z } = schur(A);  
{ S, Z } = schur(A, flag);
```

Input

<i>A</i>	KxK matrix.
<i>flag</i>	String, to control whether output should be in real or complex schur form. Valid options include: <ul style="list-style-type: none">• "complex"• "real"

Output

<i>S</i>	KxK matrix, Schur form.
<i>Z</i>	KxK matrix, transformation matrix.

Examples

Example 1: Real matrix with all real eigenvalues

```
//Create a 2 x 2 matrix  
A = { 7 -2,  
      12 -5 };
```



```
//Calculate eigenvalues of 'A'
lambda = eig(A);
```

After the code above, *lambda* should equal:

```
4.4641
-2.4641
```

```
//Continuing with 'A' from above
{ S, Z } = schur(A);
```

Now *S* and *Z* should equal:

```
S = 4.4641    -14.000    Z = 0.6193    -0.7852
      0      -2.4641      0.7852    0.6193
```

Example 2: Real matrix with some complex eigenvalues

```
//Create a 3 x 3 matrix
A = { 1  -4  -1,
      3  -1   9,
     -9   1  -2 };
```

```
//Calculate real schur form, with complex eigenvalues
//stored as 2 x 2 blocks on the diagonal
{ S_r, Z_r } = schur(A, "real");
```

```
//Calculate complex schur form
{ S_c, Z_c } = schur(A, "complex");
```

After the code above:

```
S_r = -4.1991    10.2489    0.8409    Z_r = -0.2103    -0.6911    -
0.6915
      -2.8394    -4.1991    -2.2342      -0.8736    -0.1847
0.4502
           0           0    6.3981      0.4389    -0.6988
```

schur

0.5649

```
S_c = -4.1991+5.3945i  -2.5084+6.9720i   1.0168-0.7763i
        0+0i  -4.1991-5.3945i   1.9825-0.3630i
        0+0i           0+0i       6.3981+0i
```

```
Z_c =  0.3275 - 0.1759i   0.5326 + 0.3160i  -0.6915
        0.1102 - 0.7700i   0.0023 + 0.4385i   0.4502
        0.3132 + 0.3984i   0.6502 + 0.0373i   0.5649
```

Remarks

If a real matrix is passed in without a flag variable, the real Schur form will be returned. If a complex matrix is passed in without a flag variable, **GAUSS** will check to see if any of the imaginary elements are greater than *imagtol* (2.23e-16 by default). If any imaginary elements are greater than *imagtol*, the complex Schur form will be calculated, otherwise the real Schur form will be returned. If a real flag is passed in with a complex matrix, the flag will be ignored and the complex Schur factorization will be returned.

The real Schur form is an upper quasi-triangular matrix, that is, it is block triangular where the blocks are 2x2 submatrices which correspond to complex eigenvalues of A . If A has no complex eigenvalues, S will be strictly upper triangular. To convert the real Schur form of S to the complex Schur form, use the **Run-Time Library** function **schtoc**.

Z is an orthogonal matrix that transforms A into S and vice versa. Thus

$$S = Z' * A * Z;$$

and since Z is orthogonal,

$$A = Z * S * Z';$$

See Also

[hess](#), [shtoc](#)

screen

Purpose

Controls output to the screen.

Format

```
screen on;  
screen off;  
screen;
```

Remarks

- When this is **on**, the results of all print statements will be directed to the window. When this is **off**, print statements will not be sent to the window. This is independent of the statement **output on**, which will cause the results of all print statements to be routed to the current auxiliary output file.
- If you are sending a lot of output to the auxiliary output file on a disk drive, turning the window off will speed things up.
- The **end** statement will automatically perform **output off** and **screen on**.
- **screen** with no arguments will print "Screen is on" or "Screen is off" on the console.
- Changing the **screen** setting is NOT threadsafe and therefore, should not be done inside of **threadbegin**, **threadstat** or **threadfor** blocks.

Example

```
output file = mydata.asc reset;  
screen off;
```

searchsourcepath

```
format /m1/rz 1,8;
open fp = mydata;
do until eof(fp);
    print readr(fp,200);;
endo;
fp = close(fp);
end;
```

The program above will write the contents of the **GAUSS** file `mydata.dat` into an ASCII file called `mydata.asc`. If `mydata.asc` already exists, it will be overwritten.

Turning the window off will speed up execution. The `end` statement above will automatically perform output off and **screen on**.

See Also

[output](#), [end](#), [new](#)

searchsourcepath

Purpose

Searches the source path and (if specified) the `src` subdirectory of the **GAUSS** installation directory for a specified file.

Format

```
fpath = searchsourcepath(fname, srcdir);
```

Input

<i>fname</i>	string, name of file to search for.
--------------	-------------------------------------

<i>srcdir</i>	scalar, one of the following: 0 do not search in the <code>src</code> subdirectory of the GAUSS installation directory. 1 search the <code>src</code> subdirectory first. 2 search the <code>src</code> subdirectory last.
---------------	--

Output

<i>fpath</i>	string, the path of <i>fname</i> , or null string if <i>fname</i> is not found.
--------------	---

Remarks

The source path is set by the *src_path* configuration variable in your **GAUSS** configuration file, `gauss.cfg`.

seekr

Purpose

Moves the pointer in a `.dat` or `.fmt` file to a particular row.

Format

y = **seekr**(*fh*, *r*);

Input

<i>fh</i>	scalar, file handle of an open file.
<i>r</i>	scalar, the row number to which the pointer is to be moved.

select (dataloop)

Output

y	scalar, the row number to which the pointer has been moved.
-----	---

Remarks

If $r = -1$, the current row number will be returned.

If $r = 0$, the pointer will be moved to the end of the file, just past the end of the last row.

rowsf returns the number of rows in a file.

```
seekr(fh, 0) == rowsf(fh) + 1;
```

Do NOT try to seek beyond the end of a file.

See Also

[open](#), [readr](#), [rowsf](#)

select (dataloop)

Purpose

Selects specific rows (observations) in a data loop based on a logical expression.

Format

```
select logical_expression;
```

Remarks

Selects only those rows for which *logical_expression* is TRUE. Any

variables referenced must already exist, either as elements of the source data set, as `extern`'s, or as the result of a previous `make`, `vector`, or `code` statement.

Example

```
select age > 40 AND sex $== 'MALE';
```

See Also

[delete \(dataloop\)](#)

selif

Purpose

Selects rows from a matrix. Those selected are the rows for which there is a 1 in the corresponding row of `e`.

Format

```
y = selif(x, e);
```

Input

x	NxK matrix or string array.
e	Nx1 vector of 1's and 0's.

Output

y	MxK matrix or string array consisting of the rows of x for which there is a 1 in the corresponding row of e.
---	--

selif

Remarks

The argument `e` will usually be generated by a logical expression using "dot" operators.

`y` will be a scalar missing if no rows are selected.

Example

```
y = selif(x,x[.,2] .gt 100);
```

This example selects all rows of `x` in which the second column is greater than 100.

```
let x[3,3] = 0 10 20
             30 40 50
             60 70 80;

e =(x[.,1] .gt 0) .and (x[.,3] .lt 100);
y = selif(x,e);
```

The resulting matrix `y` is:

```
30 40 50
60 70 80
```

All rows for which the element in column 1 is greater than 0 and the element in column 3 is less than 100 are placed into the matrix `y`.

See Also

[delif](#), [scalmiss](#)

seqa, seqm

Purpose

seqa creates an additive sequence. **seqm** creates a multiplicative sequence.

Format

```
y = seqa(start, inc, n);  
y = seqm(start, inc, n);
```

Input

<i>start</i>	scalar specifying the first element.
<i>inc</i>	scalar specifying increment.
<i>n</i>	scalar specifying the number of elements in the sequence.

Output

<i>y</i>	<i>n</i> ×1 vector containing the specified sequence.
----------	---

Remarks

For **seqa**, *y* will contain a first element equal to *start*, the second equal to *start + inc*, and the last equal to *start + inc*(n-1)*.

For instance,

```
seqa(1, 1, 10);
```

will create a column vector containing the numbers 1, 2, ...10.

setarray

For **seqm**, *y* will contain a first element equal to *start*, the second equal to *start * inc*, and the last equal to *start * incⁿ⁻¹*.

For instance,

```
seqm(10,10,10);
```

will create a column vector containing the numbers 10, 100,...10¹⁰.

Example

```
a = seqa(2,2,10)';  
print a;
```

```
2 4 6 8 10 12 14 16 18 20
```

```
m = seqm(2,2,10)';  
print m;
```

```
2 4 8 16 32 64 128 512 1024
```

Note that the results have been transposed in this example. Both functions return Nx1 (column) vectors.

See Also

[recserar](#), [recsercp](#)

setarray

Purpose

Sets a contiguous subarray of an N-dimensional array.

Format

```
setarray a, loc, src;
```

Input

<i>a</i>	N-dimensional array.
<i>loc</i>	Mx1 vector of indices into the array to locate the subarray of interest, where M is a value from 1 to N.
<i>src</i>	[N-M]-dimensional array, matrix, or scalar.

Remarks

`setarray` resets the specified subarray of *a* in place, without making a copy of the entire array. Therefore, it is faster than `putarray`.

If *loc* is an Nx1 vector, then *src* must be a scalar. If *loc* is an [N-1]x1 vector, then *src* must be a 1-dimensional array or a 1xL vector, where L is the size of the fastest moving dimension of the array. If *loc* is an [N-2]x1 vector, then *src* must be a KxL matrix, or a KxL 2-dimensional array, where K is the size of the second fastest moving dimension.

Otherwise, if *loc* is an Mx1 vector, then *src* must be an [N-M]-dimensional array, whose dimensions are the same size as the corresponding dimensions of array *a*.

Example

```
a = arrayalloc(2|3|4|5|6,0);
src = arrayinit(4|5|6,5);
loc = { 2,1 };
setarray a,loc,src;
```

This example sets the contiguous 4x5x6 subarray of *a* beginning at [2,1,1,1,1] to the array *src*, in which each element is set to the specified value 5.

setdif

See Also

[putarray](#)

setdif

Purpose

Returns the unique elements in one vector that are not present in a second vector.

Format

```
y = setdif(v1, v2);  
y = setdif(v1, v2, typ);
```

Input

v1	Nx1 vector.
v2	Mx1 vector.
typ	Optional input, scalar, type of data. 0 character, case sensitive. 1 numeric (Default). 2 character, case insensitive.

Output

y	Lx1 vector containing all unique values that are in v1 and are not in v2, sorted in ascending order.
---	--

Examples

Example 1: Basic example

```
//Create a vector of years
y1 = { 1980,
      1984,
      1988,
      1992,
      1996,
      2000,
      2004,
      2008,
      2012,
      2016 };

y2 = { 1980,
      1988,
      1992,
      2000,
      2008,
      2016 };

//Set 'y_diff' equal to years in 'y1' and NOT in 'y2'
y_diff = setdif(y1, y2);
```

After the code above, *y_diff* will be equal to:

```
1984
1996
2004
2012
```

Example 2: Character data

```
//Create 2 vectors of character data using the
```

setdif

```
//numeric concatenation operator
sp500 = "aapl" | "goog" | "msft" | "xom" | "wfc" | "jnj";
nasdaq = "aapl" | "msft" | "amzn" | "goog" | "fb" | "gild";

//Set type to 'character, case insensitive'
typ = 2;

//Find characters in sp500 and NOT in nasdaq
sp_only = setdif(sp500, nasdaq, 2);

//NOTE: The $ in front of the variable name tell
//GAUSS to print the variable as character data
print $sp_only;
```

The code above will produce the following output:

```
jnj
wfc
xom
```

Remarks

Place smaller vector first for fastest operation.

When there are a lot of duplicates, it is faster to remove them first with `unique` before calling this function.

Source

setdif.src

See Also

[setdifsa](#), [union](#)

setdifsa

Purpose

Returns the unique elements in one string vector that are not present in a second string vector.

Format

```
sy = setdifsa(sv1, sv2);
```

Input

<i>sv1</i>	Nx1 or 1xN string vector.
<i>sv2</i>	Mx1 or 1xM string vector.

Output

<i>sy</i>	Lx1 vector containing all unique values that are in <i>sv1</i> and are not in <i>sv2</i> , sorted in ascending order.
-----------	---

Remarks

Place smaller vector first for fastest operation.

When there are a lot of duplicates it is faster to remove them first with **unique** before calling this function.

Example

```
string sv1 = { "mary", "jane", "linda", "john" };  
string sv2 = { "mary", "sally" };
```

setvars

```
sy = setdifsa(sv1,sv2);
```

Now *sy* is equal to:

```
jane  
john  
linda
```

Source

setdif.src

See Also

[setdif](#)

setvars

Purpose

Reads the variable names from a data set header and creates global matrices with the same names.

Format

```
nvec = setvars(dataset);
```

Input

<i>dataset</i>	string, the name of the GAUSS data set. Do not use a file extension.
----------------	---

Output

<i>nvec</i>	Nx1 character vector, containing the variable names
-------------	---

defined in the data set.

Remarks

setvars is designed to be used interactively.

Example

```
nvec = setvars ("freq");
```

Source

vars.src

See Also

[makevars](#)

setvwrmode

Purpose

Sets the graphics viewer mode. NOTE: This function is for use with the deprecated PQG graphics.

Library

pgraph

Format

```
oldmode = setvwrmode(mode);
```

setwind

Input

<i>mode</i>	string, new mode or null string.
"one"	Use only one viewer.
"many"	Use a new viewer for each graph.

Output

<i>oldmode</i>	string, previous <i>mode</i> .
----------------	--------------------------------

Remarks

If *mode* is a null string, the current *mode* will be returned with no changes made.

If "one" is set, the viewer executable will be `vwr.exe`.

Example

```
oldmode = setvwrmode("one");  
call setvwrmode(oldmode);
```

Source

`pgraph.src`

See Also

[pqgwin](#)

setwind

Purpose

Sets the current graphic panel to a previously created graphic panel number.

NOTE: This function is for use with the deprecated PQG graphics. Use

`plotLayout` instead.

Library

pgraph

Format

`setwind(n);`

Input

n scalar, graphic panel number.

Remarks

This function selects the specified graphic panel to be the current graphic panel. This is the graphic panel in which the next graph will be drawn.

See the discussion on using graphic panels in **Graphic Panels**, Section 1.1.

Source

pwindow.src

See Also

[begwind](#), [endwind](#), [getwind](#), [nextwind](#), [makewind](#), [window](#)

shell

Purpose

Executes an operating system command.

shell

Format

```
shell stmt;
```

Input

<i>stmt</i>	literal or ^string, the command to be executed.
-------------	---

Remarks

`shell` lets you run shell commands and programs from inside **GAUSS**. If a command is specified, it is executed; when it finishes, you automatically return to **GAUSS**. If no command is specified, the shell is executed and control passes to it, so you can issue commands interactively. You have to type **exit** to get back to **GAUSS** in that case.

If you specify a command in a string variable, precede it with the ^ (caret) as shown in the examples below.

Example

```
comstr = "ls ./src";  
shell ^comstr;
```

This lists the contents of the `./src` subdirectory, then returns to **GAUSS**.

```
shell cmp n1.fmt n1.fmt.old;
```

This compares the matrix file `n1.fmt` to an older version of itself, `n1.fmt.old`, to see if it has changed. When **cmp** finishes, control is returned to **GAUSS**.

```
shell;
```

This executes an interactive shell. The OS prompt will appear and OS commands or other programs can be executed. To return to **GAUSS**, type **exit**.

See Also

[exec](#)

shiftr

Purpose

Shifts the rows of a matrix.

Format

$y = \text{shiftr}(x, s, f);$

Input

x	NxK matrix to be shifted.
s	scalar or Nx1 vector specifying the amount of shift.
f	scalar or Nx1 vector specifying the value to fill in.

Output

y	NxK shifted matrix.
-----	---------------------

Remarks

The shift is performed within each row of the matrix, horizontally. If the shift value is positive, the elements in the row will be moved to the right. A negative shift value causes the elements to be moved to the left. The elements that are pushed off the end of the row are lost, and the fill value will be used for the new elements on the other end.

shiftr

Example

s

```
x = { 1 2,  
      3 4 };  
s = { 1,  
      1 };  
f = { 99,  
      999 };  
y = shiftr(x,s,f);
```

Now *y* is equal to:

```
99  1  
4   999
```

```
x = { 1 2 3,  
      4 5 6,  
      7 8 9 };  
s = { 0,  
      1,  
      2 };  
f = 0;  
y2 = shiftr(x,s,f);
```

Now *y2* is equal to:

```
1  2  3  
0  4  5  
0  0  7
```

See Also

[rotater](#)

show

Purpose

Displays the global symbol table.

Format

```
show -flags symbol;  
show -flags;  
show symbol;  
show;
```

Input

<i>flags</i>		flags to specify the symbol type that is shown.
	<i>k</i>	keywords
	<i>p</i>	procedures
	<i>f</i>	fn functions
	<i>m</i>	matrices
	<i>s</i>	strings
	<i>g</i>	show only symbols with global references
	<i>l</i>	show only symbols with all local references
<i>symbol</i>		the name of the symbol to be shown. If the last character is an asterisk (*), all symbols beginning with the supplied characters will be shown.

Remarks

If there are no arguments, the entire symbol table will be displayed.

show is directed to the auxiliary output if it is open.

show

Here is an example listing with an explanation of the columns. Note that `show` does not display the column titles shown here:

```
Memory used Name Cplx Type References Info
128 bytes a MATRIX 4,4
672 bytes add KEYWORD global refs 0=1
192 bytes area FUNCTION local refs 1=1
256 bytes c C MATRIX 4,4
296 bytes p1 PROCEDURE local refs 1=1
384 bytes p2 PROCEDURE global refs 0=1
8 bytes ps1 STRUCT sdat *
16 bytes s STRING 8 char
312 bytes s1 STRUCT sdat 1,1
40 bytes sa STRING ARRAY 3,1
56 bytes sm SPARSE MATRIX 15,15
2104 bytes token PROCEDURE local refs 2=1
216 bytes y ARRAY 3 dims 2,3,4
672 bytes program space used
12 global symbols, 2000 maximum, 12 shown
0 active locals, 2000 maximum
1 active structure
```

The 'Memory used' column gives the amount of memory used by each item.

The 'Name' column gives the name of each symbol.

The 'Cplx' column contains a 'C' if the symbol is a complex matrix.

The 'Type' column specifies the type of the symbol. It can be ARRAY, FUNCTION, KEYWORD, MATRIX, PROCEDURE, STRING, STRING ARRAY, or STRUCT.

If the symbol is a procedure, keyword or function, the 'References' column will show if it makes any global references. If it makes only local references, the procedure or function can be saved to disk in an `.fcg` file with the `save` command. If the function or procedure makes any global references, it cannot be saved in an `.fcg` file.

If the symbol is a structure, the 'References' column will contain the structure type. A structure pointer is indicated by a * following the structure type.

The 'Info' column depends on the type of the symbol. If the symbol is a procedure or a function, it gives the number of values that the function or procedure returns and the number of arguments that need to be passed to it when it is called. If the symbol is a matrix, sparse matrix, string array or array of structures, then the 'Info' column gives the number of rows and columns. If the symbol is a string, then it gives the number of characters in the string. If the symbol is an N-dimensional array, then it gives the orders of each dimension. As follows:

Rets=Args	if procedure, keyword, or function
Row,Col	if matrix, sparse matrix, string array, or structure
Length	if string
OrdN,...,Ord2,Ord1	if array, where N is the slowest moving dimension of the array, and Ord is the order (or size) of a dimension

sin

If the symbol is an array of structures, the 'Info' column will display the size of the array. A scalar structure instance is treated as a 1x1 array of structures. If the symbol is a structure pointer, the 'Info' column will be blank.

The program space is the area of space reserved for all nonprocedure, nonfunction program code. The maximum program space can be controlled by the [new](#) command.

The maximum number of global and local symbols is controlled by the *maxglobals* and *maxlocals* configuration variables in *gauss.cfg*.

Example

```
show -fpg eig*;
```

This command will show all functions and procedures that have global references and begin with **eig**.

```
show -m;
```

This command will show all matrices.

See Also

[new](#), [delete](#)

sin

Purpose

Returns the sine of its argument.

Format

```
y = sin(x);
```

Input

x	$N \times K$ matrix or N -dimensional array.
-----	--

Output

y	$N \times K$ matrix or N -dimensional array containing the sine of x .
-----	--

Remarks

For real data, x should contain angles measured in radians.
To convert degrees to radians, multiply the degrees by $\pi/180$.

Example

```
let x = { 0, .5, 1, 1.5 };
y = sin(x);
print y;
```

```
0.000000
0.479426
0.841471
0.997495
```

See Also

[atan](#), [cos](#), [sinh](#), [pi](#)

singleindex

Purpose

Converts a vector of indices for an N -dimensional array to a scalar vector index.

singleindex

Format

```
si = singleindex(i, o);
```

Input

<i>i</i>	Nx1 vector of indices into an N-dimensional array.
<i>o</i>	Nx1 vector of orders of an N-dimensional array.

Output

<i>si</i>	scalar, index of corresponding element in 1-dimensional array or vector.
-----------	--

Remarks

This function and its opposite, **arrayindex**, allow you to convert between an N-dimensional index and its corresponding location in a 1-dimensional object of the same size.

Example

```
orders = { 2,3,4 };

a = arrayalloc(orders,0);
ai = { 2,1,3 };
setarray a, ai, 49;
v = vecr(a);
vi = singleindex(ai,orders);

print "ai = " ai;
print "vi = " vi;
print "getarray(a,ai) = " getarray(a,ai);
print "v[vi] = " v[vi];
```

produces:

```
ai =  
    2.0000000  
    1.0000000  
    3.0000000  
vi = 15.000000  
getarray(a,ai) = 49.000000  
v[vi] = 49.000000
```

This example allocates a 3-dimensional array *a* and sets the element corresponding to the index vector *ai* to 49. It then creates a vector, *v*, with the same data. The element in the array *a* that is indexed by *ai* corresponds to the element of the vector *v* that is indexed by *vi*.

See Also

[arrayindex](#)

sinh

Purpose

Computes the hyperbolic sine.

Format

```
y = sinh(x);
```

Input

<i>x</i>	NxK matrix.
----------	-------------

sleep

Output

y

NxK matrix containing the hyperbolic sines of the elements of *x*.

Example

```
let x = { -0.5, -0.25, 0, 0.25, 0.5, 1 };  
x = x * pi;  
y = sinh(x);
```

The above statement produces, *y* equal to:

```
-2.301299  
-0.868671  
0.000000  
0.868671  
2.301299  
11.548739
```

Source

trig.src

sleep

Purpose

Sleeps for a specified number of seconds.

Format

```
unslept = sleep(secs);
```

Input

<i>secs</i>	scalar, number of seconds to sleep.
-------------	-------------------------------------

Output

<i>unslept</i>	scalar, number of seconds not slept.
----------------	--------------------------------------

Remarks

secs does not have to be an integer. If your system does not permit sleeping for a fractional number of seconds, *secs* will be rounded to the nearest integer, with a minimum value of 1.

If a program sleeps for the full number of *secs* specified, **sleep** returns 0; otherwise, if the program is awakened early (e.g., by a signal), **sleep** returns the amount of time not slept.

A program may sleep for longer than *secs* seconds, due to system scheduling.

solpd

Purpose

Solves a set of positive definite linear equations.

Format

$x = \text{solpd}(b, A);$

Input

<i>b</i>	NxK matrix or M-dimensional array where the last two dimensions are NxK.
----------	--

solpd

A

$N \times N$ symmetric positive definite matrix or M -dimensional array where the $N \times N$ 2-dimensional arrays described by the last two dimensions are symmetric and positive definite.

Output

x

$N \times K$ matrix or M -dimensional array where the last two dimensions are $N \times K$, the solutions for the system of equations, $Ax = b$.

Remarks

b can have more than one column. If so, the system of equations is solved for each column, i.e., $A * x[:, i] = b[:, i]$.

This function uses the Cholesky decomposition to solve the system directly. Therefore it is more efficient than using `inv(A)*b`.

If b and A are M -dimensional arrays, the sizes of their corresponding $M-2$ leading dimensions must be the same. The resulting array will contain the solutions for the system of equations given by each of the corresponding 2-dimensional arrays described by the two trailing dimensions of b and A . In other words, for a $10 \times 4 \times 2$ array b and a $10 \times 4 \times 4$ array A , the resulting array x will contain the solutions for each of the 10 corresponding 4×2 arrays contained in b and 4×4 arrays contained in A . Therefore, $A[n,...] * x[n,...] = b[n,...]$, for $1 \leq n \leq 10$.

solpd does not check to see that the matrix A is symmetric. **solpd** will look only at the upper half of the matrix including the principal diagonal.

If the A matrix is not positive definite:

- trap 1** return scalar error code 30.
- trap 0** terminate with an error message.

One obvious use for this function is to solve for least squares coefficients. The effect of this function is thus similar to that of the `/` operator.

If X is a matrix of independent variables, and Y is a vector containing the dependent variable, then the following code will compute the least squares coefficients of the regression of Y on X :

```
b = solpd(X'Y,X'X);
```

Example

```
n = 5;
format /lo 16,8;

A = randn(n,n);
A = A'A;
x = randn(n,1);
b = A*x;

x2 = solpd(b,A);

print " X solpd(b,A) Difference";
print x~x2~x-x2;
```

produces:

```
X solpd(b,A) Difference

0.32547881    0.32547881   -4.9960036e-16
1.5190182     1.5190182    -1.7763568e-15
0.88099266    0.88099266     1.5543122e-15
1.8192784     1.8192784    -2.2204460e-16
-0.060848175 -0.060848175   -1.4710455e-15
```

See Also

[chol](#), [invpd](#), [trap](#)

sortc, sortcc

sortc, sortcc

Purpose

Sorts a numeric matrix, character matrix or string array.

Format

```
y = sortc(x, c);  
y = sortcc(x, c);
```

Input

x	NxK matrix, or string array.
c	scalar specifying one column of x to sort on.

Output

y	NxK matrix equal to x and sorted on the column c .
-----	--

Examples

Example 1: Sort rows of a matrix based upon first column

```
x = { 4 7 3,  
      1 3 2,  
      3 4 8 };  
  
//Sort 'x' based upon the first row  
y = sortc(x,1);
```

The above example code produces, y equal to:

```
1 3 2
3 4 8
4 7 3
```

Example 2: Sort rows of a 5x1 string vector

```
//Create a 5x1 string array, using the string
//vertical concatenation operator '$|'
letters = "epsilon" $|
          "gamma" $|
          "beta" $|
          "alpha" $|
          "delta";

//Sort 'letters'
letters_s = sortc(letters,1);
```

The above example code produces, *letters_s* equal to:

```
alpha
beta
delta
epsilon
gamma
```

Remarks

- These functions will sort the rows of a matrix with respect to a specified column. That is, they will sort the elements of a column and will arrange all rows of the matrix in the same order as the sorted column.
- **sortc** assumes that the column to sort on is numeric. **sortcc** assumes that the column to sort on contains character data.
- The matrix may contain both character and numeric data, but the sort column must be all of one type.
- Missing values will sort as if their value is below $-\infty$.

sortd

- The sort will be in ascending order.
- This function uses the Quicksort algorithm.
- If you need to obtain the matrix sorted in descending order, you can use:

```
rev(sortc(x, c))
```

See Also

[rev](#), [sortind](#), [unique](#)

sortd

Purpose

Sorts a data file on disk with respect to a specified variable.

Format

```
sortd(infile, outfile, keyvar, keytyp);
```

Input

<i>infile</i>	string, name of input file.
<i>outfile</i>	string, name of output file, must be different.
<i>keyvar</i>	string, name of key variable.
<i>keytyp</i>	scalar, type of key variable. <ul style="list-style-type: none">1 numeric key, ascending order.2 character key, ascending order.-1 numeric key, descending order.-2 character key, descending order.

Remarks

The data set *infile* will be sorted on the variable *keyvar*, and will be placed in *outfile*.

If the inputs are null ("" or 0), the procedure will ask for them.

Source

sortd.src

See Also

[sortmc](#), [sortc](#), [sortec](#), [sorthc](#), [sorthcc](#)

sorthc, sorthcc

Purpose

Sorts a matrix of numeric or character data, or a string array.

Format

```
y = sorthc(x, c);
y = sorthcc(x, c);
```

Input

<i>x</i>	NxK matrix or string array.
<i>c</i>	scalar specifying one column of <i>x</i> to sort on.

Output

<i>y</i>	NxK matrix or string array equal to <i>x</i> and sorted on the column <i>c</i> .
----------	--

sorthc, sorthcc

Remarks

These functions will sort the rows of a matrix or string array with respect to a specified column. That is, they will sort the elements of a column and will arrange all rows of the object in the same order as the sorted column.

sorthc assumes that the column to sort on is numeric. **sorthcc** assumes that the column to sort on contains character data.

If *x* is a matrix, it may contain both character and numeric data, but the sort column must be all of one type. Missing values will sort as if their value is below $-\infty$.

The sort is in ascending order. This function uses the heap sort algorithm.

If you need to obtain the matrix sorted in descending order, you can use:

```
rev(sorthc(x, c))
```

Example

```
let x[3,3]= 4 7 3
            1 3 2
            3 4 8;

//Sort x based upon the values in the third column
y = sorthc(x,3);
```

This produces *y* equal to:

```
1 3 2
4 7 3
3 4 8
```

See Also

[sortc](#), [rev](#)

sortind, sortindc

Purpose

Returns the sorted index of x .

Format

```
ind = sortind(x);  
ind = sortindc(x);
```

Input

x	$N \times 1$ column vector.
-----	-----------------------------

Output

<i>ind</i>	$N \times 1$ vector representing sorted index of x .
------------	--

Remarks

sortind assumes that x contains numeric data. **sortindc** assumes that x contains character data.

This function can be used to sort several matrices in the same way that some other reference matrix is sorted. To do this, create the index of the reference matrix, then use **submat** to rearrange the other matrices in the same way.

Example

```
//Create uniform random integers between 0 and 10  
x = round(10*randu(10, 1));  
  
ind = sortind(x);
```

sortmc

```
y = x[ind];
```

After running the above code:

```
          9.00
          8.00
x  =    0.00
          4.00
          6.00

          3.00
          4.00
ind =    5.00
          2.00
          1.00

          0.00
          4.00
y  =    6.00
          8.00
          9.00
```

sortmc

Purpose

Sorts a matrix on multiple columns.

Format

```
y = sortmc(x, v);
```


Input

x	NxK matrix to be sorted.
v	Lx1 vector containing integers specifying the columns, in order, that are to be sorted. If an element is negative, that column will be interpreted as character data.

Output

y	NxK sorted matrix.
---	--------------------

Example

sortmc keeps all rows together. After it sorts on the first specified column, it will continue to sort the rows of the matrix using the other specified columns ONLY when there is a tie in the first column. For example:

```
x = { 9 2 5 6,
      3 6 1 9,
      3 7 4 1,
      1 2 8 9 };

s1 = sortc(x,1);

sm = sortmc(x, 1|2);
```

will return:				
	1	2	8	9
s1 =	3	7	4	1
	3	6	1	9
	9	2	5	6
	1	2	8	9

sortr, sortrc

```
sm = 3      6      1      9
      3      7      4      1
      9      2      5      6
```

In the output above, we see that the difference between *s1* and *sm* is that the second and third rows have been switched. This is because **sortmc** first sorted the matrix based upon row one like **sortc**. Then **sortmc** sorted the rows in which the first column was the same (in our example they are both threes), based upon the values in the second column.

Source

sortmc.src

See Also

[sortd](#), [sortc](#), [sortcc](#), [sorthc](#), [sorthec](#)

sortr, sortrc

Purpose

Sorts the columns of a matrix of numeric or character data, with respect to a specified row.

Format

```
y = sortr(x, r);
y = sortrc(x, r);
```

Input

<i>x</i>	NxK matrix.
<i>r</i>	scalar, row of <i>x</i> on which to sort.

Output

y NxK matrix equal to x and sorted on row r .

Remarks

These functions sort the columns of a matrix with respect to a specified row. That is, they sort the elements of a row and arrange all rows of the matrix in the same order as the sorted column.

sortr assumes the row on which to sort is numeric. **sortrc** assumes that the row on which to sort contains character data.

The matrix may contain both character and numeric data, but the sort row must be all of one type. Missing values will sort as if their value is below $-\infty$.

The sort will be in left to right ascending order. This function uses the Quicksort algorithm. If you need to obtain the matrix sorted left to right in descending order (i.e., ascending right to left), use:

```
rev(sortr(x, r)')
```

Example

```
//Create a 5 x 3 matrix of random integers
//between 1 and 30
x = ceil(30*randu(5, 3));

//Sort the columns based upon the first row
y = sortr(x,1);
```

Examine the variables after the code above. Notice that the columns remain the same, but their order has changed.

spBiconjGradSol

```
      10.000  21.000  18.000
      11.000  30.000  20.000
x = 10.000  23.000   7.000
      6.000   9.000  20.000
      7.000   4.000  30.000
```

```
      10.000  18.000  21.000
      11.000  20.000  30.000
y = 10.000   7.000  23.000
      6.000  20.000   9.000
      7.000  30.000   4.000
```

If we were to use the same x , but sort on the 5th row:

```
y2 = sortr(x, 5);
```

We get the following result:

```
      21.000  10.000  18.000
      30.000  11.000  20.000
y2 = 23.000  10.000   7.000
      9.000   6.000  20.000
      4.000   7.000  30.000
```

spBiconjGradSol

Purpose

Attempts to solve the system of linear equations $Ax = b$ using the biconjugate gradient method where A is a sparse matrix.

Format

```
x = spBiconjGradSol(a, b, epsilon, maxit);
```

Input

<i>a</i>	NxN, sparse matrix.
<i>b</i>	Nx1, dense vector.
<i>epsilon</i>	Method tolerance: If epsilon is set to 0, the default tolerance is set to 1e-6.
<i>maxit</i>	Maximum number of iterations. If maxit is set to 0, the default setting is 300 iterations.

Output

<i>x</i>	Nx1 dense vector.
----------	-------------------

Example

```
nz = { 33.446  82.641 -12.710 -25.062   0.000,  
       0.000 -26.386  17.016  21.576 -45.273,  
       0.000 -42.331 -47.902   0.000   0.000,  
       0.000 -26.517 -22.135 -76.827  31.920,  
       10.364 -29.843 -20.277   0.000  65.816 };  
b = { 10.349,  
      -3.117,  
       4.240,  
       0.013,  
       2.115 };  
  
sparse matrix a;  
a = densetosp(nz,0);  
  
//Setting the third and fourth arguments to 0 employs the  
//default tolerance and maxit settings  
x = spBiconjGradSol(a,b,0,0);
```

spBiconjGradSol

```
//Solve the system of equations using the '/' operator for
//comparison
x2 = b/a;
```

The output from the above code:

```
      0.135
      0.055
x =   -0.137
      0.018
     -0.006

      0.135
      0.055
x2 =  -0.137
      0.018
     -0.006
```

Remarks

If convergence is not reached within the maximum number of iterations allowed, the function will either terminate the program with an error message or return an error code which can be tested for with the **scalerr** function. This depends on the trap state as follows:

trap 1	return error code: 60
trap 0	terminate with error message: Unable to converge in allowed number of iterations.

If matrix A is not well conditioned use the / operator to perform the solve. If the matrix is symmetric, **spConjGradSol** will be approximately twice as fast as **spBiconjGradSol**.

See Also

[spConjGradSol](#)

spChol

Purpose

Computes the LL' decomposition of a sparse matrix A.

Format

```
l = spChol(a);
```

Input

<i>a</i>	NxN, symmetric, positive definite sparse matrix.
----------	--

Output

<i>l</i>	NxN lower-triangular sparse matrix.
----------	-------------------------------------

Example

```
sparse matrix A;  
sparse matrix L;  
  
//Create a small, simple positive-definite matrix  
let x = { 9.53984224e+001 -5.84272701e+000 1.99970335e+001,  
          -5.84272701e+000 1.09765831e+002 2.52038945e+000,  
          1.99970335e+001 2.52038945e+000 4.71834812e+000  
};  
  
//Create the sparse matrix A from x, keeping all elements
```

spConjGradSol

```
A = denseToSp(x, 0);

//Create matrix factorization
L = spChol(A);
```

See Also

[spLDL](#), [spLU](#)

Technical Notes

spChol implements functions from the TAUCS library: TAUCS Version 2.2. Copyright ©2001, 2002, 2003 by Sivan Toledo, Tel-Aviv University, stoledo@tau.ac.il. All Rights Reserved.

spConjGradSol

Purpose

Attempts to solve the system of linear equations $Ax = b$ using the conjugate gradient method where A is a symmetric sparse matrix.

Format

```
x = spConjGradSol(a, b, epsilon, maxit);
```

Input

<i>a</i>	NxN, symmetric sparse matrix.
<i>b</i>	Nx1, dense vector.
<i>epsilon</i>	Method tolerance: If epsilon is set to 0, the default tolerance is set to 1e-6.
<i>maxit</i>	Maximum number of iterations. If maxit is set to 0, the

default setting is 300 iterations.

Output

x Nx1 dense vector

Example

```

nz = {   0.000   2845.607    0.000    0.000    0.000,
        2845.607  10911.430    0.000    0.000    0.000,
          0.000    0.000  3646.798  2736.338 -2674.440,
          0.000    0.000  2736.338  7041.526 -3758.528,
          0.000    0.000 -2674.440 -3758.528  7457.899 };

sparse matrix a;

//Set 'a' to be a sparse matrix with the same contents as
//the dense matrix 'nz'
a = densetosp(nz,0);

//Create our right-hand-side
b = { 10.349,
      -3.117,
       4.240,
       0.013,
       2.115 };

//Setting the third and fourth arguments to 0 employs the
//default tolerance maxit settings
x = spConjGradSol(a,b,0,0);

newb = a*x;

```

The results from the above code are:

spConjGradSol

s

```

-0.01504075
 0.00363683
x  =  0.00203504
-0.00033936
 0.00084234

10.34900000
-3.11700000
newb = 4.24000000
 0.01300000
 2.11500000
```

Remarks

If convergence is not reached within the maximum number of iterations allowed, the function will either terminate the program with an error message or return an error code which can be tested for with the **scalerr** function. This depends on the trap state as follows:

trap 1	return error code: 60
trap 0	terminate with error message: Unable to converge in allowed number of iterations.

If matrix A is not symmetric or well conditioned use the **/** operator to perform the solve. For a nonsymmetric, but well conditioned matrix A, use **spBiconjGradSol**.

See Also

[spBiconjGradSol](#)

spCreate

Purpose

Creates a sparse matrix from vectors of non-zero values, row indices, and column indices.

Format

```
y = spCreate(r, c, vals, rinds, cinds);
```

Input

<i>r</i>	scalar, rows of output matrix.
<i>c</i>	scalar, columns of output matrix.
<i>vals</i>	Nx1 vector, non-zero values.
<i>rinds</i>	Nx1 vector, row indices of corresponding non-zero values.
<i>cinds</i>	Nx1 vector, column indices of corresponding non-zero values.

Output

<i>y</i>	<i>r</i> x <i>c</i> sparse matrix.
----------	------------------------------------

Remarks

Since sparse matrices are strongly typed in **GAUSS**, *y* must be defined as a sparse matrix before the call to **spCreate**.

spDenseSubmat

Example

```
//Declare 'y' to be a sparse matrix
sparse matrix y;

//Create the non-zero values to place in the sparse matrix
vals = { 1.7, 2.4, 3.2, 4.5 };

//Set the row and column indices for the location in which
//to place each successive element of 'vals' into the new
//matrix
rinds = { 2,5,8,13 };
cinds = { 4,1,9,5 };

y = spCreate(15,10,vals,rinds,cinds);
```

This example creates a 15x10 sparse matrix *y*, containing the following non-zero values:

Non-zero value	Index
1.7	(2,4)
2.4	(5,1)
3.2	(8,9)
4.5	(13,5)

See Also

[packedToSp](#), [denseToSp](#), [spEye](#)

spDenseSubmat

Purpose

Returns a dense submatrix of a sparse matrix.

Format

```
y = spDenseSubmat(x, rinds, cinds);
```

Input

<i>x</i>	MxN sparse matrix.
<i>rinds</i>	Kx1 vector, row indices.
<i>cinds</i>	Lx1 vector, column indices.

Output

<i>y</i>	KxL dense matrix, the intersection of <i>rinds</i> and <i>cinds</i> .
----------	---

Remarks

If *rinds* or *cinds* are scalar zeros, all rows or columns will be returned.

Example

```
sparse matrix y;  
x = { 0  0  0 10,  
      0  2  0  0,  
      0  0  0  0,  
      5  0  0  0,  
      0  0  0  3 };  
  
//Set 'y' to be a sparse matrix with the same values as 'x'  
y = denseToSp(x,0);  
  
//Extract a submatrix from 'y' with all rows of 'y' and  
//columns 1, 3 and 4
```

spDiagRvMat

```
d = spDenseSubmat(y, 0, 1 | 3 | 4);
```

Now d is equal to:

```
0  0  10
0  0   0
0  0   0
5  0   0
0  0   3
```

See Also

[spSubmat](#)

spDiagRvMat

Purpose

Inserts submatrices along the diagonal of a sparse matrix.

Format

```
y = spDiagRvMat(x, inds, size, a);
```

Input

x	MxN sparse matrix.
$inds$	Kx2 vector or scalar 0, row and column indices into x at which to place the corresponding submatrices in a .
$size$	Kx2 vector or scalar 0, sizes of the corresponding submatrices in a .
a	KxLxP array, containing the submatrices to insert into x .

Output

y	MxN sparse matrix, a copy of x containing the specified insertions.
-----	---

Remarks

Each row of *inds* must contain the row and column indices, respectively, that form the starting point for the insertion of the corresponding submatrix in *a*. If *inds* is a scalar 0, the starting point for the insertion of each submatrix will be one row and one column past the ending point of the previous insertion. The first insertion will begin at the [1,1] element.

Each row of *size* must contain the number of rows and columns in the corresponding submatrix in *a*. This allows you to insert submatrices of different sizes $L_i \times P_i$ by inserting them into the planes of an array that is $K \times \text{MAX}(L) \times \text{MAX}(P)$ and padding the submatrices with zeros to $\text{MAX}(L) \times \text{MAX}(P)$. For each plane in *a*, **spDiagRvMat** extracts the submatrix $a[i, 1:\text{size}[i,1], 1:\text{size}[i,2]]$ and inserts that into *x* at the location indicated by the corresponding row of *inds*. If *size* is a scalar 0, then each $L \times P$ plane of *a* is inserted into *x* as is.

Example

```
declare sparse matrix x,y;

//Create a 10x10 sparse identity matrix
x = spEye(10);

sx1 = { 2 3, 5 8 };
sx2 = { 8 2 3 4, 7 9 5 6, 3 2 8 4 };
sx3 = { 4 7 2, 6 5 3 };
sx4 = { 9, 3 };

//Create a 4x3x4 dimensional array with every element set
```

5

```
//to 0
a = arrayinit(4|3|4,0);

//Set some of the array values
a[1,1:2,1:2] = sx1;
a[2,...] = sx2;
a[3,1:2,1:3] = sx3;
a[4,1:2,1] = sx4;
```

The value of a is now:

Plane [1,...]

2.00000000	3.00000000	0.00000000	0.00000000
5.00000000	8.00000000	0.00000000	0.00000000
0.00000000	0.00000000	0.00000000	0.00000000

Plane [2,...]

8.00000000	2.00000000	3.00000000	4.00000000
7.00000000	9.00000000	5.00000000	6.00000000
3.00000000	2.00000000	8.00000000	4.00000000

Plane [3,...]

4.00000000	7.00000000	2.00000000	0.00000000
6.00000000	5.00000000	3.00000000	0.00000000
0.00000000	0.00000000	0.00000000	0.00000000

Plane [4,...]

9.00000000	0.00000000	0.00000000	0.00000000
3.00000000	0.00000000	0.00000000	0.00000000
0.00000000	0.00000000	0.00000000	0.00000000


```
inds = 0;
siz = { 2 2, 3 4, 2 3, 2 1 };

y = spDiagRvMat(x,inds,siz,a);
```

The output, in variable *y*, is:

```
2 3 0 0 0 0 0 0 0 0
5 8 0 0 0 0 0 0 0 0
0 0 8 2 3 4 0 0 0 0
0 0 7 9 5 6 0 0 0 0
0 0 3 2 8 4 0 0 0 0
0 0 0 0 0 1 4 7 2 0
0 0 0 0 0 0 6 5 3 0
0 0 0 0 0 0 0 1 0 9
0 0 0 0 0 0 0 0 1 3
0 0 0 0 0 0 0 0 0 1
```

spEigv

Purpose

Computes a specified number of eigenvalues and eigenvectors of a square, sparse matrix *a*.

Format

```
{ va, ve } = spEigv(a, nev, which, tol, maxit, ncv);
```

Input

<i>a</i>	NxN square, sparse matrix.
<i>nev</i>	Scalar, number of eigenvalues to compute.
<i>which</i>	String, may be one of the following: "LM" largest

spEigv

s

	magnitude, "LR" largest real, "LI" largest imaginary, "SR" smallest real, or "SI" smallest imaginary. Default input 0, sets <i>which</i> to "LM."
<i>tol</i>	Scalar, tolerance for eigenvalues. Default input 0, sets <i>tol</i> to 1e-15.
<i>maxit</i>	Scalar, maximum number of iterations. Default input 0, sets <i>maxit</i> to <i>nevx</i> (columns of <i>a</i>)x100.
<i>ncv</i>	Scalar, size of Arnoldi factorization. The minimum setting is the greater of <i>nev</i> +2 and 20. See Remarks on how to set <i>ncv</i> . Default input 0, sets <i>ncv</i> to 2x <i>nev</i> +1.

Output

<i>va</i>	<i>nevx</i> 1 dense vector containing the computed eigenvalues of input matrix <i>a</i> .
<i>ve</i>	Nx <i>nev</i> dense matrix containing the corresponding eigenvectors of input matrix <i>a</i> .

Example

<pre>rndseed 3456; sparse matrix a; x = 10*rndn(5,5); a = densetosp(x,4);</pre>				
a =				
21.276135	5.4078872	-19.817044	9.6771132	-19.211952
0.0000000	-4.4011007	10.445221	-5.1742289	-16.336474
-12.637055	8.1227002	0.0000000	-8.7817892	0.0000000
0.0000000	-7.8181517	15.326816	0.0000000	0.0000000

```
{ va, ve } = spEigv(a,2,0,0,0,0);
/* equivalent to call { va, ve } = spEigv(a,2,"LM",1e-
15,2*5*100,5); */
```

```
va = 21.089832
     -3.4769986 + 20.141970i

ve = -0.92097057    0.29490584 - 0.38519280i
     -0.10091920   -0.18070330 - 0.38405816i
     0.061241324    0.24121182 - 0.56419722i
     0.36217049    0.017643612 + 0.26254313i
     0.081917964   -0.31466284 - 0.19936942i
```

Below we show that the first eigenvalue times the corresponding eigenvector (1) equals the input matrix times the first eigenvector (2).

(1) va[1]*ve[:,1]	=	(2) a*ve[:,1] =
-19.423115		-19.423115
-2.1283690		-2.1283690
1.2915693		1.2915693
7.6381149		7.6381149
1.7276361		1.7276361

Remarks

The ideal setting for input `ncv` is problem dependent and cannot be easily predicted ahead of time. Increasing `ncv` will increase the amount of memory used during computation. For a large, sparse matrix, `ncv` should be small compared to the order of input matrix `a`. **spEigv** is *not* thread-safe.

Technical Notes

spEigv implements functions from the ARPACK library.

spEye

spEye

Purpose

Creates a sparse identity matrix.

Format

```
y = spEye(n);
```

Input

n scalar, order of identity matrix.

Output

y $n \times n$ sparse identity matrix.

Remarks

Since sparse matrices are strongly typed in **GAUSS**, y must be defined as a sparse matrix before the call to **spEye**.

Example

```
//Declare 'y' a sparse matrix
sparse matrix y;

//Create 3x3 sparse identity matrix
y = spEye(3);
```

y is now equal to:

```
1  0  1
0  1  0
0  0  1
```

See Also

[spCreate](#), [spOnes](#), [denseToSp](#)

spGetNZE

Purpose

Returns the non-zero values in a sparse matrix, as well as their corresponding row and column indices.

Format

```
{ vals, rowinds, colinds } = spNumNZE(x);
```

Input

<i>x</i>	MxN sparse matrix.
----------	--------------------

Output

<i>vals</i>	Nx1 vector, non-zero values in <i>x</i> .
<i>rinds</i>	Nx1 vector, row indices of corresponding non-zero values.
<i>cinds</i>	Nx1 vector, column indices of corresponding non-zero values.

spGetNZE

Example

```
s  
  
sparse matrix y;  
x = { 0 0 0 10,  
      0 2 0  0,  
      0 0 0  0,  
      5 0 0  0,  
      0 0 0  3 };  
  
//Create sparse matrix from 'x'  
y = denseToSp(x,0);  
  
//Get non-zero values, row indices and column indices  
{ v,r,c } = spGetNZE(y);
```

v, the non-zero values, is equal to:

```
10  
2  
5  
3
```

r, the row indices, is equal to:

```
1  
2  
4  
5
```

c, the column indices, is equal to:

```
4  
2  
1  
4
```

See Also

[spNumNZE](#)

spline

Purpose

Computes a two-dimensional interpolatory spline.

Format

$\{ u, \quad v, \quad w \} = \textbf{spline}(x, y, \quad z, \textit{sigma}, g);$

Input

x	$K \times 1$ vector, x-abscissae (x-axis values).
y	$N \times 1$ vector, y-abscissae (y-axis values).
z	$K \times N$ matrix, ordinates (z-axis values).
\textit{sigma}	scalar, tension factor.
g	scalar, grid size factor.

Output

u	$(K * g) \times 1$ vector, x-abscissae, regularly spaced.
v	$(N * g) \times 1$ vector, y-abscissae, regularly spaced.
w	$(K * g) \times (N * g)$ matrix, interpolated ordinates.

Remarks

\textit{sigma} contains the tension factor. This value indicates the curviness desired. If

spLDL

σ is nearly zero (e.g., .001), the resulting surface is approximately the tensor product of cubic splines. If σ is large (e.g., 50.0), the resulting surface is approximately bi-linear. If σ equals zero, tensor products of cubic splines result. A standard value for σ is approximately 1.

g is the grid size factor. It determines the fineness of the output grid. For $g = 1$, the output matrices are identical to the input matrices. For $g = 2$, the output grid is twice as fine as the input grid, i.e., u will have twice as many columns as x , v will have twice as many rows as y , and w will have twice as many rows and columns as z .

Source

spline.src

spLDL

Purpose

Computes the LDL decomposition of a symmetric sparse matrix A.

Format

$\{ \textit{l}, \textit{d} \} = \text{spLDL}(a);$

Input

a N x N, symmetric sparse matrix.

Output

\textit{l} NxN lower-triangular sparse matrix.
 \textit{d} NxN diagonal sparse matrix.

Example

```
declare sparse matrix a, l, d;  
nz = { 142 13 56 57 0,  
      13  0  0  0 0,  
      56  0 94 47 0,  
      57  0 47 35 0,  
      0  0  0  0 0 };  
  
a = densetosp(nz,0);  
{ l, d } = spLDL(a);
```

Remarks

spLDL will not check to see if the input matrix is symmetric. The function looks only at the lower triangular portion of the input matrix.

See Also

[spLU](#)

Technical Notes

spLDL implements functions from the TAUCS library:

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spLU

Purpose

Computes the LU decomposition of a sparse matrix A with partial pivoting.

spLU

Format

```
{ l, u } = spLU(a);
```

Input

a N x N, non-singular sparse matrix.

Output

l NxN "scrambled" lower-triangular sparse matrix. This is a lower triangular matrix that has been reordered based upon the row pivoting.

u NxN "scrambled" upper-triangular sparse matrix. This is an upper triangular matrix that has been reordered based upon column pivoting to preserve sparsity.

Example

```
declare sparse matrix a, l, u;

nz = {-5.974 0 -13.37 6.136          0,
      0 5.932 7.712          0 -6.549,
      0 -5.728          0 14.227          0,
      0 -12.164 9.916 13.902 6.182,
      13.425          0 -12.654 -16.534          0 };

a = densetosp(nz,0);
{ l, u } = spLU(a);
```

Remarks

If the input matrix or either of the factors L and U are singular, the function will either

terminate the program with an error message or return an error code which can be tested for with the `scalerr` function. This depends on the trap state as follows:

trap 1	return error code: 50
trap 0	terminate with error message: Matrix singular

See Also

[spLDL](#)

Technical Notes

spLU implements functions from the SuperLU 4.0 library written by James W. Demmel, John R. Gilbert and Xiaoye S. Li.

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spNumNZE

Purpose

Returns the number of non-zero elements in a sparse matrix.

Format

$n = \text{spNumNZE}(x);$

Input

x	MxN sparse matrix.
-----	--------------------

spOnes

Output

n scalar, the number of non-zero elements in x .

Example

```
sparse matrix y;  
x = { 0 0 0 10,  
      0 2 0 0,  
      0 0 0 0,  
      5 0 0 0,  
      0 0 0 3 };  
  
y = denseToSp(x, 0);  
n = spNumNZE(y);  
print "The number of nonzeros is" n;
```

4.00

See Also

[spGetNZE](#)

spOnes

Purpose

Generates a sparse matrix containing only ones and zeros

Format

$y = \text{spOnes}(r, c, \text{rinds}, \text{cinds});$

Input

<i>r</i>	scalar, rows of output matrix.
<i>c</i>	scalar, columns of output matrix.
<i>rinds</i>	Nx1 vector, row indices of ones.
<i>cinds</i>	Nx1 vector, column indices of ones.

Output

<i>y</i>	$r \times c$ sparse matrix of ones.
----------	-------------------------------------

Remarks

Since sparse matrices are strongly typed in **GAUSS**, *y* must be defined as a sparse matrix before the call to **spOnes**.

Example

```
//declare sparse matrix
sparse matrix y;

//Set row indices and column indices
rinds = { 1, 3, 5 };
cinds = { 2, 1, 3 };

//Create a 5x4 sparse matrix with ones at the intersection
//of the 'rind' and 'cind'
y = spOnes(5,4,rinds,cinds);
```

The resulting *y* is equal to:

```
0  1  0  0
0  0  0  0
```

spreadSheetReadM

```
1  0  0  0
0  0  0  0
0  0  1  0
```

See Also

[spCreate](#), [spEye](#), [spZeros](#), [denseToSp](#)

spreadSheetReadM

Purpose

Reads and writes Excel files.

Format

```
x = spreadSheetReadM(file);
x = spreadSheetReadM(file, range);
x = spreadSheetReadM(file, range, sheet);
```

Input

<i>file</i>	string, name of .xls, or .xlsx file.
<i>range</i>	string, range to read or write; e.g., "A1:B20". Default = "A1".
<i>sheet</i>	scalar, sheet number. Default = 1.

Output

<i>x</i>	matrix of numbers read from Excel.
----------	------------------------------------

Examples

Example 1: Basic Example

Read all contents from the file `myfile.xlsx` located in your current GAUSS working directory.

```
x = spreadSheetReadM("myfile.xlsx");
```

Example 2: Read From a Range

```
x = spreadSheetReadM("myfile.xlsx", "B2:D110");
```

Example 3: Specify Path and Sheet Number

```
x = spreadSheetReadM("C:\\mydata\\myfile.xlsx", "A1", 1);
```

Portability

Windows, Linux and Mac

Remarks

1. If range is a null string, then by default the read will begin at cell "A1".
2. If `spreadSheetReadM` fails, it will either terminate and print an error message or return a scalar error code, which can be decoded with `scalerr`, depending on the state of the `trap` flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code.

```
//Will end the program and print an error message
x = spreadSheetReadM("nonexistent_file.xlsx");
```

S

spreadSheetReadSA

```
//Turn error trapping on
trap 1;
x = spreadSheetReadM("nonexistent_file.xlsx");

//Check to see if 'x' is a scalar error code
if scalmiss(x);
    //Code to handle error case here
endif;

//Turn error trapping off
trap 0;
```

See Also

[scalerr](#), [error](#), [SpreadsheetReadSA](#), [SpreadsheetWrite](#)

spreadSheetReadSA

Purpose

Reads and writes Excel files.

Format

```
sa = spreadSheetReadSA(file);
sa = spreadSheetReadSA(file, range);
sa = spreadSheetReadSA(file, range, sheet);
```

Input

<i>file</i>	string, name of .xls file.
<i>range</i>	string, range to read or write; e.g., "A1:B20". Default = "A1".

<i>sheet</i>	scalar, sheet number. Default = 1.
--------------	------------------------------------

Output

<i>sa</i>	string array read from Excel.
-----------	-------------------------------

Examples

Example 1: Basic Example

Read all contents from the file `myfile.xlsx` located in your current GAUSS working directory as a string array.

```
s = spreadSheetReadSA("myfile.xlsx");
```

Example 2: Read From a Range

```
s = spreadSheetReadSA("myfile.xlsx", "B2:D110");
```

Example 3: Specify Path and Sheet Number

```
s = spreadSheetReadSA("C:\\mydata\\myfile.xlsx", "A1", 1);
```

Portability

Windows, Linux and Mac

Remarks

1. If range is a null string, then by default the read will begin at cell "A1".
2. If `spreadSheetReadSA` fails, it will either terminate and print an error message or return a scalar error code, which can be decoded with `scalerr`, depending on the state of the `trap` flag.

spreadSheetWrite

trap 0	Print error message and terminate program.
trap 1	Return scalar error code.

```
//Will end the program and print an error message
x = spreadSheetReadSA("nonexistent_file.xlsx");
```

```
//Turn error trapping on
trap 1;
x = spreadSheetReadSA("nonexistent_file.xlsx");
```

```
//Check to see if 'x' is a scalar error code
if scalmiss(x);
    //Code to handle error case here
endif;
```

```
//Turn error trapping off
trap 0;
```

See Also

[scalerr](#), [error](#), [spreadSheetReadM](#), [spreadSheetWrite](#)

spreadSheetWrite

Purpose

Reads and writes Excel files.

Format

```
ret = spreadSheetWrite(data, file);
ret = spreadSheetWrite(data, file, range);
ret = spreadSheetWrite(data, file, range, sheet);
```

Input

<i>data</i>	matrix, string or string array, data to write.
<i>file</i>	string, name of .xls file.
<i>range</i>	string, range to read or write; e.g., "A1:B20". Default = "A1".
<i>sheet</i>	scalar, sheet number. Default = 1.

Output

<i>ret</i>	success code, 0 if successful, else error code.
------------	---

spreadSheetWrite

Examples

Example 1: Basic Example

```
x = { 0 1,
      1 2,
      3 5 };

//Write contents of 'x' to 'myfile.xlsx'
//from cell 'A1' to 'B3'
ret = spreadSheetWrite(x, "myfile.xlsx");
```

Example 2: Write To a Range

```
//Create a 1x4 string array of variable names
head = "Real GDP" $~ "Unemployment" $~ "CPI" $~ "PPI";

//Write the variable names to the cells 'C1:F1'
ret = spreadSheetWrite(head, "myfile.xlsx", "C1");
```

Example 3: Specify Path and Sheet Number

```
//Create a 10x3 matrix of Bernoulli random variables
x = rndBernoulli(10, 3, 0.6);

//Write the data from 'x' to cells 'B4:D13' on sheet 2 of
'myfile.xlsx'
ret = spreadSheetWrite(x, "C:\\mydata\\myfile.xlsx", "B4",
2);
```

Portability

Windows, Linux and Mac

Remarks

If **spreadSheetWrite** fails, it will either terminate and print an error message or

return a scalar error code, which can be detected with **scalmiss**, depending on the state of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code.

```
//If this fails, it will end the program and print an error  
message
```

```
x = spreadsheetWrite("myfile.xlsx");
```

```
//Turn error trapping on
```

```
trap 1;
```

```
x = spreadsheetWrite("myfile.xlsx");
```

```
//Check to see if 'x' is a scalar error code
```

```
if scalmiss(x);
```

```
    //Code to handle error case here
```

```
endif;
```

```
//Turn error trapping off
```

See Also

[scalerr](#), [error](#), [SpreadsheetReadM](#), [SpreadsheetReadSA](#)

spScale

Purpose

Scales a sparse matrix.

spScale

Format

`{ a, r, s } = spScale(x);`

Input

<code>x</code>	MxN sparse matrix.
----------------	--------------------

Output

<code>a</code>	MxN scaled sparse matrix.
<code>r</code>	Mx1 vector, row scale factors.
<code>s</code>	Nx1 vector, column scale factors.

Remarks

`spScale` scales the elements of the matrix by powers of 10 so that they are all within (-10,10).

Example

```
x = { 25  -12   0,
      3   0  -11,
      8 -100   0 };

declare sparse matrix sm, smsc;
sm = denseToSp(x,0);

{ smsc, r, c } = spScale(sm);
```

The results:

```
smc =
    2.50  -0.12  0.00
    0.30   0.00 -0.11
    0.80  -1.00  0.00

c =
    1.00
    0.10
    0.10

r =
    0.10
    0.10
    0.10
```

s

spSubmat

Purpose

Returns a sparse submatrix of a sparse matrix.

Format

`y = spSubmat(x, rinds, cinds);`

Input

<code>x</code>	MxN sparse matrix.
<code>rinds</code>	Kx1 vector, row indices.
<code>cinds</code>	Lx1 vector, column indices.

Output

<code>s</code>	KxL sparse matrix, the intersection of <code>rinds</code> and <code>cinds</code> .
----------------	--

spSubmat

Remarks

If *rinds* or *cinds* are scalar zeros, all rows or columns will be returned.

Since sparse matrices are strongly typed in **GAUSS**, *y* must be defined as a sparse matrix before the call to **spSubmat**.

Example

```
sparse matrix y;  
sparse matrix z;  
  
x = { 0 0 0 10,  
      0 2 0 0,  
      0 0 0 0,  
      5 0 0 0,  
      0 0 0 3 };  
  
y = denseToSp(x,0);  
  
//Extract all columns; rows 1, 3 and 4  
z = spSubmat(y,1|3|4,0);  
  
//Extract all values from 'z' into a dense matrix 'd'  
d = spDenseSubmat(z,0,0);
```

Now *d* is equal to:

0.00	0.00	0.00	10.00
0.00	0.00	0.00	0.00
5.00	0.00	0.00	0.00

See Also

[spDenseSubmat](#)

spToDense

Purpose

Converts a sparse matrix to a dense matrix.

Format

```
y = spToDense(x);
```

Input

<code>x</code>	MxN sparse matrix.
----------------	--------------------

Output

<code>y</code>	MxN dense matrix.
----------------	-------------------

Remarks

A dense matrix is just a normal format matrix.

Example

```
sparse matrix y;  
  
//Create a 4x4 sparse identity matrix  
y = spEye(4);  
  
//Create a dense matrix with the same values as 'y'  
d = spToDense(y);
```

The dense matrix `d` is equal to:

spTrTDense

```
1  0  0  0
0  1  0  0
0  0  1  0
0  0  0  1
```

See Also

[spDenseSubmat](#), [denseToSp](#)

spTrTDense

Purpose

Multiplies a sparse matrix transposed by a dense matrix.

Format

```
y = spTrTDense(s, d);
```

Input

s	NxM sparse matrix.
d	NxL dense matrix.

Output

y	MxL dense matrix, the result of $s' * d$.
-----	--

Remarks

This may also be accomplished by the following code:

```
y = s' * d;
```

However, **spTrTDense** will be more efficient.

See Also

[spTScalar](#)

spTScalar

Purpose

Multiplies a sparse matrix by a scalar.

Format

```
y = spTScalar(s, scal, rinds, cinds);
```

Input

<i>s</i>	NxM sparse matrix.
<i>scal</i>	scalar.
<i>rinds</i>	Kx1 vector of row indices.
<i>cinds</i>	Lx1 vector of column indices.

Output

<i>y</i>	KxL sparse matrix.
----------	--------------------

Remarks

Only the elements of *s* specified by *rinds* and *cinds* will be multiplied by *scal*. All other elements will be unchanged in the result.

To select all rows or all columns, input a scalar 0 for *rinds* or *cinds*.

spZeros

Since sparse matrices are strongly typed in **GAUSS**, *y* must be defined as a sparse matrix before the call to **spTScalar**.

Example

```
s  
  
sparse matrix y;  
x = { 3 0 2 1,  
      0 4 0 0,  
      5 0 0 3,  
      0 1 2 0 };  
  
rinds = 0;  
cinds = { 2,4 };  
  
//Multiply all elements in the second and fourth column  
//by 'scal'  
y = spTScalar(x,10,rinds,cinds);  
d = spDenseSubmat(y,0,0);
```

The result, in *d* is:

```
3 0 2 1  
0 40 0 0  
5 0 0 3  
0 10 2 0
```

See Also

[spTrTDense](#)

spZeros

Purpose

Creates a sparse matrix containing no non-zero values.

Format

```
y = spZeros(r, c);
```

Input

<i>r</i>	scalar, rows of output matrix.
<i>c</i>	scalar, columns of output matrix.

Output

<i>y</i>	<i>r</i> x <i>c</i> sparse matrix.
----------	------------------------------------

Remarks

Since sparse matrices are strongly typed in **GAUSS**, *y* must be defined as a sparse matrix before the call to **spZeros**.

Example

```
sparse matrix y;  
  
//Create a 4x3 sparse matrix with all elements set to 0  
y = spZeros(4,3);  
  
//Create a dense matrix with the same values as 'y'  
d = spToDense(y);
```

The contents of *d* are equal to:

```
0 0 0  
0 0 0  
0 0 0  
0 0 0
```

sqpSolve

See Also

[spOnes](#), [spEye](#)

sqpSolve

Purpose

Solves the nonlinear programming problem using a sequential quadratic programming method.

Format

```
{ x, f, lagr, retcode } = sqpSolve(&fct, start);
```

Input

<i>&fct</i>	pointer to a procedure that computes the function to be minimized. This procedure must have one input argument, a vector of parameter values, and one output argument, the value of the function evaluated at the input vector of parameter values.
<i>start</i>	Kx1 vector of start values.

Global Input

<i>_sqp_A</i>	MxK matrix, linear equality constraint coefficients.
<i>_sqp_B</i>	Mx1 vector, linear equality constraint constants.
	These globals are used to specify linear equality constraints of the following type:
	$_sqp_A * x = _sqp_B$

`_sqp_EqProc`

where x is the $K \times 1$ unknown parameter vector.
scalar, pointer to a procedure that computes the
nonlinear equality constraints. For example, the
statement:

```
_sqp_EqProc = &eqproc;
```

tells **sqpSolve** that nonlinear equality constraints are
to be placed on the parameters and where the
procedure computing them is to be found. The
procedure must have one input argument, the $K \times 1$
vector of parameters, and one output argument, the
 $R \times 1$ vector of computed constraints that are to be equal
to zero. For example, suppose that you wish to place
the following constraint:

```
p[1] * p[2] = p[3]
```

The procedure for this is:

```
proc eqproc(p);  
  retp(p[1]*p[2]-p[3]);  
endp;
```

`_sqp_C`

$M \times K$ matrix, linear inequality constraint
coefficients.

`_sqp_D`

$M \times 1$ vector, linear inequality constraint constants.

These globals are used to specify linear inequality
constraints of the following type:

```
_sqp_C * X >= _sqp_D
```

where x is the $K \times 1$ unknown parameter vector.
scalar, pointer to a procedure that computes the

`_sqp_`

sqpSolve

IneqProc

nonlinear inequality constraints. For example the statement:

```
_sqp_EqProc = &ineqproc;
```

tells **sqpSolve** that nonlinear equality constraints are to be placed on the parameters and where the procedure computing them is to be found. The procedure must have one input argument, the Kx1 vector of parameters, and one output argument, the Rx1 vector of computed constraints that are to be equal to zero. For example, suppose that you wish to place the following constraint:

```
p[1] * p[2] >= p[3]
```

The procedure for this is:

```
proc ineqproc(p);  
    retp(p[1]*[2]-p[3]);  
endp;
```

_sqp_Bounds

Kx2 matrix, bounds on parameters. The first column contains the lower bounds, and the second column the upper bounds. If the bounds for all the coefficients are the same, a 1x2 matrix may be used. Default is:

```
[1] -1e256 [2] 1e256
```

*_sqp_
GradProc*

scalar, pointer to a procedure that computes the gradient of the function with respect to the parameters. For example, the statement:

```
_sqp_GradProc = &gradproc;
```


*_sqp_
HessProc*

tells **sqpSolve** that a gradient procedure exists and where to find it. The user-provided procedure has two input arguments, a Kx1 vector of parameter values and an NxP matrix of data. The procedure returns a single output argument, an NxK matrix of gradients of the log-likelihood function with respect to the parameters evaluated at the vector of parameter values.

Default = 0, i.e., no gradient procedure has been provided.

scalar, pointer to a procedure that computes the Hessian, i.e., the matrix of second order partial derivatives of the function with respect to the parameters. For example, the instruction:

```
_sqp_HessProc = &hessproc;
```

will tell **sqpSolve** that a procedure has been provided for the computation of the Hessian and where to find it. The procedure that is provided by the user must have two input arguments, a Px1 vector of parameter values and an NxK data matrix. The procedure returns a single output argument, the PxP symmetric matrix of second order derivatives of the function evaluated at the parameter values.

*_sqp_
MaxIters*

scalar, maximum number of iterations. Default = 1e+5. Termination can be forced by pressing C on the keyboard.

_sqp_DirTol

scalar, convergence tolerance for gradient of estimated coefficients. Default = 1e-5. When this criterion has been satisfied, **sqpSolve** will exit the iterations.

*_sqp_
ParamNames*

Kx1 character vector, parameter names.

sqpSolve

s

<i>ParNames</i>	
<i>_sqp_ PrintIters</i>	scalar, if nonzero, prints iteration information. Default = 0. Can be toggled during iterations by pressing P on the keyboard.
<i>_sqp_ FeasibleTest</i>	scalar, if nonzero, parameters are tested for feasibility before computing function in line search. If function is defined outside inequality boundaries, then this test can be turned off.
<i>_sqp_ RandRadius</i>	scalar, if zero, no random search is attempted. If nonzero it is the radius of random search which is invoked whenever the usual line search fails. Default = .01.
<i>__output</i>	scalar, if nonzero, results are printed. Default = 0.

Output

<i>x</i>	Kx1 vector of parameters at minimum.				
<i>f</i>	scalar, function evaluated at <i>x</i> .				
<i>lagr</i>	vector, created using vput . Contains the Lagrangean for the constraints. They may be extracted with the vread command using the following strings: <table><tr><td><i>"lineq"</i></td><td>Lagrangeans of linear equality constraints,</td></tr><tr><td><i>"nlineq"</i></td><td>Lagrangeans of nonlinear equality</td></tr></table>	<i>"lineq"</i>	Lagrangeans of linear equality constraints,	<i>"nlineq"</i>	Lagrangeans of nonlinear equality
<i>"lineq"</i>	Lagrangeans of linear equality constraints,				
<i>"nlineq"</i>	Lagrangeans of nonlinear equality				

	constraints
"linineq"	Lagrangeans of linear inequality constraints
"nlinineq"	Lagrangeans of nonlinear inequality constraints
"bounds"	Lagrangeans of bounds

Whenever a constraint is active, its associated Lagrangean will be nonzero.

<i>retcode</i>	return code:
0	normal convergence
1	forced exit
2	maximum number of iterations exceeded
3	function calculation failed
4	gradient calculation failed
5	Hessian calculation failed
6	line search failed
7	error with

5

sqpSolve

constraints

Remarks

Pressing C on the keyboard will terminate iterations, and pressing P will toggle iteration output.

sqpSolve is recursive, that is, it can call itself with another function and set of global variables,

Example

```
//Reset all sqpSolve global variables

sqpSolveSet;

proc fct(x);
    retp( (x[1] + 3*x[2] + x[3])^2 + 4*(x[1] - x[2])^2);
endp;

proc ineqp(x);
    retp(6*x[2] + 4*x[3] - x[1]^3 - 3);
endp;

proc eqp(x);
    retp(1-sumc(x));
endp;

_sqp_Bounds = { 0 1e256 };

start = { .1, .7, .2 };

_sqp_IneqProc = &ineqp;
_sqp_EqProc = &eqp;
```

```
{ x,f,lagr,ret } = sqpSolve(&fct,start);
```

Source

sqpsolve.src

sqpSolveMT

Purpose

Solves the nonlinear programming problem.

Format

```
out = sqpSolveMT(&fct, par1);
out = sqpSolveMT(&fct, par1,...);
out = sqpSolveMT(&fct, par1,...,ctl);
out = sqpSolveMT(&fct, par1,ctl);
```

Input

<i>&fct</i>	pointer to a procedure that computes the function to be minimized. The first input to this procedure must be an instance of structure of type PV .
<i>par1</i>	an instance of structure of type PV . The <i>par1</i> instance is passed to the user-provided procedure pointed to by <i>&fct</i> . <i>par1</i> is constructed using the " pack " functions.
<i>...</i>	Optional extra arguments. These arguments are passed untouched to the user-provided objective function, by sqpSolveMT .
<i>ctl</i>	an instance of an sqpSolveMTControl structure.

Normally an instance is initialized by calling **sqpSolveMTControlCreate** and members of this instance can be set to other values by the user. For an instance named `ctl`, the members are:

<code>ctl.A</code>	MxK matrix, linear equality constraint coefficients: $ctl.A * p = ctl.B$ where p is a vector of the parameters.
<code>ctl.B</code>	Mx1 vector, linear equality constraint constants: $ctl.A * p = ctl.B$ where p is a vector of the parameters.
<code>ctl.C</code>	MxK matrix, linear inequality constraint coefficients: $ctl.C * p \geq ctl.D$ where p is a vector of the parameters.
<code>ctl.D</code>	Mx1 vector, linear inequality constraint constants: $ctl.C * p \geq ctl.D$ where p is a vector of the parameters.
<code>ctl.eqProc</code>	scalar, pointer to a procedure that computes the nonlinear equality constraints. When such a procedure has been provided, it has one input argument, a structure of type SQPdata, and one output argument, a vector of computed equality constraints. For more details see Remarks below. Default = ., i.e., no

<i>ctl.weights</i>	equality procedure. vector, weights for objective function returning a vector. Default = 1.
<i>ctl.ineqProc</i>	scalar, pointer to a procedure that computes the nonlinear inequality constraints. When such a procedure has been provided, it has one input argument, a structure of type SQPdata, and one output argument, a vector of computed inequality constraints. For more details see Remarks below. Default = ., i.e., no inequality procedure.
<i>ctl.bounds</i>	1x2 or Kx2 matrix, bounds on parameters. If 1x2 all parameters have same bounds. Default = -1e256 1e256 .
<i>ctl.covType</i>	scalar, if 2, QML covariance matrix, else if 0, no covariance matrix is computed, else ML covariance matrix is computed.
<i>ctl.gradProc</i>	scalar, pointer to a procedure that computes the gradient of the function with respect to the parameters. Default = ., i.e., no gradient procedure has been provided.
<i>ctl.hessProc</i>	scalar, pointer to a procedure that

ctl.maxIters

computes the Hessian, i.e., the matrix of second order partial derivatives of the function with respect to the parameters. Default = ., i.e., no Hessian procedure has been provided.

scalar, maximum number of iterations. Default = 1e+5.

ctl.dirTol

scalar, convergence tolerance for gradient of estimated coefficients. Default = 1e-5. When this criterion has been satisfied SQPSolve exits the iterations.

ctl.feasibleTest

scalar, if nonzero, parameters are tested for feasibility before computing function in line search. If function is defined outside inequality boundaries, then this test can be turned off. Default = 1.

ctl.randRadius

scalar, If zero, no random search is attempted. If nonzero, it is the radius of random search which is invoked whenever the usual line search fails. Default = .01.

ctl.output

scalar, if nonzero, results are printed. Default = 0.

ctl.printIters

scalar, if nonzero, prints iteration information. Default = 0.

Output

<i>out</i>	an instance of an sqpSolveMTout structure. For an instance named <i>out</i> , the members are:
<i>outx.par</i>	an instance of structure of type PV containing the parameter estimates will be placed in the member matrix <i>out.par</i> .
<i>out.fct</i>	scalar, function evaluated at x.
<i>out.lagr</i>	an instance of a SQPLagrange structure containing the Lagrangeans for the constraints. The members are:
<i>out.lagr.lineq</i>	Mx1 vector, Lagrangeans of linear equality constraints.
<i>out.lagr.nlineq</i>	Nx1 vector, Lagrangeans of nonlinear equality constraints.
<i>out.lagr.linineq</i>	Px1 vector, Lagrangeans of linear inequality constraints.
<i>out.lagr.nlinineq</i>	Qx1 vector, Lagrangeans of nonlinear inequality constraints.
<i>out.lagr.bounds</i>	Kx2 matrix, Lagrangeans of bounds.

S

Whenever a constraint is active, its associated Lagrangean will be nonzero. For any constraint that is inactive throughout the iterations as well as at convergence, the corresponding Lagrangean matrix will be set to a scalar missing value.

return code:

<i>out.retcode</i>	
0	normal convergence.
1	forced exit.
2	maximum number of iterations exceeded.
3	function calculation failed.
4	gradient calculation failed.
5	Hessian calculation failed.
6	line search failed.
7	error with constraints.
8	function complex.

Remarks

There is one required user-provided procedure, the one computing the objective function to be minimized, and four other optional functions, one each for computing the equality constraints, the inequality constraints, the gradient of the objective function, and the Hessian of the objective function.

All of these functions must take exactly the same input arguments. The first input argument is an instance of a structure of type **PV**. On input to the call to **sqpSolveMT**, this **PV** structure contains starting values for the parameters.

Both of the structures of type **PV** are set up using the **PV "pack"** procedures, **pvPack**, **pvPackm**, **pvPacks**, and **pvPacksm**. These procedures allow for setting up a parameter vector in a variety of ways.

For example, we might have the following objective function for fitting a nonlinear curve to data:

```
proc (1) = micherlitz(struct PV par1, y, x);
    local p0,e,s2,x,y;
    p0 = pvUnpack(par1, "parameters");
    e = y - p0[1] - p0[2]*exp(-p0[3] * x);
    retp(e'*e);
endp;
```

In this example the dependent and independent variables are passed to the procedure as the second and third arguments to the procedure.

The other optional procedures must take exactly the same arguments as the objective function. For example, to constrain the squared sum of the first two parameters to be greater than one in the above problem, provide the following procedure:

```
proc (1) = ineqConst(struct PV par1, y, x);
    local p0;
    p0 = pvUnpack(p0, "parameters");
    retp( (p0[2]+p0[1])^2 - 1);
endp;
```

The following is a complete example for estimating the parameters of the Micherlitz equation in data with bounds constraints on the parameters and where an optional gradient procedure has been provided:

sqpSolveMT

S

```
//Create data needed by 'Micherlitz' procedure
y = { 3.183,
      3.059,
      2.871,
      2.622,
      2.541,
      2.184,
      2.110,
      2.075,
      2.018,
      1.903,
      1.770,
      1.762,
      1.550 };

x = seqa(1,1,13);

//Declare control structure
struct sqpSolveMTControl c0;

//Initialize structure to default values
c0 = sqpSolveMTControlCreate();

//Constrain parameters to be positive
c0.bounds = 0~100;

//Declare 'par1' to be a PV structure
struct PV par1;

//Initialize 'par1'
par1 = pvCreate();

//Add 3x1 vector named 'parameters' to 'p1'
par1 = pvPack(par1,.92|2.62|.114, "parameters");
```

```

//Declare 'out' to be an sqpsolvemt control structure
//to hold the results from sqpsolvemt
struct sqpSolveMTout out;

//Estimate the model parameters
out = sqpSolveMT(&Micherlitz,par1,y,x,c0);

//Print returned parameter estimates
print "parameter estimates ";
print pvUnPack(out.par, "parameters");

proc Micherlitz(struct PV par1, y, x);
    local p0,e,s2;
    p0 = pvUnpack(par1, "parameters");
    e = y - p0[1] - p0[2]*exp(-p0[3] * x);
    retp(e'*e);
endp;

```

Source

sqpsolvemt.src

See Also

[sqpSolveMTControlCreate](#), [sqpSolveMTlagrangeCreate](#), [CR-sqpSolveMToutCreate](#)

sqpSolveMTControlCreate

Purpose

Creates an instance of a structure of type **sqpSolveMTcontrol** set to default values.

Include

sqpsolvemt.sdf

sqpSolveMTlagrangeCreate

Format

```
s = sqpSolveMTControlCreate();
```

Output

s instance of structure of type **sqpSolveMTControl**.

Example

```
//Declare instance of structure
struct sqpSolveMTControl s;

//Initialize the structure to default values
s = sqpSolveMTControlCreate();
```

Source

sqpsolvemt.src

See Also

[sqpSolve](#)

sqpSolveMTlagrangeCreate

Purpose

Creates an instance of a structure of type **sqpSolveMTlagrange** set to default values.

Include

sqpsolvemt.sdf

Format

```
s = sqpSolveMTlagrangeCreate();
```

Output

```
s                    instance of structure of type sqpSolveMTlagrange.
```

Example

```
//Declare instance of structure  
struct sqpSolveMTlagrange sla;  
  
//Initialize the structure to default values  
sla = sqpSolveMTlagrangeCreate ();
```

Source

sqpsolvemt.src

See Also

[sqpSolve](#)

sqpSolveMToutCreate**Purpose**

Creates an instance of a structure of type **sqpSolveMTout** set to default values.

Include

sqpsolvemt.sdf

sqpSolveSet

Format

```
s = sqpSolveMToutCreate();
```

Output

s instance of structure of type **sqpSolveMTout**.

Example

```
//Declare instance of structure
struct sqpSolveMTout out;

//Initialize the structure to default values
out = sqpSolveMToutCreate();
```

Source

sqpsolvemt.src

See Also

[sqpSolve](#)

sqpSolveSet

Purpose

Resets global variables used by **sqpSolve** to default values.

Format

```
sqpSolveSet;
```


Source

sqpsolve.src

sqrt

Purpose

Computes the square root of every element in x .

Format

$y = \text{sqrt}(x);$

Input

x	NxK matrix or N-dimensional array.
-----	------------------------------------

Output

y	NxK matrix or N-dimensional array, the square roots of each element of x .
-----	--

Remarks

If x is negative, complex results are returned by default. You can turn the generation of complex numbers for negative inputs on or off in the **GAUSS** configuration file, and with the **sysstate** function, case 8. If you turn it off, **sqrt** will generate an error for negative inputs.

If x is already complex, the complex number state does not matter; **sqrt** will compute a complex result.

stdc

Example

```
let x[2,2] = 1 2 3 4;  
y = stdc(x);
```

The output, in variable y is equal to:

```
1.00000000  
1.41421356  
1.73205081  
2.00000000
```

stdc

Purpose

Computes the standard deviation of the elements in each column of a matrix.

Format

```
 $y$  = stdc( $x$ );
```

Input

x	$N \times K$ matrix.
-----	----------------------

Output

y	$K \times 1$ vector, the standard deviation of each column of x .
-----	---

Remarks

This function essentially computes sample standard deviation, s :

$$s = \sqrt{\frac{1}{n-1} \times \sum_{i=1}^n (X_i - \bar{X})^2}$$

Thus, the divisor is N-1 rather than N, where N is the number of elements being summed.

To convert to the population's standard deviation, multiply by $\sqrt{\frac{n-1}{n}}$:

$$\sigma = s \times \sqrt{\frac{n-1}{n}}$$

Example

```
//Set the rng seed so that the random numbers produced will
//be repeatable

rndseed 94243524;

//Create a vector of random normal numbers
y = rndn(8100,1);

//Compute the standard deviation of the column vector 'y'
std = stdc(y);
```

The standard deviation, in variable `std`, is equal to:

```
1.00183907
```

See Also

[meanc](#)

stdsc

stdsc

Purpose

Computes the standard deviation of the elements in each column of a matrix.

Format

$y = \text{stdsc}(x);$

Input

x NxK matrix.

Output

y Kx1 vector, the standard deviation of each column of x .

Remarks

This function essentially computes:

$$\text{sqrt} (1 / (N) * \text{sumc} ((x - \text{meanc} (x)) ^ 2))$$

Thus, the divisor is N rather than N-1, where N is the number of elements being summed. See **stdc** for the alternate definition.

Example

```
//Create 3 columns of random normal numbers
y = rndn(8100,3);

//Calculate the standard deviation of each column
```

```
std = stdsc(y);
```

The return, in variable *std* is equal to:

```
1.00095980  
0.99488832  
1.00201375
```

See Also

[stdc](#), [astds](#), [meanc](#)

stocv

Purpose

Converts a string to a character vector.

Format

```
v = stocv(s);
```

Input

s string, to be converted to character vector.

Output

v Nx1 character vector, contains the contents of *s*.

Remarks

stocv breaks *s* up into a vector of 8-character length matrix elements. Note that the

stof

character information in the vector is not guaranteed to be null-terminated.

Example

s

```
s = "Now is the time for all good men";  
v = stof(s);
```

```
    "Now is t"  
  
    "the time "  
v =  
    "for all "  
  
    "good men"
```

See Also

[cvtos](#), [vget](#), [vlist](#), [vput](#), [vread](#)

stof

Purpose

Converts a string to floating point.

Format

```
y = stof(x);
```

Input

x	string or NxK matrix containing character elements to be converted.
---	---

Output

y	matrix, the floating point equivalents of the ASCII numbers in x .
-----	--

Remarks

If x is a string containing "1 2 3", then **stof** will return a 3x1 matrix containing the numbers 1, 2 and 3.

If x is a null string, **stof** will return a 0.

This uses the same input conversion routine as [loadm](#) and [let](#). It will convert character elements and missing values. **stof** also converts complex numbers in the same manner as [let](#).

See Also

[ftos](#), [ftocv](#), [chrs](#)

stop

Purpose

Stops a program and returns to the command prompt. Does not close files.

Format

```
stop;
```

Remarks

This command has the same effect as [end](#), except it does not close files or the auxiliary output.

strcombine

It is not necessary to put a `stop` or an `end` statement at the end of a program. If neither is found, an implicit `stop` is executed.

See Also

[end](#), [new](#), [system](#)

strcombine

Purpose

Converts an NxM string array to an Nx1 string vector by combining each element in a column separated by a user-defined delimiter string.

Format

```
y = strcombine(sa, delim, qchar);
```

Input

<i>sa</i>	NxM string array.
<i>delim</i>	1x1, 1xM, or Mx1 delimiter string.
<i>qchar</i>	scalar, 2x1, or 1x2 string vector containing quote characters as required: scalar: Use this character as quote character. If this is 0, no quotes are added. 2x1 or 1x2 Contains left and right quote string characters. vector:

Output

y Nx1 string vector result.

Examples

Example 1: Basic example

```
//Create 1x3 string array
sa_dir = "C:" $~ "gauss" $~ "myProject";

//Combine 1x3 string array with '/' at end of each element
path = strcombine(sa_dir, "/", 0);
```

After the above code, *path* is equal to:

```
"C:/gauss/myProject/"
```

Remarks

Note that **strcombine** adds a delimiter after the final element. To combine strings with the delimiter added only between tokens, see **strjoin**.

Source

strfns.src

See Also

[satostrC](#), [strjoin](#)

strindx

strindx

Purpose

Finds the index of one string within another string.

Format

y = **strindx**(*where*, *what*, *start*);

Input

<i>where</i>	string or scalar, the data to be searched.
<i>what</i>	string or scalar, the substring to be searched for in <i>where</i> .
<i>start</i>	scalar, the starting point of the search in <i>where</i> for an occurrence of <i>what</i> . The index of the first character in a string is 1.

Output

<i>y</i>	scalar containing the index of the first occurrence of <i>what</i> , within <i>where</i> , which is greater than or equal to <i>start</i> . If no occurrence is found, it will be 0.
----------	--

Remarks

An example of the use of this function is the location of a name within a string of names:

```
z = "nameagepaysex";  
x = "pay";  
y = strindx(z,x,1);
```

The above code will set *y* equal to:

```
8.00
```

This function is used with **strsect** for extracting substrings.

See Also

[strindx](#), [strlen](#), [strsect](#), [strput](#)

strlen

Purpose

Returns the length of a string.

Format

```
y = strlen(x);
```

Input

<i>x</i>	string, NxK matrix of character data, or NxK string array.
----------	--

Output

<i>y</i>	scalar containing the exact length of the string <i>x</i> , or NxK matrix or string array containing the lengths of the elements in <i>x</i> .
----------	--

strput

Remarks

The null character (ASCII 0) is a legal character within strings and so embedded nulls will be counted in the length of strings. The final terminating null byte is not counted, though.

For character matrices, the length is computed by counting the characters (maximum of 8) up to the first null in each element of the matrix. The null character, therefore, is not a valid character in matrices containing character data and is not counted in the lengths of the elements of those matrices.

Example

```
x1 = "How long?";  
x2 = "Classification";  
len1 = strlen(x1);  
len2 = strlen(x2);
```

After running the code above:

```
len1 = 9  
  
len2 = 14
```

See Also

[strsect](#), [strindx](#), [strrindx](#)

strput

Purpose

Lays a substring over a string.

Format

```
y = strput(substr, str, off);
```

Input

<i>substr</i>	string, the substring to be laid over the other string.
<i>str</i>	string, the string to receive the substring.
<i>off</i>	scalar, the offset in <i>str</i> to place <i>substr</i> . The offset of the first byte is 1.

Output

<i>y</i>	string, the new string.
----------	-------------------------

Example

```
str = "max";  
sub = "imum";  
loc = 4;  
y = strput(sub, str, loc);  
print y;
```

produces:

```
maximum
```

Source

strput.src

strrindx

strrindx

Purpose

Finds the index of one string within another string. Searches from the end of the string to the beginning.

Format

`y=`**strrindx**(*where*, *what*, *start*);

Input

<i>where</i>	string or scalar, the data to be searched.
<i>what</i>	string or scalar, the substring to be searched for in <i>where</i> .
<i>start</i>	scalar, the starting point of the search in <i>where</i> for an occurrence of <i>what</i> . <i>where</i> will be searched from this point backward for <i>what</i> .

Output

<i>y</i>	scalar containing the index of the last occurrence of <i>what</i> , within <i>where</i> , which is less than or equal to <i>start</i> . If no occurrence is found, it will be 0.
----------	--

Remarks

A negative value for *start* causes the search to begin at the end of the string. An example of the use of **strrindx** is extracting a file name from a complete path specification:

```
path = "/gauss/src/ols.src";
ps = "/";
pos = strrindx(path,ps,-1);
if pos;
    name = strsect(path,pos+1,strlen(path)-pos);
else;
    name = "";
endif;
```

The above code makes the following assignments:

```
pos = 11

name = ols.src
```

See Also

[strindx](#), [strlen](#), [strsect](#), [strput](#)

strjoin

Purpose

Converts an NxM string array to an Nx1 string vector by combining each element in a column separated by a user-defined delimiter string.

Format

```
y = strjoin(sa, delim);
y = strjoin(sa, delim, qchar);
```

Input

<i>sa</i>	NxM string array.
-----------	-------------------

strjoin

<i>delim</i>	1x1, 1xM, or Mx1 delimiter string.	
<i>qchar</i>	Optional input, scalar, 2x1, or 1x2 string vector containing quote characters as required:	
	scalar:	Use this character as quote character. If this is 0, no quotes are added.
	2x1 or 1x2 string vector:	Contains left and right quote characters.
	Default value is 0 (no quotes).	

Output

<i>y</i>	Nx1 string vector result.
----------	---------------------------

Example

Example 1

```
//Create a 1x4 string array
s = "alpha" $~ "beta" $~ "gamma" $~ "delta";

//Combine the string array into a single comma-separated string
varnames = strjoin(s, ",");
```

After the above code, *varnames* will be a single string with the following contents:

```
"alpha,beta,gamma,delta"
```

Example 2


```
//Create 1x3 string array
s = "GDP" $~ "Gross Exports" $~ "Net Exports";

//Create single string separated by spaces
//with each element surrounded by a single tic '
names = strjoin(s, " ", "'");
```

After the above code, *names* should be equal to the string:

```
"'GDP' 'Gross Exports' 'Net Exports'"
```

Remarks

- **strjoin** differs from **strcombine** by not adding a delimiter after the last element.
- In the case where the input has only 1 column, the delimiter is ignored.

Source

strfns.src

See Also

[strcombine](#)

strsect

Purpose

Extracts a substring of a string.

Format

```
y = strsect(str, start, len);
```

strsect

Input

<i>str</i>	string or scalar from which the segment is to be obtained.
<i>start</i>	scalar, the index of the substring in <i>str</i> . The index of the first character is 1.
<i>len</i>	scalar, the length of the substring.

Output

<i>y</i>	string, the extracted substring, or a null string if <i>start</i> is greater than the length of <i>str</i> .
----------	--

Remarks

If there are not enough characters in a string for the defined substring to be extracted, then a short string or a null string will be returned.

If *str* is a matrix containing character data, it must be scalar.

Example

```
strng = "This is an example string.";
y = strsect(strng,12,7);
```

The above code assigns the variable *y* to be:

example

See Also

[strlen](#), [strindx](#), [strindx](#)

strsplit

Purpose

Splits a string into individual tokens.

Format

```
sa = strsplit(str);  
sa = strsplit(str, sep);
```

Input

str	String or Nx1 string array to be split.
sep	Optional argument, string containing the character used to separate the input string into individual tokens.

Output

sa	1xK or NxK string array.
----	--------------------------

Remarks

Case 1: No supplied separator

If **strsplit** is called with only one input (i.e. a separator is not passed in as the second argument), each of the following characters are considered delimiters:

space	ASCII 32
tab	ASCII 9
comma	ASCII 44
newline	ASCII 10
carriage return	ASCII 13

strsplit

The input string will be split at each occurrence of ANY of the separators listed in the table above. For example:

```
sa = "alpha 1,beta 2,gamma 3";  
strsplit(s);
```

will return a 1x6 string array with the following contents:

```
"alpha"    "1"    "beta"    "2"    "gamma"    "3"
```

Tokens containing delimiters must be enclosed in single or double quotes or parentheses. Tokens enclosed in single or double quotes will NOT retain the quotes upon translation. Tokens enclosed in parentheses WILL retain the parentheses after translation. Parentheses cannot be nested.

Case 2: Supplied separator

If a separator is passed to **strsplit**, the input string will be split into individual tokens at each instance of the specified separator. Only the supplied separator will be used to separate the tokens. Separators may only be 1 character. Any remaining white-space will be preserved. For example:

```
strsplit("alpha 1,beta 2,gamma 3", ",");
```

will return a 1x3 string array with the following contents:

```
"alpha 1"    "beta 2"    "gamma 3"
```

Rows with fewer tokens will be padded on the right. For example:

```
string s = { "1982-04-19", "1994-06" };  
strsplit(s, "-");
```

will return:

```
"1982"    "04"    "19"  
"1994"    "06"    ""
```

Examples

Example 1: Dates

```
dt = "1977/04/03";  
dt_split = strsplit(dt, "/");
```

After the code above, *dt_split* will be a 1x3 string array with the following contents:

```
"1977"    "04"    "03"
```

Example 2: Comma-separated list of variables

```
vars = "CPI,PPI,Employment,Oil:Brent blend,Oil:WTI";  
vars = strsplit(vars, ",");
```

After the code above, *vars* will be a 1x5 string array with the following contents:

```
"CPI"      "PPI"      "Employment"    "Oil:Brent blend"  
"Oil:WTI"
```

Example 3: String array with supplied separator

```
//Create a 3x1 string array  
string dow_str = { "apple:technology",  
                  "goldman sachs:finance",  
                  "home depot:retail" };  
  
//Split 'dow_str' into a 3x2 string array  
dow_sa = strsplit(dow_str, ":");
```

The above code sets *dow_sa* to be equal to:

strsplit

```
"apple"           "technology"
"goldman sachs"   "finance"
"home depot"      "retail"
```

Example 4: String array without supplied separator

Elements that contain spaces may be grouped with single ticks, like this:

```
ss = "classification 'scientific taxonomy'";
ss2 = strsplit(ss);

print "ss2[1] = " ss2[1];
print "ss2[2] = " ss2[2];
```

In this program, 'scientific taxonomy' is kept as one token, and thus the output from the above code is:

```
ss2[1] = classification
ss2[2] = scientific taxonomy
```

Example 5: String array with multi-character delimiter

```
ss = "h5://example.h5";
ss2 = strsplit(ss, "://");

print "ss2[1] = " ss2[1];
print "ss2[2] = " ss2[2];
```

The output from the above code is:

```
ss2[1] = h5
ss2[2] = example.h5
```

See Also

[strsplitPad](#)

strsplitPad

Purpose

Splits a string vector into a string array of the individual tokens. Pads on the right with null strings.

Format

```
sa = strsplitPad(sv, n_cols);
```

Input

sv	Nx1 string array.
n_cols	scalar, number of columns of output string array.

Output

sa	Nx n_cols string array.
----	-------------------------

Remarks

Rows containing more than n_cols tokens are truncated and rows containing fewer than n_cols tokens are padded on the right with null strings. The following characters are considered delimiters between tokens:

space	ASCII 32
tab	ASCII 9
comma	ASCII 44
newline	ASCII 10
carriage return	ASCII 13

strtodt

Tokens containing delimiters must be enclosed in single or double quotes or parentheses. Tokens enclosed in single or double quotes will NOT retain the quotes upon translation. Tokens enclosed in parentheses WILL retain the parentheses after translation. Parentheses cannot be nested.

Example

```
string sv = {  
    "alpha beta gamma",  
    "delta, epsilon, zeta, eta",  
    "theta iota kappa"  
};  
  
sa = strsplitPad(sv, 4);
```

After the code above, *sa* will be equal to:

```
"alpha"    "beta" "gamma"    ""  
"delta" "epsilon" "zeta" "eta"  
"theta"    "iota" "kappa"    ""
```

See Also

[strsplit](#)

strtodt

Purpose

Converts a string array of dates to a matrix in DT scalar format.

Format

```
x = strtodt(sa, fmt);
```


Input

<i>sa</i>	NxK string array containing dates.
<i>fmt</i>	string containing date/time format characters.

Output

<i>x</i>	NxK matrix of dates in DT scalar format.
----------	--

Remarks

The DT scalar format is a double precision representation of the date and time. In the DT scalar format, the number:

```
20120921223505
```

represents 22:35:05 or 10:35:05 PM on September 21, 2012.

The following formats are supported:

YYYY	Four digit year
YR	Last two digits of year
MO	Number of month, 01-12
DD	Day of month, 01-31
HH	Hour of day, 00-23
MI	Minute of hour, 00-59
SS	Second of minute, 00-59

Example

```
x = strtodt("2012-07-12 10:18:32", "YYYY-MO-DD HH:MI:SS");
print x;
```

strtof

produces:

20120712101832.0

```
x = strtodt("2012-07-12 10:18:32", "YYYY-MO-DD");  
print x;
```

produces:

20120712000000.0

```
x = strtodt("10:18:32", "HH:MI:SS");  
print x;
```

produces:

101832.0

```
x = strtodt("05-28-10", "MO-DD-YR");  
print x;
```

produces:

20100528000000.0

See Also

[dttostr](#), [dttoutc](#), [utctodt](#)

strtof

Purpose

Converts a string array to a numeric matrix.

Format

```
 $x = \text{strtof}(sa);$ 
```

Input

Input	Description
sa	$N \times K$ string array containing numeric data.

Output

Output	Description
x	$N \times K$ matrix.

Remarks

Elements with more than one numerical character separated by a delimiter such as a comma or a space will be interpreted as complex data. For example, the string:

```
"1.2 1.9"
```

will be converted into the number:

```
1.2 + 1.9i
```

Parentheses surrounding the numerical elements in the string will be ignored as will be a following *i*. The following strings will be interpreted as the same by **strtof**.

```
"(2.31 4.72)" "2.31 4.73" "2.31,4.73i"
```

Example

```
//Create a string array  
  
string sa = { "1.1""2.2""3.3", "4.4""5.5""6.6" };
```

strtofcp1x

```
num = strtof(sa);
```

After the code above, *num* is a numeric matrix with the following values:

```
1.100  2.200  3.300
4.400  5.500  6.600
```

See Also

[strtofcp1x](#), [f1ostrC](#)

strtofcp1x

Purpose

Converts a string array to a complex numeric matrix.

Format

```
x = strtofcp1x(sa);
```

Input

<i>sa</i>	NxK string array containing numeric data.
-----------	---

Output

<i>x</i>	NxK complex matrix.
----------	---------------------

Remarks

strtofcp1x supports both real and complex data. It is slower than **strtof** for real matrices. **strtofcp1x** requires the presence of the real part. The imaginary part can

be absent.

See Also

[strtof](#), [fstrC](#)

strtrim

Purpose

Strips all white space characters from the left and right side of each element in a string array.

Format

```
y = strtrim(sa);
```

Input

<i>sa</i>	NxM string array.
-----------	-------------------

Output

<i>y</i>	NxM string array.
----------	-------------------

Examples

Example 1: Basic example

```
//Create a string with leading and trailing spaces  
str = "    Time Series Estimation    ";
```

strtrim

```
//Remove leading and trailing spaces from string
str_mod = strtrim(str);
```

After the code above, *str* should contain:

```
Time Series Estimation
```

while *str_mod* should contain the same characters, but have all spaces on the right and left removed:

```
Time Series Estimation
```

Example 2: Create a string array of variable names

strtrim can be useful when parsing tokens from a text file. For example, you may read the header row of a CSV file, containing something like the *header_vars* variable in the example below and want to create a string array in which each variable name is an element in the string array.

```
//Create string similar to a messy header row
header_vars = "alpha, beta ,   gamma";

//Split string into 3x1 string array at comma locations//
(notice the transpose operator ' ' at the end of the state-
ment
header_sa = strsplit(header_vars, ",")';
```

After the above code, *header_sa* will equal (NOTE: the print function will automatically align the string array, so 'print header_sa' will make it appear as if the leading and trailing spaces are gone. To see the spaces, you will need to print individual elements i.e. 'print header_sa[1]; print header_sa[2];', etc):

```
alpha
beta
gamma
```

You can remove the leading and trailing spaces with **strtrim**, like this:

```
//Remove leading and trailing spaces  
header_sa = strtrim(header_sa);
```

Which will transform *header_sa* into:

```
alpha  
beta  
gamma
```

Source

strfns.src

See Also

[strtriml](#), [strtrimr](#), [strtrunc](#), [strtrunc1](#), [strtruncpad](#), [strtruncr](#)

strtriml

Purpose

Strips all whitespace characters from the left side of each element in a string array.

Format

```
y = strtriml(sa);
```

Input

<i>sa</i>	NxM string array.
-----------	-------------------

strtrimr

Output

y NxM string array.

Source

strfns.src

See Also

[strtrimr](#), [strtrunc](#), [strtrunc1](#), [strtruncpad](#), [strtruncr](#)

strtrimr

Purpose

Strips all whitespace characters from the right side of each element in a string array.

Format

```
y = strtrimr(sa);
```

Input

sa NxM string array.

Output

y NxM string array.

Source

strfns.src

See Also

[strtriml](#), [strtrunc](#), [strtrunc1](#), [strtruncpad](#), [strtruncr](#)

strtrunc

Purpose

Truncates all elements of a string array to not longer than the specified number of characters.

Format

```
y = strtrunc(sa, maxlen);
```

Input

<i>sa</i>	NxK string array.
<i>maxlen</i>	1xK or 1x1 matrix, maximum length.

Output

<i>y</i>	NxK string array result.
----------	--------------------------

Example

```
string s = { "best", "linear", "unbiased", "estimator" };  
ss = strtrunc(s, 6);
```

After the code above, the variables **s** and **ss** are equal to:

```
best  
linear
```

strtrunc1

```
s = unbiased
    estimator

    best
    linear
ss = unbiased
    estima
```

See Also

[strtriml](#), [strtrimr](#), [strtrunc1](#), [strtruncpad](#), [strtruncr](#)

strtrunc1

Purpose

Truncates the left side of all elements of a string array by a user-specified number of characters.

Format

```
y = strtrunc1(sa, ntrunc);
```

Input

<i>sa</i>	NxM, Nx1, 1xM, or 1x1 string array.
<i>ntrunc</i>	NxM, Nx1, 1xM, or 1x1 matrix containing the number of characters to strip.

Output

<i>y</i>	string array result.
----------	----------------------

Source

strfns.src

See Also

[strtriml](#), [strtrimr](#), [strtrunc](#), [strtruncpad](#), [strtruncr](#)

strtruncpad

Purpose

Truncates all elements of a string array to the specified number of characters, adding spaces on the end as needed to achieve the exact length.

Format

$y = \text{strtruncpad}(sa, maxlen);$

Input

sa	NxK string array.
$maxlen$	1xK or 1x1 matrix, maximum length.

Output

y	NxK string array result.
-----	--------------------------

See Also

[strtriml](#), [strtrimr](#), [strtrunc](#), [strtruncr](#), [strtruncpad](#)

strtruncr

strtruncr

Purpose

Truncates the right side of all elements of a string array by a user-specified number of characters.

Format

```
y = strtruncr(sa, ntrunc);
```

Input

<i>sa</i>	NxM, Nx1, 1xM, or 1x1 string array.
<i>ntrunc</i>	NxM, Nx1, 1xM, or 1x1 matrix containing the number of characters to strip.

Output

<i>y</i>	String array result.
----------	----------------------

Source

strfns.src

See Also

[strtriml](#), [strtrimr](#), [strtrunc](#), [strtruncr](#), [strtruncpad](#)

submat

Purpose

Extracts a submatrix of a matrix, with the appropriate rows and columns given by the elements of vectors.

Format

```
y = submat(x, r, c);
```

Input

x	NxK matrix.
r	LxM matrix of row indices.
c	PxQ matrix of column indices.

Output

y	(L*M)x(P*Q) submatrix of x , y may be larger than x .
----------	--

Remarks

If $r = 0$, then all rows of x will be used. If $c = 0$, then all columns of x will be used.

Example

```
//Create 12x1 vector with consecutive numbers
x = seqa(1, 1, 12);

//Reshape the 12x1 vector into a 3x4 matrix
x = reshape(x, 3, 4);

v1 = 1 3;
v2 = 2 4;

//Extract sub-matrices
y = submat(x, v1, v2);
z = submat(x, 0, v2);
```

After the code above, the matrix values are:

subscat

```
      1  2  3  4
x =  5  6  7  8
     9 10 11 12

      2  4
y = 10 12

      2  4
z =  6  8
     10 12
```

See Also

[diag](#), [vec](#), [reshape](#)

subscat

Purpose

Changes the values in a vector depending on the category a particular element falls in.

Format

```
y = subscat(x, breaks, levels);
```

Input

<i>x</i>	Nx1 vector.
<i>breaks</i>	Px1 numeric vector, containing breakpoints specifying the ranges within which substitution is to be made. This MUST be sorted in ascending order. <i>breaks</i> can contain a missing value as a separate

levels

category if the missing value is the first element in *breaks*.

If *breaks* is a scalar, all matches must be exact for a substitution to be made.

Px1 vector, containing values to be substituted.

Output

y

Nx1 vector, with the elements in *levels* substituted for the original elements of *x* according to which of the regions the elements of *x* fall into:

```

 $x \leq \text{breaks}[1] \rightarrow \text{levels}[1]$ 
 $\text{breaks}[1] < x \leq \text{breaks}[2] \rightarrow \text{levels}[2]$ 
...
 $\text{breaks}[p - 1] < x \leq \text{breaks}[p] \rightarrow \text{levels}[p]$ 
 $x > \text{breaks}[p] \rightarrow \text{the original value of } x$ 

```

If missing is not a category specified in *breaks*, missings in *x* are passed through without change.

Example

Example 1

```

//BMI Data
bmi = { 36,
        19,
        24,
        38,
        34,
        16,
        26,

```

subscat

```
37,  
20,  
34 };  
  
//Set the breakpoints for the new categories  
breaks = { 18.5, 25, 30, 40 };  
  
//The categorical levels  
levels = { 0, 1, 2, 3 };  
  
bmi_levels = subscat(bmi, breaks, levels);
```

The above code assigns the following values:

bmi = 36	bmi_levels = 3
19	1
24	1
38	3
34	3
16	0
26	2
37	3
20	1
34	3

Example 2

This example combines 2 levels in a categorical label into one category.

```
//Create categorical vector with 3 levels  
x = { 1,  
      1,  
      2,  
      2,  
      1,
```



```
1,  
2,  
0,  
2,  
0 };  
  
//Assign all instances of 2 to 1, merging the second and  
third categories  
x = subscat(x, 2, 1);
```

After the code above, `x` is equal to:

```
1  
1  
1  
1  
1  
1  
1  
1  
0  
1  
0
```

Replacing instances of one particular value with another value can also be accomplished with **reclassify** and **substute**

Remarks

reclassifyCuts offers functionality similar to **subscat**, but:

- Also assigns values to data past the final breakpoint.
- Offers the option of whether the breakpoints are open or closed on the right(e.g., $<$ or \leq).
- Assigns the input to two categories in the case of a single breakpoint, (e.g., $level_1 < break < level_2$). Whereas, **subscat** tests for equality in the case of a single breakpoint.

substute

See Also

[reclassify](#), [reclassifyCuts](#), [substute](#)

substute

Purpose

Substitutes new values for old values in a matrix, depending on the outcome of a logical expression.

Format

$y = \text{substute}(x, \text{ } e, \text{ } v);$

Input

x	$N \times K$ matrix containing the data to be changed.
e	$L \times M$ matrix, $E \times E$ conformable with x containing 1's and 0's.
v	$P \times Q$ matrix, $E \times E$ conformable with x and e , containing the values to be substituted for the original values of x when the corresponding element of e is 1.

Output

y $\max(N,L,P)$ by $\max(K,M,Q)$ matrix.

Remarks

The e matrix is usually the result of an expression or set of expressions using dot

conditional and boolean operators.

Example

Example 1

Set all elements between 0 and 2.25×10^{-16} equal to 0.

```
//Create example vector
x = { 3.8e-21,
      1.0,
      3.5,
      2.7e-18,
      0.5,
      3.0,
      1.1e-16,
      0.5,
      2.2,
      4.0 };

//Substitute all values less than  $2.2 \times 10^{-16}$  with a zero
x = substitute(x, x .<  $2.2 \times 10^{-16}$ , 0);
```

After the code above, x is equal to:

```
0.0
1.0
3.5
0.0
0.5
3.0
0.0
0.5
```

substitute

2.2
4.0

Example 2

```
//Create a matrix with character elements
//in the first column
x = { Y 55 30,
      N 57 18,
      Y 24 3,
      N 63 38,
      Y 55 32,
      N 37 11 };

//Create a rows(x) by 1 vector with a '1' for each row
// that:
// 1) The first element is a Y
// 2) The second element is greater than or equal to 55
// 3) The third element is greater than or equal to 30
//If the row does not meet ALL of these conditions a 0 will
//be returned.
e = (x[.,1] == "Y") .and (x[.,2] >= 55) .and (x[.,3] >=
30);

//Substitute an 'R' for the first element in every row that
//meets the conditions specified in the assignment to 'e'
x[.,1] = substitute(x[.,1],e, "R");
```

The vector e is equal to:

1
0
0

```
0  
1  
0
```

Here is what x looks like after substitution:

```
R 55 30  
N 57 18  
Y 24  3  
N 63 38  
R 55 32  
N 37 11
```

Source

`datatran.src`

See Also

[code](#), [recode](#), [reclassifyCuts](#), [reclassify](#), [rescale](#)

subvec

Purpose

Extracts an $N \times 1$ vector of elements from an $N \times K$ matrix.

Format

```
 $y = \text{subvec}(x, ci);$ 
```

subvec

Input

x	$N \times K$ matrix.
ci	$N \times 1$ vector of column indices.

Output

y	$N \times 1$ vector containing the elements in x indicated by ci .
-----	--

Remarks

Each element of y is from the corresponding row of x and the column set by the corresponding row of ci . In other words, $y[i] = x[i, ci[i]]$.

Example

```
//Create an additive sequence from 1-12, i.e. 1, 2, 3,...12
x = seqa(1, 1, 12);

//Reshape the sequential vector 'x' into a 4x3 matrix
x = reshape(x, 4, 3);

//The column indices (one per row of 'x') indicating which
//values to extract from 'x'
ci = { 2, 3, 1, 3 };

//Extract subvector from 'x' and assign it to 'y'
y = subvec(x, ci);
```

After the above code, x and y are equal to:

	1	2	3
$x =$	4	5	6

```
      7  8  9
    10 11 12

      2
    y = 6
        7
      12
```

sumc

Purpose

Computes the sum of each column of a matrix or the sum across the second-fastest moving dimension of an L-dimensional array.

Format

```
y = sumc(x);
```

Input

x	$N \times K$ matrix or L-dimensional array where the last two dimensions are $N \times K$.
-----	---

Output

y	$K \times 1$ vector or L-dimensional array where the last two dimensions are $K \times 1$.
-----	---

Example

```
//Create a 12x1 vector containing an additive sequence
```

sumc

```
//counting by twos, from 0-22, i.e. 2, 4, 6, 8...22
x = seqa(0,2,12);

//Reshape the 12x1 vector 'x' into a 3x4 matrix
x = reshape(x,3,4);

//Sum the columns
y = sumc(x);
```

After the above code, the variables *x* and *y* are equal to:

```
      0  2  4  6
x =   8 10 12 14
     16 18 20 22

      24
y =   30
     36
     42
```

```
//Create an additive sequence from 1-24 and reshape it into
//a 2x3x4 array
a = areshape(seqa(1,1,24),2|3|4);

//Sum the columns across the second fastest moving
//dimension
z = sumc(a);
```

a is a 2x3x4 array such that:

```
Plane [1,...]
```

1.0000000	2.0000000	3.0000000	4.0000000
5.0000000	6.0000000	7.0000000	8.0000000
9.0000000	10.000000	11.000000	12.000000

Plane [2,...]

13.000000	14.000000	15.000000	16.000000
17.000000	18.000000	19.000000	20.000000
21.000000	22.000000	23.000000	24.000000

Variable *z* is a 2x4x1 array equal to:

Plane [1,...]

15.000000
18.000000
21.000000
24.000000

Plane [2,...]

51.000000
54.000000
57.000000
60.000000

See Also

[cumsumc](#), [meanc](#), [stdc](#)

sumr

Purpose

Computes the sum of each row of a matrix or the sum of the fastest moving dimension of an L-dimensional array.

sumr

Format

`y = sumr(x);`

Input

`x` NxK matrix or L-dimensional array where the last two dimensions are NxK.

Output

`y` Nx1 vector or L-dimensional array where the last two dimensions are Nx1.

Example

```
//Create an additive sequence from 1-12 and reshape it into
//a 3x4 matrix
x = reshape(seqa(1,1,12),3,4);

//Sum the rows
y = sumr(x);
```

After the above code, the variables `x` and `y` will be:

```
1  2  3  4      10
x = 5  6  7  8    y = 26
    9 10 11 12    42
```

```
//Reshape an additive sequence from 1-24 into a 2x3x4
//dimensional array
a = areshape(seqa(1,1,24),2|3|4);
z = sumr(a);
```

`a` is a 2x3x4 array such that:

```
Plane [1,...]

      1.0000000    2.0000000    3.0000000    4.0000000
      5.0000000    6.0000000    7.0000000    8.0000000
      9.0000000   10.000000    11.000000    12.000000

Plane [2,...]

      13.000000    14.000000    15.000000    16.000000
      17.000000    18.000000    19.000000    20.000000
      21.000000    22.000000    23.000000    24.000000
```

The variable *z* is equal to:

```
Plane [1,...]

      10.000000
      26.000000
      42.000000

Plane [2,...]

      58.000000
      74.000000
      90.000000
```

See Also

[sumc](#)

surface

Purpose

Graphs a 3-D surface. NOTE: This function is for use with the deprecated PQG graphics. Use **plotSurface** instead.

surface

Library

pgraph

Format

```
surface(x, y, z);
```

Input

<i>x</i>	1xK vector, the X axis data.
<i>y</i>	Nx1 vector, the Y axis data.
<i>z</i>	NxK matrix, the matrix of height data to be plotted.

Global Input

<i>_psurf</i>	2x1 vector, controls 3-D surface characteristics. [1] if 1, show hidden lines. Default 0. [2] color for base (default 7). The base is an outline of the X-Y plane with a line connecting each corner to the surface. If 0, no base is drawn.
<i>_pticout</i>	scalar, if 0 (default), tick marks point inward, if 1, tick marks point outward.
<i>_pzclr</i>	Z level color control. There are 3 ways to set colors for the Z levels of a surface graph. 1. To specify a single color for the entire surface plot, set the color control variable to a scalar value 1-15. For example:

```
_pzclr = 15;
```

2. To specify multiple colors distributed evenly over the entire Z range, set the color control variable to a vector containing the desired colors only. **GAUSS** will automatically calculate the required corresponding Z values for you. The following example will produce a three color surface plot, the Z ranges being lowest=blue, middle=light blue, highest=white:

```
_pzclr = { 1, 10, 15 };
```

3. To specify multiple colors distributed over selected ranges, the Z ranges as well as the colors must be manually input by the user. The following example assumes -0.2 to be the minimum value in the `z` matrix:

```
_pzclr = { -0.2 1,
/* z >= -0.2 blue */
0.0 10,
/* z >= 0.0 light blue */
0.2 15 };
/* z >= 0.2 white */
```

Since a Z level is required for each selected color, the user must be responsible to compute the minimum value of the `z` matrix as the first Z range element. This may be most easily accomplished by setting the `_pzclr` matrix as shown above (the first element being an arbitrary value), then resetting the first element to the minimum `z` value as follows:

svd

s

```
_pzclr = { 0.0 1,  
          0.0 10,  
          0.2 15 };  
_pzclr[1,1] = minc(minc(z));
```

See **PQG Graphics Colors**, for the list of available colors.

Remarks

surface uses only the minimum and maximum of the X axis data in generating the graph and tick marks.

Source

psurface.src

See Also

[volume](#), [view](#)

svd

Purpose

Computes the singular values of a matrix.

Format

```
s = svd(x);
```

Input

x NxP matrix whose singular values are to be computed.

Output

s Mx1 vector, where $M = \min(N,P)$, containing the singular values of *x* arranged in descending order.

Global Input

_svderr scalar, if the singular values cannot be computed, *_svderr* will be nonzero.

Examples

```
//Create a 10x3 matrix
x = {  -0.60    3.50    0.47,
       8.40    16.50    0.27,
       11.40    6.50    0.17,
       7.40    -0.50   -2.43,
       -9.60   -10.50    0.57,
       -17.60   -5.50    0.67,
       -12.60  -14.50    0.87,
       18.40    12.50   -1.43,
       -11.60  -19.50    0.77,
       6.40    11.50    0.07 };

//Calculate the singular values
s = svd(x);
```

After the code above, *s* will be equal to:

```
49.58
14.96
2.24
```

svd1

Remarks

1. **svd** is not threadsafe. New code should use **svds** instead.
2. Error handling is controlled with the low bit of the trap flag.

s

trap	set <code>_svderr</code> to a non-zero value and terminate with message
0	
trap	set <code>_svderr</code> to a non-zero value and continue execution
1	

Source

`svd.src`

See Also

[svd2](#), [svds](#)

svd1

Purpose

Computes the singular value decomposition of a matrix so that: $x = u * s * v'$.

Format

$\{ u, s, v \} = \mathbf{svd1}(x);$

Input

x	NxP matrix whose singular values are to be computed.
-----	--

Output

<i>u</i>	NxN matrix, the left singular vectors of <i>x</i> .
<i>s</i>	NxP diagonal matrix, containing the singular values of <i>x</i> arranged in descending order on the principal diagonal.
<i>v</i>	PxP matrix, the right singular vectors of <i>x</i> .

Global Output

`_svderr` scalar, if the singular values cannot be computed, `_svderr` will be non-zero.

Example

```
//Create 6x3 matrix
x = { -9.35    15.67   -41.75,
      -13.55   40.97    15.55,
      -0.95   -17.03   40.15,
       8.15    -9.73   13.15,
       2.35   -36.73  -43.55,
      13.35    6.87   16.45 };

//Perform matrix decomposition
{ u, s, v } = svd1(x);
```

After the code above, the outputs will have the following values;

```
u =  0.44    -0.49    -0.06    0.36    -0.24    0.61
     -0.35    -0.60    -0.28    0.12     0.65   -0.08
     -0.41     0.46    -0.53    0.07     0.03    0.58
     -0.12     0.25     0.24    0.91     0.08   -0.18
       0.67     0.35    -0.13   -0.02     0.64     0.05
```

svd2

s

```
      -0.23      0.04      0.75      -0.17      0.33      0.50
s = 79.03      0.00      0.00
      0.00     60.19      0.00
      0.00      0.00     17.16
      0.00      0.00      0.00
      0.00      0.00      0.00
      0.00      0.00      0.00
v = -0.02      0.26      0.97
     -0.32     -0.91      0.24
     -0.95      0.31     -0.10
```

Remarks

1. **svd1** is not threadsafe. New code should use **svdusv** instead.
2. Error handling is controlled with the low bit of the trap flag.

trap 0 set `_svderr` to a non-zero value and terminate with message
trap 1 set `_svderr` to a non-zero value and continue execution

Source

`svd.src`

See Also

[svd](#), [svd2](#), [svdusv](#)

svd2

Purpose

Computes the singular value decomposition of a matrix so that: $x = u * s * v'$
(compact u).

Format

```
{ u, s, v } = svd2(x);
```

Input

x	$N \times P$ matrix whose singular values are to be computed.
-----	---

Output

u	$N \times N$ or $N \times P$ matrix, the left singular vectors of x . If $N > P$, then u will be $N \times P$, containing only the P left singular vectors of x .
s	$N \times P$ or $P \times P$ diagonal matrix, containing the singular values of x arranged in descending order on the principal diagonal. If $N > P$, then s will be $P \times P$.
v	$P \times P$ matrix, the right singular vectors of x .

Global Output

<code>_svderr</code>	scalar, if all of the singular values are correct, <code>_svderr</code> is 0. If the singular values cannot be computed, <code>_svderr</code> is set to a non-zero value.
----------------------	---

Examples

```
//Create a 10x3 matrix
x = {  -0.60    3.50    0.47,
        8.40    16.50    0.27,
        11.40    6.50    0.17,
        7.40    -0.50   -2.43,
```

svd2

s

```
-9.60  -10.50   0.57,  
-17.60  -5.50   0.67,  
-12.60  -14.50   0.87,  
 18.40   12.50  -1.43,  
-11.60  -19.50   0.77,  
  6.40   11.50   0.07 };
```

```
//Calculate the singular values  
{ u, s, v } = svd2(x);
```

After the code above, u , s and v will be equal to:

```
u =  0.04    0.20   -0.11  
     0.36    0.38   -0.14  
     0.25   -0.23   -0.44  
     0.10   -0.39    0.75  
    -0.29   -0.04   -0.06  
    -0.33    0.57    0.35  
    -0.39   -0.08   -0.14  
     0.44   -0.29    0.10  
    -0.44   -0.37   -0.25  
     0.26    0.24   -0.07  
  
s = 49.58    0.00    0.00  
     0.00   14.96    0.00  
     0.00    0.00    2.24  
  
v =  0.70   -0.70   -0.10  
     0.71    0.70    0.05  
    -0.04    0.10   -0.99
```

Remarks

1. **svd2** is not threadsafe. New code should use **svdcusv** instead.
2. Error handling is controlled with the low bit of the trap flag. If the singular

values cannot be computed, `_svderr` will be set to a non-zero value.

trap	set <code>_svderr</code> to a non-zero value and terminate with
0	message
trap	set <code>_svderr</code> to a non-zero value and continue execution
1	

Source

svd.src

See Also

[svd](#), [svd1](#), [svdcusv](#)

svdcusv

Purpose

Computes the singular value decomposition of `x` so that: $x = u * s * v'$ (compact u).

Format

$\{ \ u, \ s, \ v \} = \text{svdcusv}(x);$

Input

<code>x</code>	NxP matrix or K-dimensional array where the last two dimensions are NxP, whose singular values are to be computed.
----------------	--

svdcusv

Output

s

u	NxN or NxP matrix or K-dimensional array where the last two dimensions are NxN or NxP, the left singular vectors of x . If $N > P$, u is NxP, containing only the P left singular vectors of x .
s	NxP or PxP diagonal matrix or K-dimensional array where the last two dimensions describe NxP or PxP diagonal arrays, the singular values of x arranged in descending order on the principal diagonal. If $N > P$, s is PxP.
v	PxP matrix or K-dimensional array where the last two dimensions are PxP, the right singular vectors of x .

Examples

```
//Create a 10x3 matrix
x = {  -0.60    3.50    0.47,
       8.40    16.50    0.27,
       11.40    6.50    0.17,
        7.40   -0.50   -2.43,
       -9.60  -10.50    0.57,
      -17.60   -5.50    0.67,
      -12.60  -14.50    0.87,
       18.40   12.50   -1.43,
      -11.60  -19.50    0.77,
        6.40   11.50    0.07 };

//Calculate the singular values
{ u, s, v } = svdcusv(x);
```

After the code above, u , s and v will be equal to:

```

u =  0.04      0.20     -0.11
      0.36      0.38     -0.14
      0.25     -0.23     -0.44
      0.10     -0.39      0.75
     -0.29     -0.04     -0.06
     -0.33      0.57      0.35
     -0.39     -0.08     -0.14
      0.44     -0.29      0.10
     -0.44     -0.37     -0.25
      0.26      0.24     -0.07

s = 49.58      0.00      0.00
      0.00     14.96      0.00
      0.00      0.00      2.24

v =  0.70     -0.70     -0.10
      0.71      0.70      0.05
     -0.04      0.10     -0.99

```

s

Remarks

- If x is an array, the resulting arrays u , s and v will contain their respective results for each of the corresponding 2-dimensional arrays described by the two trailing dimensions of x . In other words, for a 10x4x5 array x :
 - u will be a 10x4x4 array containing the left singular vectors of each of the 10 corresponding 4x5 arrays contained in x .
 - s will be a 10x4x5 array containing the singular values.
 - v will be a 10x5x5 array containing the right singular vectors
- Error handling is controlled by the `trap` command. If not all of the singular values can be computed:

```

trap      terminate with an error message
0

```

svds

trap set the first element of s to a scalar missing value and
1 continue execution

```
//Turn on error trapping
trap 1;

//Compute singular value decomposition
{ u, s, v } = svdcusv(x);

//Check for failure or success
if scalmiss(s[1,1]);
    //Code for failure case
endif;
```

Note that in the `trap 1` case, if the input to `svdcusv` is a multi-dimensional array and the singular values for a submatrix fail to compute, only the first value of that s submatrix will be set to a missing value. For a 3 dimensional array, you could change the `if` check in the above example to:

```
//Check for success or failure of each submatrix
if ismiss(s[:,1,1]);
```

See Also

[svd2](#), [svds](#), [svdusv](#)

svds

Purpose

Computes the singular values of a x .

Format

```
s = svds(x);
```

Input

x	$N \times P$ matrix or K-dimensional array where the last two dimensions are $N \times P$, whose singular values are to be computed.
-----	---

Output

s	$\min(N,P) \times 1$ vector or K-dimensional array where the last two dimensions are $\min(N,P) \times 1$, the singular values of x arranged in descending order.
-----	--

Examples

```
//Create a 10x3 matrix
x = {  -0.60      3.50      0.47,
        8.40      16.50      0.27,
       11.40       6.50      0.17,
        7.40     -0.50     -2.43,
       -9.60    -10.50      0.57,
      -17.60     -5.50      0.67,
      -12.60    -14.50      0.87,
       18.40     12.50     -1.43,
      -11.60    -19.50      0.77,
        6.40     11.50      0.07 };

//Calculate the singular values
s = svds(x);
```

After the code above, s will be equal to:

svds

```
49.58
14.96
2.24
```

Remarks

1. If x is an array, the result will be an array containing the singular values of each of the 2-dimensional arrays described by the two trailing dimensions of x . In other words, for a 10x4x5 array x , s will be a 10x4x1 array containing the singular values of each of the 10 4x5 arrays contained in x .
2. If the singular values cannot be computed, either the program will be terminated with an error message, or the first element of the return, $s[1]$, is set to a missing value. This behavior is controlled by the `trap` command. Below is an example with error trapping:

```
//Turn on error trapping
trap 1;

//Calculate singular values
s = svds(x);

//Check for success or failure
if ismiss(s);
//Code to handle failure case
endif;
```

Note that in the `trap 1` case, if the input to `svds` is a multi-dimensional array and the singular values for a submatrix fail to compute, only the first value of that s submatrix will be set to a missing value. For a 3 dimensional array, you could change the `if` check in the above example to:

```
//Check for success or failure of each submatrix
if ismiss(s[.,1,1]);
```

3. Call either **svdcusv** or **svdusv**, to also calculate the right and left singular vectors

See Also

[svd](#), [svdcusv](#), [svdusv](#)

svdusv

Purpose

Computes the singular value decomposition of x so that: $x = u * s * v'$.

Format

$$\{ \ u, \ s, \ v \} = \text{svdusv}(x);$$

Input

x	NxP matrix or K-dimensional array where the last two dimensions are NxP, whose singular values are to be computed.
-----	--

Output

u	NxN matrix or K-dimensional array where the last two dimensions are NxN, the left singular vectors of x .
s	NxP diagonal matrix or K-dimensional array where the last two dimensions describe NxP diagonal arrays, the singular values of x arranged in descending order on the principal diagonal.
v	PxP matrix or K-dimensional array where the last two dimensions are PxP, the right singular vectors of x .

Example

s

```
//Create 6x3 matrix
x = { -9.35    15.67   -41.75,
      -13.55   40.97    15.55,
      -0.95   -17.03   40.15,
       8.15    -9.73   13.15,
       2.35   -36.73  -43.55,
      13.35     6.87   16.45 };

//Perform matrix decomposition
{ u, s, v } = svdusv(x);
```

After the code above, the outputs will have the following values;

```
u =  0.44    -0.49    -0.06    0.36    -0.24    0.61
     -0.35    -0.60    -0.28    0.12     0.65   -0.08
     -0.41     0.46    -0.53    0.07     0.03    0.58
     -0.12     0.25     0.24    0.91     0.08   -0.18
       0.67     0.35    -0.13   -0.02     0.64     0.05
     -0.23     0.04     0.75   -0.17     0.33     0.50

s = 79.03     0.00     0.00
     0.00    60.19     0.00
     0.00     0.00    17.16
     0.00     0.00     0.00
     0.00     0.00     0.00
     0.00     0.00     0.00

v = -0.02     0.26     0.97
     -0.32    -0.91     0.24
     -0.95     0.31    -0.10
```

Remarks

1.

If x is an array, the resulting arrays u , s and v will contain their respective results for each of the corresponding 2-dimensional arrays described by the two trailing dimensions of x . In other words, for a $10 \times 4 \times 5$ array x :

- u will be a $10 \times 4 \times 4$ array, containing the left singular vectors of each of the 10 corresponding 4×5 arrays contained in x .
- s will be a $10 \times 4 \times 5$ array, containing the singular values.
- v will be a $10 \times 5 \times 5$ array containing, the right singular vectors.

2. Error handling is controlled by the `trap` command. If not all of the singular values can be computed:

trap terminate with an error message

0

trap set the first element of s to a scalar missing value and

1 continue execution

```
//Turn on error trapping
trap 1;

//Compute singular value decomposition
{ u, s, v } = svdusv(x);

//Check for failure or success
if scalmiss(s[1,1]);
    //Code for failure case
endif;
```

Note that in the `trap 1` case, if the input to **svdusv** is a multi-dimensional array and the singular values for a submatrix fail to compute, only the first value

sylvester

of that *s* submatrix will be set to a missing value. For a 3 dimensional array, you could change the `if` check in the above example to:

```
//Check for success or failure of each submatrix
if ismiss(s[.,1,1]);
```

See Also

[svdl](#), [svdcusv](#), [svds](#)

sylvester

Purpose

Computes the solution to the Sylvester matrix equation, $AX + XB = C$.

Format

$X = \text{sylvester}(A, B, C);$

Input

<i>A</i>	MxM real or complex matrix.
<i>B</i>	NxN real or complex matrix.
<i>C</i>	MxN real or complex matrix.

Output

<i>X</i>	MxN matrix, solution to the equation $AX + XB = C$.
----------	--

Examples

Example 1: Real input

```
//Create a 3 x 3 real matrix
A = { 0.9069 -0.3150 -0.9732,
      0.6023 0.6848 0.4925,
      -0.8555 -0.7430 0.6521 };

//Create a 2 x 2 real matrix
B = { -0.9876 0.4503 ,
      -0.3043 0.9807 };

//Create a 3 x 2 real matrix
C = { -0.8625 0.5247,
      0.6331 -0.3334,
      0.7912 0.0711 };

//Solve the Sylvester matrix equation
X = sylvester(A, B, C);
```

After the code above, X will equal the 3 x 2 matrix:

```
X = -0.4279 0.3246
     -1.0525 -0.0013
      1.1609 -0.1071
```

Example 2: Complex input

```
//Create a 3 x 3 complex matrix
A = { 7 + 7i 4 + 10i 2 + 8i,
      10 - 3i -7 - 5i -10 - 7i,
      3 + 5i -10 - 2i 2 - 4i };

//Create a 2 x 2 complex matrix
B = { 5 + 1i -5 - 8i,
      8 - 10i 8 - 1i };

//Create a 3 x 2 complex matrix
```

sysstate

```
C = { -9 - 3i    -1 - 1i,  
      9 - 8i     -5 + 8i,  
      -1 - 2i    -5 + 5i };  
  
//Solve the Sylvester matrix equation  
X = sylvester(A, B, C);
```

After the code above, X will equal the 3 x 2 complex matrix:

```
X =    0.1697 - 0.2242i    -0.5923 + 0.2221i  
      -0.5684 + 0.4562i     0.3670 - 0.7153i  
      -0.7502 + 0.2470i    -0.0636 - 0.4208i
```

Remarks

The equation $AX + XB = C$ will not have a unique solution if the eigenvalues of the matrices A and $-B$ are equal. In this case an error will be returned.

See Also

[hess](#), [schur](#)

sysstate

Purpose

Gets or sets general system parameters.

Format

```
{ rets... } = sysstate(case, y);
```

Remarks

The available cases are as follows:

Case 1	Version Information Returns the current GAUSS version information in an 8-element numeric vector.
Cases 2-7	GAUSS System Paths Gets or sets GAUSS system path.
Case 8	Complex Number Toggle Controls automatic generation of complex numbers in sqrt , ln , and log for negative arguments.
Case 9	Complex Trailing Character Gets or sets trailing character for the imaginary part of a complex number.
Case 10	Printer Width Gets or sets lprint width.
Case 11	Auxiliary Output Width Gets or sets the auxiliary output width.
Case 13	LU Tolerance Gets or sets singularity tolerance for LU decomposition in current thread.
Case 14	Cholesky Tolerance Gets or sets singularity tolerance for Cholesky decomposition in current thread.
Case 15	Screen State Gets or sets window state as controlled by screen command.
Case 18	Auxiliary Output Gets auxiliary output parameters.

Case 19	Get/Set Format Gets or sets format parameters.
Case 21	Imaginary Tolerance Gets or sets imaginary tolerance in current thread.
Case 22	Source Path Gets or sets the path the compiler will search for source files.
Case 24	Dynamic Library Directory Gets or sets the path for the default dynamic library directory.
Case 25	Temporary File Path Gets or sets the path GAUSS will use for temporary files.
Case 26	Interface Mode Returns the current interface mode.
Case 28	Random Number Generator Parameters Gets or sets parameters used by the random number generation commands.
Case 30	Base Year Toggle Specifies whether year value returned by date is to include base year (1900) or not.
Case 32	Global LU Tolerance Gets or sets global singularity tolerance for LU decomposition.
Case 33	Global Cholesky Tolerance Gets or sets global singularity tolerance for Cholesky decomposition.

Case 34

Global Imaginary Tolerance
Gets or sets global imaginary tolerance.

Case 1: Version Information

Purpose

Returns the current **GAUSS** version information in an 8-element numeric vector.

Format

```
vi = sysstate(1,0);
```

Output

vi

8x1 numeric vector containing version information:
[1] Major version number.
[2] Minor version number.
[3] Revision.
[4] Machine type.
[5] Operating system.
[6] Runtime module.
[7] Light version.
[8] Always 0.
vi[4] indicates the type of machine on which **GAUSS** is running:
1 Intel x86
2 Sun SPARC
4 HP 9000

sysstate

S

7	Mac 32-bit PowerPC
<i>vi</i> [5] indicates the operating system on which GAUSS is running:	
3	Solaris
5	HP-UX
9	Windows
10	Linux
12	Mac OS

Cases 2-7: GAUSS System Paths

Purpose

Gets or sets **GAUSS** system path.

Format

oldpath = **sysstate**(*case*, *path*);

Input

<i>case</i>	scalar 2-7, path to set.
2	.exe file location.
3	<i>loadexe</i> path.
4	<i>save</i> path.
5	<i>load</i> , <i>loadm</i> path.
6	<i>loadf</i> , <i>loadp</i> path.
7	loads path.
<i>path</i>	scalar 0 to get path, or string containing the new path.

Output

oldpath string, original path.

Remarks

If *path* is of type matrix, the path will be returned but not modified.

Case 8: Complex Number Toggle

Purpose

Controls automatic generation of complex numbers in **sqrt**, **ln** and **log** for negative arguments.

Format

oldstate = **sysstate**(8, *state*);

Input

state scalar, 1, 0, or -1

Output

oldstate scalar, the original state.

Remarks

If *state* = 1, **log**, **ln**, and **sqrt** will return complex numbers for negative arguments. If *state* = 0, the program will terminate with an error message when negative numbers are passed to **log**, **ln**, and **sqrt**. If *state* = -1, the current state is returned and left unchanged. The default state is 1.

Case 9: Complex Trailing Character

sysstate

Purpose

Gets or sets trailing character for the imaginary part of a complex number.

Format

```
oldtrail = sysstate(9, trail);
```

Input

<i>trail</i>	scalar 0 to get character, or string containing the new trailing character.
--------------	---

Output

<i>oldtrail</i>	string, the original trailing character.
-----------------	--

Remarks

The default character is "i".

Case 10: Printer Width

Purpose

Gets or sets `lprint` width.

Format

```
oldwidth = sysstate(10, width);
```

Input

<i>width</i>	scalar, new printer width.
--------------	----------------------------

Output

oldwidth scalar, the current original width.

Remarks

If *width* is 0, the printer width will not be changed.

Case 11: Auxiliary Output Width

Purpose

Gets or sets the auxiliary output width.

Format

```
oldwidth = sysstate(11, width);
```

Input

width scalar, new output width.

Output

oldwidth scalar, the original output width.

Remarks

If *width* is 0 then the output width will not be changed.

This may also be set with the [outwidth](#) command.

See Also

[outwidth](#)

sysstate

Case 13: LU Tolerance

Purpose

Gets or sets singularity tolerance for LU decomposition in current thread.

Format

```
oldtol = sysstate(13, tol);
```

Input

tol scalar, new tolerance.

Output

oldtol scalar, the original tolerance.

Remarks

The tolerance must be ≥ 0 . If *tol* is negative, the tolerance is returned and left unchanged.

This tolerance is thread-safe. It must be set in the same thread in which it is to be referenced. To set the global singularity tolerance for LU decomposition, use case 32.

See Also

[croutp](#), [inv](#)

Case 14: Cholesky Tolerance

Purpose

Gets or sets singularity tolerance for Cholesky decomposition in current thread.

Format

```
oldtol = sysstate(14, tol);
```

Input

<i>tol</i>	scalar, new tolerance.
------------	------------------------

Output

<i>oldtol</i>	scalar, the original tolerance.
---------------	---------------------------------

Remarks

The tolerance must be ≥ 0 . If *tol* is negative, the tolerance is returned and left unchanged.

This tolerance is thread-safe. It must be set in the same thread in which it is to be referenced. To set the global singularity tolerance for Cholesky decomposition, use case 33.

This affects the following functions:

solpd

invpd for matrices $\leq 12 \times 12$

See Also

[chol](#), [invpd](#), [solpd](#)

Case 15: Screen State

Purpose

Gets or sets window state as controlled by [screen](#) command.

sysstate

Format

```
oldstate = sysstate(15, state);
```

Input

<i>state</i>	scalar, new window state.
--------------	---------------------------

Output

<i>oldstate</i>	scalar, the original window state.
-----------------	------------------------------------

Remarks

If *state* = 1, window output is turned on. If *state* = 0, window output is turned off. If *state* = -1, the state is returned unchanged.

See Also

[screen](#)

Case 18: Auxiliary Output

Purpose

Gets auxiliary output parameters.

Format

```
{ state, name } = sysstate(18,0);
```

Output

<i>state</i>	scalar, auxiliary output state, 1 - on, 0 - off.
<i>name</i>	string, auxiliary output filename.

See Also

[output](#)

Case 19: Get/Set Format

Purpose

Gets or sets format parameters.

Format

```
oldfmt = sysstate(19, fmt);
```

Input

<i>fmt</i>	scalar or 11x1 column vector containing the new formatparameters. Usually this will have come from a previous sysstate (19,0) call. See Output for description of matrix.
------------	--

Output

<i>oldfmt</i>	11x1 vector containing the current format parameters. The characters in quotes are components of the format string that gets passed through to the C library sprintf function: <div><div>[1]</div><div>format conversion type:</div><div><div>0</div><div>string format ("s")</div></div><div><div>1</div><div>compact format ("g").</div></div><div><div>2</div><div>auto format ("#g").</div></div><div><div>3</div><div>scientific format ("e").</div></div><div><div>4</div><div>decimal format ("f").</div></div></div>
---------------	--

s

	5	compact format, upper case ("G").
	6	auto format, upper case ("#G").
	7	scientific format, upper case ("E").
[2]		justification:
	0	right justification.
	1	left justification ("–").
[3]		sign:
	0	sign used only for negative numbers.
	1	sign always used (" +").
[4]		leading zero:
	0	no leading zero.
	1	leading zero ("0").
[5]		trailing character:
	0	no trailing character.
	1	trailing space (" ").
	2	trailing comma (" , ").
	3	trailing tab (" \ t").
[6]		row delimiter:
	0	no row delimiter.
	1	one newline between rows (" \ n").
	2	two newlines between rows (" \ n \ n").
	3	print "Row 1, Row 2, ..." before each row (" \ nRow %u \ n", where "%u" is the row number).

[7]	carriage line feed position:	
	0	newline row delimiters positioned before rows.
	1	newline row delimiters positioned after rows.
[8]	automatic line feed for row vectors.	
	0	newline row delimiters occur between rows of a matrix only if that matrix has more than one row.
	1	newline row delimiters occur between rows of a matrix, regardless of number of rows.
[9]	field width.	
[10]	precision.	
[11]	formatted flag.	
	0	formatting disabled.
	1	formatting enabled.

Remarks

If `fmt` is scalar 0, then the format parameters will be left unchanged.

See the `format` and `print` commands for more information on the formatting parameters.

See Also

[format](#), [print](#)

Case 21: Imaginary Tolerance

sysstate

Purpose

Gets or sets imaginary tolerance in current thread.

Format

```
oldtol = sysstate(21, tol);
```

Input

tol scalar, the new tolerance.

Output

oldtol scalar, the original tolerance.

Remarks

The imaginary tolerance is used to test whether the imaginary part of a complex matrix can be treated as zero or not. Functions that are not defined for complex matrices check the imaginary part to see if it can be ignored. The default tolerance is 2.23e-16, or machine epsilon.

If *tol*<0, the current tolerance is returned.

This tolerance is thread-safe. It must be set in the same thread in which it is to be referenced. To set the global imaginary tolerance, use case 34.

See Also

[hasimag](#)

Case 22: Source Path

Purpose

Gets or sets the path the compiler will search for source files.

Format

```
oldpath = sysstate(22, path);
```

Input

<i>path</i>	scalar 0 to get path, or string containing the new path.
-------------	--

Output

<i>oldpath</i>	string, original path.
----------------	------------------------

Remarks

If *path* is a matrix, the current source path is returned.

This resets the *src_path* configuration variable. *src_path* is initially defined in the **GAUSS** configuration file, `gauss.cfg`.

path can list a sequence of directories, separated by semicolons.

Resetting *src_path* affects the path used for subsequent `run` and `compile` statements.

Case 24: Dynamic Library Directory

Purpose

Gets or sets the path for the default dynamic library directory.

Format

```
oldpath = sysstate(24, path);
```

sysstate

Input

<i>path</i>	scalar 0 to get path, or string containing the new path.
-------------	--

Output

<i>oldpath</i>	string, original path.
----------------	------------------------

Remarks

If *path* is a matrix, the current path is returned.

path should list a single directory, not a sequence of directories.

Changing the dynamic library path does not affect the state of any DLL's currently linked to **GAUSS**. Rather, it determines the directory that will be searched the next time `dlibrary` is called.

UNIX

Changing the path has no effect on **GAUSS**'s default DLL, `libgauss.so`. `libgauss.so` must always be located in the GAUSSHOME directory.

Windows

Changing the path has no effect on **GAUSS**'s default DLL, `gauss.dll`. `gauss.dll` must always be located in the GAUSSHOME directory.

See Also

[dlibrary](#), [dllcall](#)

Case 25: Temporary File Path

Purpose

Gets or sets the path **GAUSS** will use for temporary files.

Format

```
oldpath = sysstate(25, path);
```

Input

path scalar 0 to get path, or string containing the new path.

Output

oldpath string, original path.

Remarks

If *path* is of type matrix, the path will be returned but not modified.

Case 26: Interface Mode

Purpose

Returns the current interface mode.

Format

```
mode = sysstate(26,0);
```

Output

<i>mode</i>	scalar, interface mode flag
0	non-X mode
1	terminal (-v) mode
2	X Windows mode

sysstate

Remarks

A mode of 0 indicates that you're running a non-X version of **GAUSS**; i.e., a version that has no X Windows capabilities. A mode of 1 indicates that you're running an X Windows version of **GAUSS**, but in terminal mode; i.e., you started **GAUSS** with the -v flag. A mode of 2 indicates that you're running **GAUSS** in X Windows mode.

Case 28: Random Number Generator Parameters

Purpose

Gets or sets the random number generator (RNG) parameters.

Format

oldprms = **sysstate**(28, *prms*);

Input

<i>prms</i>	scalar 0 to get parameters, or 3x1 matrix of new parameters.
[1]	seed, $0 < \text{seed} < 2^{32}$
[2]	multiplier, $0 < \text{mult} < 2^{32}$
[3]	constant, $0 \leq \text{const} < 2^{32}$

Output

<i>oldprms</i>	3x1 vector, the original parameters.
----------------	--------------------------------------

Remarks

If *prms* is a scalar 0, the current parameters will be returned without being changed.

The modulus of the RNG cannot be changed; it is fixed at 2^{32} .

See Also

[rndcon](#), [rndmult](#), [rndseed](#), [rndn](#), [rndu](#)

Case 30: Base Year Toggle

Purpose

Specifies whether year value returned by **date** is to include base year (1900) or not.

Format

```
oldstate = sysstate(30, state);
```

Input

state scalar, 1, 0, or missing value.

Output

oldstate scalar, the original state.

Remarks

Internally, **date** acquires the number of years since 1900. **sysstate** case 30 specifies whether **date** should add the base year to that value or not. If *state* = 1, **date** adds 1900, returning a fully-qualified 4-digit year.

If *state* = 0, **date** returns the number of years since 1900. If *state* is a missing value, the current state is returned. The default state is 1.

sysstate

Case 32: Global LU Tolerance

Purpose

Gets or sets global singularity tolerance for LU decomposition.

Format

```
oldtol = sysstate(32, tol);
```

Input

tol scalar, new tolerance.

Output

oldtol scalar, the original tolerance.

Remarks

The tolerance must be ≥ 0 . If *tol* is negative, the tolerance is returned and left unchanged.

This is a global tolerance and therefore not thread-safe. To set the singularity tolerance for LU decomposition in the current thread, use case 13.

See Also

[croutp](#), [inv](#)

Case 33: Global Cholesky Tolerance

Purpose

Gets or sets global singularity tolerance for Cholesky decomposition.

Format

```
oldtol = sysstate(33, tol);
```

Input

<i>tol</i>	scalar, new tolerance.
------------	------------------------

Output

<i>oldtol</i>	scalar, the original tolerance.
---------------	---------------------------------

Remarks

The tolerance must be ≥ 0 . If *tol* is negative, the tolerance is returned and left unchanged.

This is a global tolerance and therefore not thread-safe. To set the singularity tolerance for Cholesky decomposition in the current thread, use case 14.

This affects the following functions:

solpd

invpd for matrices $\leq 12 \times 12$

See Also

[chol](#), [invpd](#), [solpd](#)

Case 34: Global Imaginary Tolerance

Purpose

Gets or sets the global imaginary tolerance.

system

Format

```
oldtol = sysstate(34, tol);
```

Input

<i>tol</i>	scalar, the new tolerance.
------------	----------------------------

Output

<i>oldtol</i>	scalar, the original tolerance.
---------------	---------------------------------

Remarks

The imaginary tolerance is used to test whether the imaginary part of a complex matrix can be treated as zero or not. Functions that are not defined for complex matrices check the imaginary part to see if it can be ignored. The default tolerance is 2.23e-16, or machine epsilon.

If $tol < 0$, the current tolerance is returned.

This is a global tolerance and therefore not thread-safe. To set the imaginary tolerance in the current thread, use case 21.

See Also

[hasimag](#)

system

Purpose

Quits **GAUSS** and returns to the operating system.

Format

```
system;  
system c;
```

Input

<i>c</i>	scalar, an optional exit code that can be recovered by the program that invoked GAUSS . The default is 0. Valid arguments are 0-255.
----------	---

Remarks

The `system` command always returns an exit code to the operating system or invoking program. If you don't supply one, it returns 0. This is usually interpreted as indicating success.

See Also

[exec](#)

tab

t

tab

Purpose

Tabs the cursor to a specified text column.

Format

```
tab(col);  
print expr1 expr2 tab(col1) expr3 tab(col2) expr4 ...;
```

Input

<i>col</i>	scalar, the column position to tab to.
------------	--

Remarks

col specifies an absolute column position. If *col* is not an integer, it will be truncated.

tab can be called alone or embedded in a **print** statement. You cannot embed it within a parenthesized expression in a **print** statement, though. For example:

```
print (tab(20) c + d * e);
```

will not give the results you expect. If you have to use parenthesized expressions, write it like this instead:

```
print tab(20) (c + d * e);
```


tan

Purpose

Returns the tangent of its argument.

Format

```
 $y = \mathbf{tan}(x);$ 
```

Input

x	NxK matrix or N-dimensional array.
-----	------------------------------------

Output

y	NxK matrix or N-dimensional array.
-----	------------------------------------

Remarks

For real matrices, x should contain angles measured in radians.

To convert degrees to radians, multiply the degrees by $\pi/180$.

Example

```
//Create an additive sequence 0.1, 0.2, 0.3...0.9  
x = seqa(0.1, 0.1, 9);  
  
y = tan(x);
```

The above code produces:

tanh

```
0.1003346
0.2027100
0.3093362
0.4227932
y = 0.5463024
0.6841368
0.8422883
1.0296386
1.2601582
```

See Also

[atan](#), [pi](#)

tanh

Purpose

Computes the hyperbolic tangent.

Format

$y = \text{tanh}(x);$

Input

x NxK matrix or N-dimensional array.

Output

y NxK matrix or N-dimensional array containing the hyperbolic tangents of the elements of x .

Example

```
//Create a sequence starting at -0.5 and increasing by  
//0.25, i.e. -0.5, -0.25, 0, 0.25...1  
x = seqa(-0.5, 0.25, 7);  
x = x * pi;  
y = tanh(x);
```

After the above code, *y* is equal to:

```
-0.46211716  
-0.24491866  
0.00000000  
0.24491866  
0.46211716  
0.63514895  
0.76159416
```

Source

trig.src

tempname

Purpose

Creates a temporary file with a unique name.

Format

```
tname = tempname(path, pre, suf);
```

ThreadBegin

Input

<i>path</i>	string, path where the file will reside.
<i>pre</i>	string, a prefix to begin the file name with.
<i>suf</i>	string, a suffix to end the file name with.

Output

<i>tname</i>	string, unique temporary file name of the form <i>path/preXXXXnnnnnsuf</i> , where XXXX are 4 letters, and nnnnn is the process id of the calling process.
--------------	--

Remarks

Any or all of the inputs may be a null string or 0. If *path* is not specified, the current working directory is used.

If unable to create a unique file name of the form requested, **tempname** returns a null string.

WARNING: **GAUSS** does not remove temporary files created by **tempname**. It is left to the user to remove them when they are no longer needed.

ThreadBegin

Purpose

Marks the beginning of a multi-line block of code to be executed as a thread.

Format

```
ThreadBegin;
```

Example

```
ThreadBegin;  
    m = n*p;  
    n = calcA(m) ;  
ThreadEnd;
```

Notice that the **writer-must-isolate** rule (see **MULTI-THREADED PROGRAMMING IN GAUSS**, CHAPTER 1) does not apply within the bounds of the [ThreadBegin/ThreadEnd](#) pair, as there is no risk of simultaneous access to a symbol. The rule only applies between the threads in a given set (and their children).

See [ThreadJoin](#) for an example of a fully-defined thread set.

See Also

[ThreadEnd](#), [ThreadJoin](#), [ThreadStat](#)

ThreadEnd

Purpose

Marks the end of a multi-line block of code to be executed as a thread.

Format

```
ThreadEnd;
```

Example

```
ThreadBegin;  
    m = n*p;  
    n = calcA(m) ;  
ThreadEnd;
```

threadfor, threadendfor

Notice that the **writer-must-isolate** rule (see **MULTI-THREADED PROGRAMMING IN GAUSS**, CHAPTER 1) does not apply within the bounds of the [ThreadBegin/ThreadEnd](#) pair, as there is no risk of simultaneous access to a symbol. The rule only applies between the threads in a given set (and their children). See [ThreadJoin](#) for an example of a fully-defined thread set.

See Also

[ThreadBegin](#), [ThreadJoin](#), [ThreadStat](#)

threadfor, threadendfor

Purpose

Begins a parallel [for](#) loop.

Format

```
threadfor i(start, stop, step);  
.  
.  
.  
threadendfor;
```

Input

<i>i</i>	literal, the name of the counter variable.
<i>start</i>	scalar expression, the initial value of the counter.
<i>stop</i>	scalar expression, the final value of the counter.
<i>step</i>	scalar expression, the increment value.

Remarks

1. The iterations of a `threadfor` loop may execute in any order.
2. Indexed assignments to global variables that use the loop counter behave the same as in a standard `for` loop.
3. Non-indexed assignments will create a temporary variable that persists only through the remainder of the current loop iteration. For example:

```
a = 34.7;
threadfor i(1, 2, 1);
    a = randu(1,1);
    print a;
threadEndfor;

print a;
```

will produce output similar to the following:

```
0.90560157
0.52594285
34.700000
```

4. `threadfor` loops may not be nested

Examples

Example 1

```
//A basic 'threadfor' loop
threadfor i (1, 4, 1);
    print i;
threadendfor;
```

The code above, will print out:

```
1.000
2.000
```

threadfor, threadendfor

3.000
4.000

Example 2: Simple bootstrap of the mean of one variable

```
//Create fully pathed name of dataset
dataset = getGAUSSHome() $+ "examples/fueleconomy.dat";

//Load all contents of dataset
x = loadadd(dataset);

//Extract 2nd column
engine_disp = x[:,2];

iters = 500;
nobs = rows(engine_disp);

//Pre-allocate vector to hold sample means
sample_means = zeros(iters, 1);

threadFor i(1, iters, 1);
    //Create tmp variable 'idx',
    //containing random integers from 1-nobs
    //'idx' exists ONLY during the loop
    idx = ceil(nobs * rndu(nobs, 1));

    //Extract random sample into tmp variable,
    //'sample'. Only exists during loop
    sample = engine_disp[idx];

    //Calculate mean of sample
    //and assign using loop counter
    //'sample_means' will persist after loop
    sample_means[i] = meanc(sample);
threadEndFor;
```


See Also

[Performance considerations](#)

ThreadJoin

Purpose

Completes the definition of a set of threads to be executed simultaneously.

Format

```
ThreadJoin;
```

Remarks

Each thread in the set must adhere to the **writer-must-isolate** rule (see **MULTI-THREADED PROGRAMMING IN GAUSS, CHAPTER 1**). Because the threads in a set execute simultaneously, there is no way of knowing in one thread the current "state" of a symbol in another, and thus no way of safely or meaningfully accessing it.

Example

```
ThreadBegin;           //Thread 1--isolates y,z
  y = x'x;
  z = y'y;
ThreadEnd;
ThreadBegin;           //Thread 2--isolates q,r
  q = r'r;
  r = q'q;
ThreadEnd;
ThreadStat n = m'm;    //Thread 3--isolates n
ThreadStat p = o'o;    //Thread 4--isolates p
ThreadJoin;            //Joins threads 1-4
```

ThreadStat

```
b = z + r + n*p;    //y,z,q,r,n,p available again,  
                    // can be read and written
```

Note how threads 1-4 isolate the various symbols they assign to--no other thread references the written symbols at all. Once the threads are joined, however, the symbols are again available for use, and can be both read and assigned to.

See Also

[ThreadBegin](#), [ThreadEnd](#), [ThreadStat](#)

ThreadStat

Purpose

Marks a single line of code to be executed as a thread.

Format

```
ThreadStat statement;
```

Example

```
ThreadStat m = n*p;
```

See [ThreadJoin](#) for an example of a fully-defined thread set.

See Also

[ThreadBegin](#), [ThreadEnd](#), [ThreadJoin](#)

time

Purpose

Returns the current system time.

Format

```
y = time;
```

Output

<i>y</i>	4x1 numeric vector, the current time in the order: hours, minutes, seconds, and hundredths of a second.
----------	---

Example

```
print time;
```

```
7.000000  
31.000000  
46.000000  
33.000000
```

See Also

[date](#), [datestr](#), [datestring](#), [datestrymd](#), [hsec](#), [timestr](#)

timedt

Purpose

Returns system date and time in DT scalar format.

timestr

Format

```
dt = timedt;
```

Output

<i>dt</i>	scalar, system date and time in DT scalar format.
-----------	---

Remarks

The DT scalar format is a double precision representation of the date and time. In the DT scalar format, the number:

```
20100306071511
```

represents:

```
07:15:11 or 7:15:11 AM on March 6, 2010.
```

Source

time.src

See Also

[todaydt](#), [timeutc](#), [dtdate](#)

timestr

Purpose

Formats a time in a vector to a string.

Format

```
ts = timestr(t);
```

Input

t	4x1 vector from the time function, or a zero. If the input is 0, the time function will be called to return the current system time.
-----	--

Output

ts	8 character string containing current time in the format: hr:mn:sc
------	--

Example

```
t = { 7, 31, 46, 33 };  
ts = timestr(t);  
print ts;
```

produces:

```
7:31:46
```

Source

time.src

See Also

[date](#), [datestr](#), [datestring](#), [datestrymd](#), [ethsec](#), [etstr](#), [time](#)

timeutc

Purpose

Returns the number of seconds since January 1, 1970 Greenwich Mean Time.

title

Format

```
tc = timeutc;
```

Output

<i>tc</i>	scalar, number of seconds since January 1, 1970 Greenwich Mean Time.
-----------	--

Example

```
//Retrieve seconds since January 1, 1970 GMT
tc = timeutc;

//Convert to a date time vector
utv = utctodtv(tc);
```

After the code above, *tc* and *utv* are equal to:

```
tc = 1340080112

utv = 2012 06 18 21 28 32 1 169
```

See Also

[dtvnormal](#), [utctodtv](#)

title

Purpose

Sets the title for the graph. NOTE: This function is for the deprecated PQG graphics. Use **plotSetTitle** instead.

Library

pgraph

Format

```
title(str);
```

Input

<i>str</i>	string, the title to display above the graph.
------------	---

Remarks

Up to three lines of title may be produced by embedding a line feed character ("`\L`") in the title string.

Example

```
title("First title line\LSecond title line\L"\  
      "Third title line");
```

Fonts may be specified in the title string. For instructions on using fonts, see **Selecting Fonts**, Section 1.0.1.

Source

pgraph.src

See Also

[xlabel](#), [ylabel](#), [fonts](#)

tkf2eps

tkf2eps

Purpose

Converts a `.tkf` file to an Encapsulated PostScript file. NOTE: This function is deprecated and does not work for the new `.plot` graphics files. Use **plotSave** to convert `.plot` files to EPS format.

Library

pgraph

Format

```
ret = tkf2eps(tekfile, epsfile);
```

Input

<i>tekfile</i>	string, name of <code>.tkf</code> file.
<i>epsfile</i>	string, name of Encapsulated PostScript file.

Output

<i>ret</i>	scalar, 0 if successful
------------	-------------------------

Remarks

The conversion is done using the global parameters in `peps.dec`. You can modify these globally by editing the `.dec` file, or locally by setting them in your program before calling **tkf2eps**.

See the header of the output Encapsulated PostScript file and a PostScript manual if you want to modify these parameters.

tkf2ps

Purpose

Converts a `.tkf` file to a PostScript file. NOTE: This function is deprecated and does not work for the new `.plot` graphics files. Use **plotSave** to convert `.plot` files to PS format.

Library

pgraph

Format

```
ret = tkf2ps(tekfile, psfile);
```

Input

<i>tekfile</i>	string, name of <code>.tkf</code> file.
<i>psfile</i>	string, name of PostScript file.

Output

<i>ret</i>	scalar, 0 if successful.
------------	--------------------------

Remarks

The conversion is done using the global parameters in `peps.dec`. You can modify these globally by editing the `.dec` file, or locally by setting them in your program before calling **tkf2ps**.

See the header of the output PostScript file and a PostScript manual if you want to modify these parameters.

tocart

tocart

Purpose

Converts from polar to Cartesian coordinates.

Format

$xy = \text{tocart}(r, \theta);$

Input

r	$N \times K$ real matrix, radius.
θ	$L \times M$ real matrix, $E \times E$ conformable with r , angle in radians.

Output

xy	$\max(N,L)$ by $\max(K,M)$ complex matrix containing the x coordinate in the real part and the y coordinate in the imaginary part.
------	--

Source

coord.src

todaydt

Purpose

Returns system date in DT scalar format. The time returned is always midnight (00:00:00), the beginning of the returned day.

Format

```
dt = todaydt;
```

Output

<i>dt</i>	scalar, system date in DT scalar format.
-----------	--

Remarks

The DT scalar format is a double precision representation of the date and time. In the DT scalar format, the number:

```
20120906130525
```

represents 13:05:25 or 1:05:25 PM on September 6, 2012.

Source

time.src

See Also

[timedt](#), [timeutc](#), [dtdate](#)

toeplitz

Purpose

Creates a Toeplitz matrix from a column vector.

Format

```
t = toeplitz(x);
```

token

Input

x $K \times 1$ vector.

Output

t $K \times K$ Toeplitz matrix.

Example

```
//Create the sequence 1, 2, 3, 4, 5 and assign it to 'x'
x = seqa(1,1,5);

//Create a diagonal-constant or Toeplitz matrix
y = toeplitz(x);
```

After the code above, y is equal to:

```
1 2 3 4 5
2 1 2 3 4
3 2 1 2 3
4 3 2 1 2
5 4 2 2 3
```

Source

toeplitz.src

token

Purpose

Extracts the leading token from a string.

Format

```
{ token, str_left } = token(str);
```

Input

<i>str</i>	string, the string to parse.
------------	------------------------------

Output

<i>token</i>	string, the first token in <i>str</i> .
<i>str_left</i>	string, <i>str</i> minus <i>token</i> .

Remarks

str can be delimited with commas or spaces.

The advantage of **token** over **parse** is that **parse** is limited to tokens of 8 characters or less; **token** can extract tokens of any length.

Example

Here is a keyword that uses **token** to parse its string parameter:

```
//Create a keyword called 'add' that takes the input
// 's' and executes all of the code from the 'keyword
//add(s)' line until the 'endp' statement each time
//it is called
keyword add(s);
    local tok,sum;
    sum = 0;

    //Continue loop until 's' equals an empty string
```

token

```
do until s $== "";

    //Remove the first token from 's' and return
    //it in 'tok'
    { tok, s } = token(s);

    //Convert the string in 'tok' to a floating
    //point number and add it to 'sum'
    sum = sum + stof(tok);
enddo;

//Set the formatting for print statements to
//create 1 space between numbers and
//to print 2 digits after the decimal point
format /rd 1,2;
print "Sum is: " sum;
endp;
```

If you type:

```
//Since it is a 'keyword' and not a 'proc', 'add'
//will take everything between 'add' and the
//semi-colon as a string input and refer to it
//internally as the 's' variable
add 1 2 3 4 5 6;
```

add will respond:

```
Sum is: 15.00
```

Source

token.src

See Also

[parse](#)

topolar

Purpose

Converts from Cartesian to polar coordinates.

Format

```
{ r, theta } = topolar(xy);
```

Input

<i>xy</i>	NxK complex matrix containing the <i>x</i> coordinate in the real part and the <i>y</i> coordinate in the imaginary part.
-----------	---

Output

<i>r</i>	NxK real matrix, radius.
<i>theta</i>	NxK real matrix, angle in radians.

Source

coord.src

trace

Purpose

Allows the user to trace program execution for debugging purposes.

trace

Format

```
trace new;  
trace new, mask;
```

Input

<i>new</i>	scalar, new value for trace flag.
<i>mask</i>	scalar, optional mask to allow leaving some bits of the trace flag unchanged.

Remarks

The `trace` command has no effect unless you are running your program under GAUSS's source level debugger. Setting the `trace` flag will not generate any debugging output during normal execution of a program.

The argument is converted to a binary integer with the following meanings:

bit	decimal	meaning
ones	1	trace calls/returns
twos	2	trace line numbers
fours	4	unused
eights	8	output to window
sixteens	16	output to print
thirty-twos	32	output to auxiliary output
sixty-fours	64	output to error log

You must set one or more of the output bits to get any output from `trace`. If you set `trace` to 2, you'll be doing a line number trace of your program, but the output will not be displayed anywhere.

The `trace` output as a program executes will be as follows:

(+GRAD)	calling function or procedure GRAD
(-GRAD)	returning from GRAD
[47]	executing line 47

Note that the line number trace will only produce output if the program was compiled with line number records.

To set a single bit use two arguments:

<code>trace 16,16;</code>	turn on output to printer
<code>trace 0,16;</code>	turn off output to printer

Example

```
trace 1+8;      //trace fn/proc calls/returns to standard
                //output
trace 2+8;      //trace line numbers to standard output
trace 1+2+8;    //trace line numbers and fn/proc calls/
                //returns to standard output
trace 1+16;     //trace fn/proc calls/returns to printer
trace 2+16;     //trace line numbers to printer
trace 1+2+16;   //trace line numbers and fn/proc calls/
                //returns to printer
```

See Also

[lineson](#)

trap

Purpose

Sets the trap flag to enable or disable trapping of numerical errors.

trap

Format

```
trap val;  
trap val, mask;
```

Input

<i>val</i>	scalar, new trap value.
<i>mask</i>	scalar, optional mask to allow leaving some bits of the trap flag unchanged.

Examples

Example 1

First we will create some code that will return an error, stopping the program. Then we will show how to trap and handle the error:

```
//Create a singular matrix  
x = { 1 1 1,  
      1 1 1,  
      1 1 1 };  
  
//Attempt to calculate inverse, but error  
//'matrix singular' stops program  
x_inv = inv(x);
```

In some cases, we would like our program to be able to detect certain errors and recover from them. The next section of code will be the same as above, with the exception of setting the `trap` flag. We will see that it will not cause an error.

```
//Create a singular matrix  
x = { 1 1 1,  
      1 1 1,  
      1 1 1 };
```

```
//Set the trap flag, to suppress the error
trap 1;

//Attempt to calculate inverse
x_inv = inv(x);
```

If you run the above code, you will notice that an error was not returned. With the trap set to 1, instead of stopping the program with an error message, **GAUSS** will set the variable `x_inv` equal to a scalar error code. A scalar error code is a missing value that contains an integer which can be used to identify the error. For more information on error codes, see **scalerr** and **error**. In this example, however, our main concern is with determining whether or not the return value is a scalar error code. We can do this with the **GAUSS** function, **scalmiss**.

```
//Create a singular matrix
x = { 1 1 1,
      1 1 1,
      1 1 1 };

//Set the trap flag, to suppress the error
trap 1;

//Attempt to calculate inverse
x_inv = inv(x);

//Check to see if 'x_inv', contains a scalar error code
if scalmiss(x_inv);
    print "matrix was singular";
endif;
```

Example 2

This example will build from the concepts in the example above to do something more useful.

trap

```
//Create a coefficient matrix with linear dependencies
x = { 1 1 0.8,
      1 1 1.5,
      1 1 0.6 };

//Create a dependent variable
y = { -0.36,
      -1.55,
      -0.02 };

//Set the trap flag, to suppress the error
trap 1;

//Attempt to compute the inverse of the moment matrix
mmi = inv(x'x);

//Check to see if 'mmi', contains a scalar error code
if scalmiss(mmi);
    //Compute the pseudo-inverse of the moment matrix
    mmi = pinv(x'x);
endif;

//Solve the linear equations
b_hat = mmi * x'y;
```

After the above code, *b_hat* is equal to:

```
0.5
0.5
-1.7
```

Remarks

The trap flag is examined by some functions to control error handling. There are 16 bits in the trap flag, but most **GAUSS** functions will examine only the lowest order bit:

trap 1;	turn trapping on
trap 0;	turn trapping off

If we extend the use of the trap flag, we will use the lower order bits of the trap flag. It would be wise for you to use the highest 8 bits of the trap flag if you create some sort of user-defined trap mechanism for use in your programs. (See the function **trapchk** for detailed instructions on testing the state of the trap flag; see **error** for generating user-defined error codes.)

To set only one bit and leave the others unchanged, use two arguments:

trap 1,1;	set the ones bit
trap 0,1;	clear the ones bit

See Also

[scalerr](#), [trapchk](#), [error](#)

trapchk

Purpose

Tests the value of the trap flag.

Format

$y = \text{trapchk}(m);$

Input

m	scalar mask value.
-----	--------------------

trapchk

Output

<i>y</i>	scalar which is the result of the bitwise logical AND of the trap flag and the mask value.
----------	--

Remarks

To check the various bits in the trap flag, add the decimal values for the bits you wish to check according to the chart below and pass the sum in as the argument to the trapchk function:		
	bit	decimal value
	0	1
	1	2
	2	4
	3	8
	4	16
	5	32
	6	64
	7	128
	8	256
	9	512
	10	1024
	11	2048
	12	4096
	13	8192
	14	16384
	15	32768

If you want to test if either bit 0 or bit 8 is set, then pass an argument of 1+256 or 257 to **trapchk**. The following table demonstrates values that will be returned for:

```
y = trapchk(257);
```

	0	1	value of bit 0 in trap flag
0	0	1	
1	256	257	
value of bit 8 in trap flag			

GAUSS functions that test the trap flag currently test only bits 0 and 1.

See Also

[scalerr](#), [trap](#), [error](#)

trigamma

Purpose

Computes trigamma function.

Format

```
y = trigamma(x);
```

Input

x	MxN matrix or N-dimensional array.
---	------------------------------------

trimr

Output

y	MxN matrix or N-dimensional array, trigamma.
-----	--

Remarks

The trigamma function is the second derivative of the log of the gamma function with respect to its argument.

trimr

Purpose

Trims rows from the top and/or bottom of a matrix.

Format

$y = \text{trimr}(x, \ t, \ b);$

Input

x	NxK matrix from which rows are to be trimmed.
t	scalar containing the number of rows which are to be removed from the top of x .
b	scalar containing the number of rows which are to be removed from the bottom of x .

Output

y	RxK matrix where $R=N-(t + b)$, containing the rows left after the trim.
-----	---

Remarks

If either t or b is zero, then no rows will be trimmed from that end of the matrix.

Example

```
//Create a 5x3 matrix of random uniform numbers
x = randu(5,3);

//Remove the top 2 rows of x and the bottom row
y = trimr(x,2,1);
```

If x is equal to:

```
0.780 0.922 0.864
0.151 0.687 0.947
0.271 0.014 0.060
0.054 0.084 0.526
0.880 0.278 0.199
```

then y will equal:

```
0.271 0.014 0.060
0.054 0.084 0.526
```

See Also

[submat](#), [rotater](#), [shiftr](#)

trunc

Purpose

Converts numbers to integers by truncating the fractional portion.

trunc

Format

```
y = trunc(x);
```

Input

x	NxK matrix or N-dimensional array.
---	------------------------------------

Output

y	NxK matrix or N-dimensional array containing the truncated elements of x.
---	---

Example

```
x = 100*rndn(2,2);  
y = trunc(x);
```

If x equals:

```
-153.373  -1.972  
109.412  127.732
```

then, y will equal:

```
-153.000  -1.000  
109.000  127.000
```

See Also

[ceil](#), [floor](#), [round](#)

type

Purpose

Returns the symbol table type (matrix, string, etc) of its input argument.

Format

$t = \text{type}(x);$

Input

x	local or global symbol, can be an expression.
-----	---

Output

t	scalar, argument type.
6	matrix
13	string
15	string array
17	structure
21	array
23	structure pointer
23	sparse matrix

Example

Example 1

//Create a matrix
x = { 1 2,

type

```
3 4 };  
  
//Find type of 'x'  
x_type = type(x);
```

After the above code, `x_type` will equal: 6, indicating that `x` is a matrix.

Example 2

```
//Create a string  
x = "myfile.dat";  
  
//Find type of 'x'  
x_type = type(x);
```

After the above code, `x_type` will equal: 13, indicating that `x` is a string.

Remarks

type is often used to verify that inputs to a user defined procedure are valued. For example, if an input is a file name, then it must be a string:

```
proc (1) = myProc(fname);  
    if type(fname) != 13;  
        errorlog "Input 'fname' must be a string";  
    end;  
endif;  
endp;
```

type returns the type of a single symbol. The related function **typecv** will take a character vector of symbol names and return a vector of either their types or the missing value code for any that are undefined. **type** works for the symbol types listed above; **typecv** works for user-defined procedures, keywords and functions as well. **type** works for global or local symbols; **typecv** works only for global symbols.

See Also

[typecv](#), [typeof](#)

typecv

Purpose

Returns the symbol table type of objects whose names are given as a string or as elements of a character vector or string array.

Format

```
y = typecv(x);
```

Input

<i>x</i>	string, or Nx1 character vector or string array which contains the names of variables whose type is to be determined.
----------	---

Output

<i>y</i>	scalar or Nx1 vector containing the types of the respective symbols in <i>x</i> .
----------	---

Remarks

The values returned by **typecv** for the various variable types are as follows:

5	keyword (keyword)
---	-------------------------------------

typeof

6	matrix (numeric, character, or mixed)
8	procedure (<code>proc</code>)
9	function (<code>fn</code>)
13	string
15	string array
17	structure
21	array
23	structure pointer

typeof will return the **GAUSS** missing value code if the symbol is not found, so it may be used to determine if a symbol is defined or not.

Example

```
xvar = sqrt(5);  
yvar = "betahat";  
fn area(r) = pi*r*r;  
let names = xvar yvar area;  
y = typeof(names);
```

This code assigns the following to `y`:

```
6 //6 for type matrix  
y = 13 //13 for string  
9 //9 for function
```

See Also

[type](#), [typeof](#), [varput](#), [varget](#)

typedef

Purpose

Returns the type of data (the number of bytes per element) in a **GAUSS** data set.

Format

```
y = typedef(fp);
```

Input

<i>fp</i>	scalar, file handle of an open file.
-----------	--------------------------------------

Output

<i>y</i>	scalar, type of data in GAUSS data set.
----------	--

Remarks

If *fp* is a valid **GAUSS** file handle, then *y* will be set to the type of the data in the file as follows:

2	2-byte signed integer
4	4-byte IEEE floating point
8	8-byte IEEE floating point

Example

```
//Assign a variable to represent each of our file names  
infile = "dat1";
```

typedef

```
outfile = "dat2";

//Open the file "dat1" for reading.
//Note: The ^ before 'infile' tells GAUSS to use the value
//of the string variable 'infile' (which is 'dat1' in this
//case) rather than name of the variable.
open fin = ^infile;

//Get the names of the variables that are saved in the
//dataset
names = getname(infile);

//Create a new data set file using the same variable names
//as 'dat1', with 1 column per data element and using the
//same size data, i.e. the number of bytes per element, as
//the data in 'dat1'
create fout = ^outfile with ^names, 0, typedef(fin);
```

In this example, a file `dat2.dat` is created which has the same variables and variable type as the input file, `dat1.dat`. **typedef** is used to return the type of the input file data for the `create` statement.

See Also

[colsf](#), [rowsf](#)

u

union

Purpose

Returns the union of two vectors with duplicates removed.

Format

```
y = union(v1, v2, flag);
```

Input

<i>v1</i>	Nx1 vector.
<i>v2</i>	Mx1 vector.
<i>flag</i>	scalar, 1 if numeric data, 0 if character.

Output

<i>y</i>	Lx1 vector containing all unique values that are in <i>v1</i> and <i>v2</i> , sorted in ascending order.
----------	--

Remarks

The combined elements of *v1* and *v2* must fit into a single vector.

Example

```
//Create two column vectors with character data
```

unionsa

```
let v1 = mary jane linda john;
let v2 = mary sally;

x = union(v1,v2,0);

//The '$' in front of 'x' tells GAUSS to print 'x' as
//character data
print $x;
```

The above code will produce the following results:

```
JANE
JOHN
LINDA
MARY
SALLY
```

unionsa

Purpose

Returns the union of two string vectors with duplicates removed.

Format

```
y = unionsa(sv1, sv2);
```

Input

<i>sv1</i>	Nx1 or 1xN string vector.
<i>sv2</i>	Mx1 or 1xM string vector.

Output

y Lx1 vector containing all unique values that are in *sv1* and *sv2*, sorted in ascending order.

Example

```
string sv1 = { "mary", "jane", "linda", "john" };
string sv2 = { "mary", "sally" };
y = unionsa(sv1,sv2);
print y;
```

The above code produces the following output:

```
jane
john
linda
mary
sally
```

Source

unionsa.src

See Also

[union](#)

uniqindx

Purpose

Computes the sorted index of *x*, leaving out duplicate elements.

uniqindx

Format

```
index = uniqindx(x, flag);
```

Input

<i>x</i>	Nx1 or 1xN vector.
<i>flag</i>	scalar, 1 if numeric data, 0 if character.

Output

<i>index</i>	Mx1 vector, indices corresponding to the elements of <i>x</i> sorted in ascending order with duplicates removed.
--------------	--

Remarks

Among sets of duplicates it is unpredictable which elements will be indexed.

Example

```
let x = 5 4 4 3 3 2 1;  
  
//Create a sorted index of all the unique elements in 'x'  
ind = uniqindx(x,1);  
  
//Use the index 'ind' to return all of the unique elements  
//of 'x' in ascending order  
y = x[ind];
```

After running the above code, *ind* and *y* are equal to:

	7.0000000	1.0000000
	6.0000000	2.0000000
ind =	4.0000000	y = 3.0000000

3.0000000	4.0000000
1.0000000	5.0000000

See Also

[unique](#), [uniqindxsa](#)

uniqindxsa

Purpose

Computes the sorted index of a string vector, omitting duplicate elements.

Format

```
ind = uniqindxsa(sv);
```

Input

<i>sv</i>	Nx1 or 1xN string vector.
-----------	---------------------------

Output

<i>ind</i>	Mx1 vector, indices corresponding to the elements of <i>sv</i> sorted in ascending order with duplicates removed.
------------	---

Remarks

Among sets of duplicates it is unpredictable which elements will be indexed.

Example

```
string sv = {"mary", "linda", "linda", "jane",
```

unique

```
                "jane", "cindy", "betty"};
ind = uniqueidxsa(sv);
y = sv[ind];
```

The above code assigns the variables *ind* and *y* as follows:

```

          7      betty
          6      cindy
ind  = 4   y =  jane
          2      linda
          1      mary
```

Source

`uniquesa.src`

See Also

[unique](#), [uniquesa](#), [uniqueidx](#)

unique

Purpose

Sorts and removes duplicate elements from a vector.

Format

```
y = unique(x);
y = unique(x, flag);
```

Input

<i>x</i>	NxM numeric matrix, NxM character data, or NxM string array.
----------	--

flag

Optional input, scalar, 1 if numeric data, 0 if character data. Default is 1. String array does not need a flag.

Output

y

Mx1 vector, sorted *x* with the duplicates removed.

Example

Example 1: Numeric

```
//Create a column vector with duplicate elements
years = { 1632,
          2012,
          1709,
          1812,
          1709,
          1989,
          1830,
          1875,
          1912,
          1912,
          1924,
          1960 };

//Sort 'years' and remove any duplicate elements
years_unique = unique(years);
```

After the code above, the variables *years* and *years_unique* are assigned as follows:

1632	
2012	1632

unique

```

        1709
        1812
        1709
years = 1989    years_unique = 1875
        1830
        1875
        1912
        1912
        1924
        1960
        1709
        1812
        1830
        1912
        1924
        2012
```

Example 2: Numeric Matrix

```
//Create a numeric matrix with duplicate elements
years = { 1632      2012,
        1709      1812,
        1709      1989,
        1830      1875,
        1912      1912,
        1924      1960 };

//Sort 'years' and remove any duplicate elements
years_unique = unique(years);
print "years: " years;
print;
print "years_unique:" years_unique;
```

After the code above, the variables *years* and *years_unique* are assigned as follows:

```
years:
1632.0000    2012.0000
1709.0000    1812.0000
1709.0000    1989.0000
```



```
1830.0000      1875.0000
1912.0000      1912.0000
1924.0000      1960.0000
```

```
years_unique:
1632.0000
1709.0000
1812.0000
1830.0000
1875.0000
1912.0000
1924.0000
1960.0000
1989.0000
2012.0000
```

Example 3: Character data

```
//Create column character vector, by using
//numeric concatenation operator
levels = "high" | "medium" | "medium" | "low" |
         "high" | "medium" | "medium";

//Set flag to indicate data is character data
flag = 0;

//Sort 'levels' alphabetically and
//remove any duplicate elements
levels_unique = unique(levels, flag);

//Note the $ used before the variable which
```

R

unique

```
//tells GAUSS to print as characters  
print $levels_unique;
```

The code above will produce the following output:

```
      high  
      low  
    medium
```

You can reorder these levels with an indexing operation, for example:

```
levels = levels_unique[2 3 1];  
print levels;
```

will produce the following output:

```
      low  
    medium  
      high
```

Example 4 : String array vector

```
//Create column string array  
string levels = { "high", "medium", "medium" , "low" ,  
                  "high" , "medium" , "medium"};  
  
//Sort 'levels' alphabetically and  
//remove any duplicate elements  
levels_unique = unique(levels);  
  
print levels_unique;
```

The code above will produce the following output:

```

    high
    low
    medium

```

Example 5 : String array matrix

```

//Create 3x2 string array

string levels = { "apple"      "watermelon",
                  "banana"    "banana",
                  "watermelon" "apple" };

//Sort 'levels' alphabetically and
//remove any duplicate elements
levels_unique = unique(levels);

print"levels: " levels;
print;
print"levels_unique:" levels_unique;

```

The code above will produce the following output:

```

levels:
apple      watermelon
banana     banana
watermelon apple

levels_unique:
apple
banana
watermelon

```

See Also

[sortc](#), [uniquesa](#), [uniqindx](#)

uniquesa

uniquesa

Purpose

Removes duplicate elements from a string vector. NOTE: **uniquesa** is deprecated. Instances of this function should be replaced by **unique**.

Format

```
y = uniquesa(sv);
```

Input

<i>sv</i>	Nx1 or 1xN string vector.
-----------	---------------------------

Output

<i>y</i>	sorted Mx1 string vector containing all unique elements found in <i>sv</i> .
----------	--

Example

```
//Create a 8x1 string array
string comTrades = { "corn", "gold", "soybeans", "silver",
                    "coffee",
                    "oil", "silver", "soybeans" };

//Return an alphabetized string array containing the
//unique elements from 'comTrades'
commodity = uniquesa(comTrades);
```

After the code above, the variables *comTrades* and *commodity* will be equal to:

```

            corn
            gold
comTrades = soybeans
            silver
            coffee
            oil
            silver
            soybeans
commodity = coffee
            corn
            gold
            oil
            silver
            soybeans

```

Remarks

It is important to note that the return from **uniquesa** will always be a column vector, even if the input string array is a row vector.

Source

`uniquesa.src`

See Also

[unique](#), [unigindxsa](#), [unigindx](#)

upmat, upmat1

Purpose

Returns the upper portion of a matrix. **upmat** returns the main diagonal and every element above. **upmat1** is the same except it replaces the main diagonal with ones.

Format

```

u = upmat(x);
u = upmat1(x);

```

upmat, upmat1

Input

X $N \times K$ matrix.

Output

u NxK matrix containing the upper elements of *x*. The lower elements are replaced with zeros. **upmat** returns the main diagonal intact. **upmat1** replaces the main diagonal with ones.

Example

$$\mathbf{x} = \begin{Bmatrix} 7 & 2 & -1, \\ 2 & 3 & -2, \\ 4 & -2 & 8 \end{Bmatrix};$$

```
u = upmat(x);  
u1 = upmat1(x);
```

The resulting matrices are:

$$\mathbf{u} = \begin{bmatrix} 7 & 2 & -1 \\ 0 & 3 & -2 \\ 0 & 0 & 8 \end{bmatrix} \quad \mathbf{u}^{-1} = \begin{bmatrix} 1 & 2 & -1 \\ 0 & 1 & -2 \\ 0 & 0 & 1 \end{bmatrix}$$

Source

diag.src

See Also

[lowmat](#), [lowmat1](#), [diag](#), [diagrv](#), [crout](#)

upper

Purpose

Converts a string, matrix of character data, or string array to uppercase.

Format

```
y = upper(x);
```

Input

x	string, or NxK matrix, or string array containing the character data to be converted to uppercase.
---	--

Output

y	string, or NxK matrix, or string array containing the uppercase equivalent of the data in x.
---	--

Remarks

If x is a numeric matrix, y will contain garbage. No error message will be generated since GAUSS does not distinguish between numeric and character data in matrices.

Example

```
//Create a lowercase string
x = "uppercase";

//Convert the string to upper case
y = upper(x);

//Adding the '$' tells GAUSS to treat the data as character
```

use

```
//data  
print $y;
```

This code produces:

```
UPPERCASE
```

See Also

[lower](#)

use

Purpose

Loads a compiled file at the beginning of the compilation of a source program.

Format

```
use fname;
```

Input

<i>fname</i>	literal or ^string, the name of a compiled file created using the <code>compile</code> or the <code>saveall</code> command.
--------------	---

Remarks

The `use` command can be used ONCE at the TOP of a program to load in a compiled file which the rest of the program will be added to. In other words, if `xy.e` had the following lines:


```
library pgraph;
external proc xy;
x = seqa(0.1,0.1,100);
```

it could be compiled to `xy.gcg`. Then the following program could be run:

```
use xy;
xy(x, sin(x));
```

which would be equivalent to:

```
new;
library pgraph;
x = seqa(0.1,0.1,100);
xy(x, sin(x));
```

The `use` command can be used at the top of files that are to be compiled with the `compile` command. This can greatly shorten compile time for a set of closely related programs. For example:

```
library pgraph;
external proc xy,logx,logy,loglog,hist;
saveall pgraph;
```

This would create a file called `pgraph.gcg` containing all the procedures, strings and matrices needed to run PQG programs. Other programs could be compiled very quickly with the following statement at the top of each:

```
use pgraph;
```

or the same statement could be executed once, for instance from the command prompt, to instantly load all the procedures for PQG.

When the compiled file is loaded with `use`, all previous symbols and procedures are deleted before the program is loaded. It is therefore unnecessary to execute a `new` before `use`'ing a compiled file.

utctodt

`use` can appear only ONCE at the TOP of a program.

See Also

[compile](#), [run](#), [saveall](#)

utctodt

Purpose

Converts UTC scalar format to DT scalar format.

Format

```
dt = utctodt(utc);
```

Input

<i>utc</i>	Nx1 vector, UTC scalar format.
------------	--------------------------------

Output

<i>dt</i>	Nx1 vector, DT scalar format.
-----------	-------------------------------

Remarks

A UTC scalar gives the number of seconds since or before January 1, 1970 Greenwich Mean Time. In DT scalar format, 08:35:52 on June 11, 2005 is 20050611083552.

Example

```
tc = 1346290409;  
print "tc = " tc;
```

```
dt = utctodt(tc);  
print "dt = " dt;
```

produces:

```
tc = 1346290409  
dt = 20120829183329
```

Source

time.src

See Also

[dtvnormal](#), [timeutc](#), [utctodtv](#), [dttodtv](#), [dttvdt](#), [dttoutc](#), [dttvdt](#), [strtdt](#), [dttostr](#)

utctodtv

Purpose

Converts UTC scalar format to DTV vector format.

Format

```
dtv = utctodtv(utc);
```

Input

<i>utc</i>	Nx1 vector, UTC scalar format.
------------	--------------------------------

Output

<i>dtv</i>	Nx8 matrix, DTV vector format.
------------	--------------------------------

utctodtv

Remarks

A UTC scalar gives the number of seconds since or before January 1, 1970 Greenwich Mean Time.

Each row of `dtv`, in DTV vector format, contains:

<i>[N, 1]</i>	Year, four digit integer.
<i>[N, 2]</i>	Month in Year, 1-12.
<i>[N, 3]</i>	Day of month, 1-31.
<i>[N, 4]</i>	Hours since midnight, 0-23.
<i>[N, 5]</i>	Minutes, 0-59.
<i>[N, 6]</i>	Seconds, 0-59.
<i>[N, 7]</i>	Day of week, 0-6, 0=Sunday.
<i>[N, 8]</i>	Days since Jan 1 of current year, 0-365.

Example

```
//Set 'tc' equal to the number of seconds since January 1,
//1970
tc = timeutc;
print "tc = " tc;

dtv = utctodtv(tc);
print "dtv = " dtv;
```

produces:

```
tc = 1340315529
dtv = 2012 6 21 14 52 9 4 172
```

See Also

[dtvnormal](#), [timeutc](#), [utctodt](#), [dttodtv](#), [dttoutc](#), [dvtodt](#), [dvtoutc](#), [strtodt](#), [dttostr](#)

utrisol

Purpose

Computes the solution of $Ux = b$ where U is an upper triangular matrix.

Format

```
 $x = \text{utrisol}(b, U);$ 
```

Input

b	PxK matrix.
U	PxP upper triangular matrix.

Output

x	PxK matrix, solution of $Ux = b$.
-----	------------------------------------

Remarks

utrisol applies a back solve to $Ux = b$ to solve for x . If b has more than one column, each column is solved for separately, i.e., **utrisol** applies a back solve to $U * x[:,i] = b[:,i]$.

U

vals

v

vals

Purpose

Converts a string into a matrix of its ASCII values.

Format

```
y = vals(s);
```

Input

<i>s</i>	string of length N where $N > 0$.
----------	------------------------------------

Output

<i>y</i>	$N \times 1$ matrix containing the ASCII values of the characters in the string <i>s</i> .
----------	--

Remarks

If the string is null, the function will fail and an error message will be given.

Example

```
//Initialize 'k' so it will be 0 for the first iteration of  
//the 'do while' loop  
k = 0;
```

```
//Prompt the user for input
print "Continue Program? [Y/N]";

//Continually check for keyboard input and exit the loop on
//keyboard input
do while (k == 0);
    k = key;
enddo;

//Follow a different code branch depending upon which key
//the user entered
if k == vals("Y") or k == vals("y");
    print "You chose to continue";
else;
    print "Exiting program now";
endif;
```

In this example the **key** function is used to read keyboard input. When **key** returns a nonzero value, meaning a key has been pressed, the ASCII value it returns is tested to see if it is an uppercase or lowercase 'Y'. If it is, the program will follow the first branch and print:

```
You chose to continue
```

otherwise, it will follow the second branch and print:

```
Exiting program now
```

See Also

[chr](#), [ftos](#), [stof](#)

varCovM, varCovX

varCovM, varCovX

Purpose

Computes the population variance-covariance matrix.

Format

```
vc = varCovM(mm);  
vc = varCovX(x);
```

Input

<i>mm</i>	KxK moment ($x'x$) matrix. A constant term MUST have been the first variable when the moment matrix was computed.
<i>x</i>	NxK matrix of data.

Output

<i>vc</i>	KxK variance-covariance matrix.
-----------	---------------------------------

Example

```
//Set rndseed for repeatable random numbers  
rndseed 7234242;  
  
//Create three randomly generated independent variables  
x = rndn(500, 3);  
  
//Create the population variance-covariance matrix from  
data matrix 'x'  
var_x = varCovX(x);
```


After the code above, `var_x` will be equal to:

1.0941806	0.0040829	-0.0024871
0.0040829	1.0606611	0.0493555
-0.0024871	0.0493555	0.8729622

where the diagonal elements in the matrix represent the population variance of the each column, while the off-diagonal elements represent the population covariance between the data columns.

The population variance can also be calculated using the moment matrix, $x'x$ and the GAUSS function **varCovM**. A constant term must be included in the data matrix x when computing the moment equation. Consider the following data matrix $x1$, consisting of the original data matrix x and a column of ones:

```
//Set rndseed so 'rndn' will return the same numbers as
above
rndseed 7234242;

//Note: the ~ operator performs horizontal concatenation
x1 = ones(500,1)~rndn(500,3);

//Create moment matrix
x2 = x1'x1;

//Calculate variance-covariance matrix using the moment mat-
rix
var_xm = varCovM(x2);
```

After the code above, `var_xm` will be equal to:

1.0941806	0.0040829	-0.0024871
0.0040829	1.0606611	0.0493555
-0.0024871	0.0493555	0.8729622

varCovMS, varCovXS

Remarks

The variance covariance matrix is that of the population data matrix. It is computed as the moment matrix of deviations about the mean divided by the number of observations N . For a sample covariance matrix which uses $N - 1$ rather than N see **varCovMS** or **varCovXS**.

Source

corrs.src

See Also

[momentd](#), [corrms](#), [corrxs](#), [corrmm](#), [corrvc](#), [corrxx](#)

varCovMS, varCovXS

Purpose

Computes a sample variance-covariance matrix.

Format

```
VC = varCovMS(mm);
VC = varCovXS(x);
```

Input

>

<i>mm</i>	KxK moment ($x'x$) matrix. A constant term MUST have been the first variable when the moment matrix was computed.
<i>x</i>	NxK matrix of data.

Output

VC

KxK variance-covariance matrix.

Example

```
//Set rndseed for repeatable random numbers
rndseed 7234242;

//Create three randomly generated independent variables
x = rndn(500, 3);

//Create the sample variance-covariance matrix from data
matrix 'x'
var_x = varCovXS(x);
```

After the code above, `var_x` will be equal to:

1.0963733	0.0040911	-0.0024921
0.0040911	1.0627867	0.0494544
-0.0024921	0.0494544	0.8747116

where the diagonal elements in the matrix represent the sample variance of the each column, while the off-diagonal elements represent the sample covariance between the data columns.

The sample variance can also be calculated using the moment matrix, $x'x$ and the GAUSS function **varCovMS**. A constant term must be included in the data matrix x when computing the moment equation. Consider the following data matrix $x1$, consisting of the original data matrix x and a column of ones:

```
//Set rndseed so 'rndn' will return the same numbers as
above
rndseed 7234242;

//Note: the ~ operator performs horizontal concatenation
```

varget

```
x1 = ones(500,1)~rndn(500,3);

//Create moment matrix
x2 = x1'*x1;

//Calculate variance-covariance matrix using the moment matrix
var_xm = varCovMS(x2);
```

After the code above, `var_xm` will be equal to:

1.0963733	0.0040911	-0.0024921
0.0040911	1.0627867	0.0494544
-0.0024921	0.0494544	0.8747116

Remarks

The variance covariance matrix is that of the sample data matrix. It is computed as the moment matrix of deviations about the mean divided by the number of observations minus one, $N - 1$. For a population covariance matrix which uses N rather than $N - 1$ see **varCovM** or **varCovX**.

Source

corrs.src

See Also

[momentd](#), [corrms](#), [corrxs](#), [corrmm](#), [corrvc](#), [corrxx](#)

varget

Purpose

Accesses a global variable whose name is given as a string argument.

Format

```
y = varget(s);
```

Input

<i>s</i>	string containing the name of the global symbol you wish to access.
----------	---

Output

<i>y</i>	contents of the variable whose name is in <i>s</i> .
----------	--

Remarks

This function searches the global symbol table for the symbol whose name is in *s* and returns the contents of the variable if it exists. If the symbol does not exist, the function will terminate with an Undefined symbol error message. If you want to check to see if a variable exists before using this function, use **typecv**.

Example

```
alpha = 1;
beta = 2;
letter = "alpha";

//Check to see if a variable named alpha exists
if typecv(letter) == miss(0,0);
    print letter " does NOT exist";
else;
    //Assign the value of the variable named alpha to 'tmp'
    tmp = varget(letter);
    print "the value of " letter " is: " tmp;
endif;
```

vargetl

The code above produces the following output:

```
the value of alpha is: 1
```

vargetl

Purpose

Accesses a local variable whose name is given as a string argument.

Format

```
y = vargetl(s);
```

Input

<i>s</i>	string containing the name of the local symbol you wish to access.
----------	--

Output

<i>y</i>	contents of the variable whose name is in <i>s</i> .
----------	--

Remarks

This function searches the local symbol list for the symbol whose name is in *s* and returns the contents of the variable if it exists. If the symbol does not exist, the function will terminate with an Undefined symbol error message.

Example

```
proc rndNormEx( r, c, loc, std, ptVar);  
local rnd1, rnd2, rnd3;
```

```

//Create random normal numbers with mean 0 and standard
//deviation 1
rnd1 = rndn(r, c);

//Change the mean to 'loc'
rnd2 = rnd1 + loc;

//Change the standard deviation to 'std'
rnd3 = std * rnd2;

//Set the contents of tmp to be equal to the contents of
//the local variable with the same name as the string
//passed in as 'ptVar'
tmp = vargetl(ptVar);

    print ptVar " is equal to: " tmp;

    retp(rnd3);
endp;

//Set the rng seed for repeatable results
rndseed 54223423;

//Passing in the final variable as the string rnd1, will
//cause the proc rndNormEx to print the contents of rnd1
r = rndNormEx( 2, 2, 0, 3, "rnd1");

```

The code above will produce the following output:

```

rnd1 is equal to:
    0.5240627925408163   1.4904799236486497
   -1.1716182730350617  -0.0519353312479753

```

See Also

[varputl](#)

varmall

varmall

Purpose

Computes log-likelihood of a Vector ARMA model.

Format

```
ll = varmall(w, phi, theta, vc);
```

Input

<i>w</i>	NxK matrix, time series.
<i>phi</i>	(K*P)xK matrix, AR coefficient matrices.
<i>theta</i>	(K*Q)xK matrix, MA coefficient matrices.
<i>vc</i>	KxK matrix, covariance matrix.

Output

<i>ll</i>	scalar, log-likelihood. If the calculation fails <i>ll</i> is set to missing value with error code:
	Error Code Reason for Failure
	1 M < 1
	2 N < 1
	3 P < 0
	4 Q < 0
	5 P = 0 and Q = 0
	7 floating point work space too small
	8 integer work space too small



9	vc is not positive definite
10	AR parameters too close to stationarity boundary
11	model not stationary
12	model not invertible
13	I+M'H'HM not positive definite

Remarks

varmall is adapted from code developed by Jose Alberto Mauricio of the Universidad Complutense de Madrid. It was published as Algorithm AS311 in Applied Statistics. Also described in "Exact Maximum Likelihood Estimation of Stationary Vector ARMA Models," JASA, 90:282-264.

varmares

Purpose

Computes residuals of a Vector ARMA model.

Format

`res = varmares(w, phi, theta);`

Input

<i>w</i>	NxK matrix, time series.
<i>phi</i>	(K*P)xK matrix, AR coefficient matrices.
<i>theta</i>	(K*Q)xK matrix, MA coefficient matrices.



Output

<i>res</i>	NxK matrix, residuals. If the calculation fails <i>res</i> is set to missing value with error code:	
	Error Code	Reason for Failure
	1	$M < 1$
	2	$N < 1$
	3	$P < 0$
	4	$Q < 0$
	5	$P = 0$ and $Q = 0$
	7	floating point work space too small
	8	integer work space too small
	10	AR parameters too close to stationarity boundary
	11	model not stationary
	12	model not invertible
	13	$I+M'H'HM$ not positive definite

Remarks

varmares is adapted from code developed by Jose Alberto Mauricio of the Universidad Complutense de Madrid. It was published as Algorithm AS311 in Applied Statistics. Also described in "Exact Maximum Likelihood Estimation of Stationary Vector ARMA Models," JASA, 90:282-264.

>

varput

Purpose

Allows a matrix, array, string, or string array to be assigned to a global symbol whose name is given as a string argument.

Format

```
 $y = \text{varput}(x, \ n);$ 
```

Input

x	matrix, array, string, or string array which is to be assigned to the target variable.
n	string containing the name of the global symbol which will be the target variable.

Output

y	scalar, 1 if the operation is successful and 0 if the operation fails.
-----	--

Remarks

x and n may be global or local. The variable, whose name is in n , that x is assigned to is always a global.

If the function fails, it will be because the global symbol table is full.

This function is useful for returning values generated in local variables within a procedure to the global symbol table.

varputl

Example

```
source = rndn(2,2);  
targname = "target";  
  
if not varput(source,targname);  
    print "Symbol table full";  
end;  
endif;
```

See Also

[varget](#), [typecv](#)

varputl

Purpose

Allows a matrix, array, string, or string array to be assigned to a local symbol given as a string argument.

Format

```
y = varputl(x, n);
```

Input

x	matrix, array, string, or string array which is to be assigned to the target variable.
n	string containing the name of the local symbol which will be the target variable.

Output

y	scalar, 1 if the operation is successful and 0 if the operation fails.
-----	--

Remarks

x and n may be global or local. The variable, whose name is in n , that x is assigned to is always a local.

Example

```
proc myProc(x);
local a,b,c,d,e,vars,putvar;
  a=1;b=2;c=3;d=5;e=7;
  vars = { a b c d e };
  putvar = 0;

  //Keep looping until the user enters a letter
  //a-e or A-E
  do while putvar $/= vars;
    //Two semi-colons at the end of a print statement,
    //prevents a 'new line' from being printed
    print "Assign x (" $vars "): ";;
    putvar = upper(cons);
    print;
  endo;

  //Assign the variable whose letter/name was entered by
  //the user to be the value passed into 'myProc'
  call varputl(x,putvar);
  retp(a+b*c-d/e);
endp;

//Format printing of numbers to allow 2 spaces between them
```

vartypef

```
//and 1 digit after the decimal place
format /rds 2,1;

z = myProc(17);
print " z is " z;
```

produces (Note: this program will ask for user input at the GAUSS command prompt):

```
Assign x ( A B C D E ): a

z is 22.3
```

See Also

[vargetl](#)

vartypef

Purpose

Returns a vector of ones and zeros that indicate whether variables in a data set are character or numeric.

Format

```
y = vartypef(f);
```

Input

<i>f</i>	file handle of an open file.
----------	------------------------------

Output

<i>y</i>	Nx1 vector of ones and zeros, 1 if variable is numeric,
----------	---

0 if character.

Remarks

This function should be used in place of older functions that are based on the case of the variable names. You should also use the `v96` data set format.

vcm, vcx

Purpose

Computes an unbiased estimate a variance-covariance matrix.

NOTE: **vcm** and **vcx** have been replaced with functions **varCovXS** and **varCovMS** whose descriptions use more standard statistical nomenclature. **vcx** and **vcm** will continue to be available for backwards compatibility.

Format

```
VC = vcm(m);
VC = vcx(x);
```

Input

m	KxK moment ($x'x$) matrix. A constant term MUST have been the first variable when the moment matrix was computed.
x	NxK matrix of data.

Output

VC	KxK variance-covariance matrix.
------	---------------------------------

vcms, vcxs

Remarks

The variance-covariance matrix is computed as an unbiased estimator of the population variance-covariance. It is computed as the moment matrix of deviations about the mean divided by the number of observations minus one, $N - 1$. For an observed variance-covariance matrix which uses N rather than $N - 1$ see **vcms** or **vcxs**.

Source

corr.src

See Also

[momentd](#)

vcms, vcxs

Purpose

Computes the observed variance-covariance matrix.

NOTE: **vcms** and **vcxs** have been replaced with functions **varCovX** and **varCovM** whose descriptions use more standard statistical nomenclature. **vcxs** and **vcms** will continue to be available for backwards compatibility.

Format

```
VC = vcms(m);
VC = vcxs(x);
```

>

Input

m	KxK moment ($x'x$) matrix. A constant term MUST have been the first variable when the moment matrix was computed.
-----	--

x	$N \times K$ matrix of data.
-----	------------------------------

Output

vc	$K \times K$ variance-covariance matrix.
------	--

Remarks

The variance covariance matrix is that of the input data matrix. It is computed as the moment matrix of deviations about the mean divided by the number of observations N . For an unbiased estimator covariance matrix which uses $N - 1$ rather than N see **vcm** or **vcx**.

Source

`corrs.src`

See Also

[momentd](#), [corrms](#), [corrxs](#)

vec, vecr

Purpose

Creates a column vector by appending the columns/rows of a matrix to each other.

Format

```
 $y_C = \mathbf{vec}(x);$   
 $y_r = \mathbf{vecr}(x);$ 
```

vec, vecr

Input

x	$N \times K$ matrix.
-----	----------------------

Output

yc	$(N \times K) \times 1$ vector, the columns of x appended to each other.
yr	$(N \times K) \times 1$ vector, the rows of x appended to each other and the result transposed.

Remarks

vecr is much faster.

Example

```
x = { 1 2,
      3 4 };
yc = vec(x);
yr = vecr(x);
```

The code above assigns the variables yc and yr :

	1		1
$yc =$	3	$yr =$	2
	2		3
	4		4

>

vech

Purpose

Vectorizes a symmetric matrix by retaining only the lower triangular portion of the matrix.

Format

```
 $v = \mathbf{vech}(x);$ 
```

Input

x	$N \times N$ symmetric matrix.
-----	--------------------------------

Output

v	$(N*(N+1)/2) \times 1$ vector, the lower triangular portion of the matrix x .
-----	---

Remarks

As you can see from the example below, **vech** will not check to see if x is symmetric. It just packs the lower triangular portion of the matrix into a column vector in row-wise order.

Example

```
//Add a 3x1 column vector containing 10, 20, 30 to a 1x3  
//row vector containing 1, 2, 3, to create a 3x3 matrix  
 $x = \mathbf{seqa}(10,10,3) + \mathbf{seqa}(1,1,3)';$   
  
//Turn the lower triangular portion of 'x' into a column
```

vector (dataloop)

```
//vector in 'v'  
v = vech(x);  
  
//Expand the vector 'v' into a symmetric matrix in 'sx'  
sx = xpnd(v);
```

After the code above:

			11		
	11	12	13	21	11 21 31
x =	21	22	23	v = 22	sx = 21 22 32
	31	32	33	31	31 32 33
			32		
			33		

See Also

[xpnd](#)

vector (dataloop)

Purpose

Specifies the creation of a new variable within a data loop.

Format

```
vector # numvar = numeric_expression;  
vector $ charvar = character_expression;
```

Remarks

A *numeric_expression* is any valid expression returning a numeric value. A *character_expression* is any valid expression returning a character value. If

neither '\$' nor '#' is specified, '#' is assumed.

`vector` is used in place of `make` when the expression returns a scalar rather than a vector. `vector` forces the result of such an expression to a vector of the correct length. `vector` could actually be used anywhere that `make` is used, but would generate slower code for expressions that already return vectors.

Any variables referenced must already exist, either as elements of the source data set, as `extern`'s, or as the result of a previous `make`, `vector`, or `code` statement.

Example

```
vector const = 1;
```

See Also

[make \(dataloop\)](#)

vget

Purpose

Extracts a matrix or string from a data buffer constructed with `vput`.

Format

```
{ x, dbufnew } = vget(dbuf, name);
```

Input

<i>dbuf</i>	Nx1 vector, a data buffer containing various strings and matrices.
<i>name</i>	string, the name of the string or matrix to extract from <i>dbuf</i> .

view

Output

x	LxM matrix or string, the item extracted from <i>dbuf</i> .
<i>dbufnew</i>	Kx1 vector, the remainder of <i>dbuf</i> after x has been extracted.

Source

pack.src

See Also

[vlist](#), [vput](#), [vread](#)

view

Purpose

Sets the position of the observer in workbox units for 3-D plots. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
view( $x$ ,  $y$ ,  $z$ );
```

Input

x	scalar, the X position in workbox units.
y	scalar, the Y position in workbox units.
z	scalar, the Z position in workbox units.

Remarks

The size of the workbox is set with **volume**. The viewer MUST be outside of the workbox. The closer the position of the observer, the more perspective distortion there will be. If $x = y = z$, the projection will be isometric.

If **view** is not called, a default position will be calculated.

Use **viewxyz** to locate the observer in plot coordinates.

Source

pgraph.src

See Also

[volume](#), [viewxyz](#)

viewxyz

Purpose

To set the position of the observer in plot coordinates for 3-D plots. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
viewxyz(x, y, z);
```

Input

<i>x</i>	scalar, the X position in plot coordinates.
----------	---

vlist

<i>y</i>	scalar, the Y position in plot coordinates.
<i>z</i>	scalar, the Z position in plot coordinates.

Remarks

The viewer MUST be outside of the workbox. The closer the observer, the more perspective distortion there will be.

If **viewxyz** is not called, a default position will be calculated.

Use **view** to locate the observer in workbox units.

Source

pgraph.src

See Also

[volume](#), [view](#)

vlist

Purpose

Lists the contents of a data buffer constructed with **vput**.

Format

```
vlist(dbuf);
```

Input

<i>dbuf</i>	Nx1 vector, a data buffer containing various strings and matrices.
-------------	--

Remarks

vlist lists the names of all the strings and matrices stored in *dbuf*.

Source

`vpack.src`

See Also

[vget](#), [vput](#), [vread](#)

vnamecv

Purpose

Returns the names of the elements of a data buffer constructed with **vput**.

Format

```
cv = vnamecv(dbuf);
```

Input

<i>dbuf</i>	Nx1 vector, a data buffer containing various strings and matrices.
-------------	--

Output

<i>cv</i>	Kx1 character vector containing the names of the elements of <i>dbuf</i> .
-----------	--

See Also

[vget](#), [vput](#), [vread](#), [vtypecv](#)

volume

volume

Purpose

Sets the length, width, and height ratios of the 3-D workbox. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
volume(x, y, z);
```

Input

<i>x</i>	scalar, the X length of the 3-D workbox.
<i>y</i>	scalar, the Y length of the 3-D workbox.
<i>z</i>	scalar, the Z length of the 3-D workbox.

Remarks

The ratio between these values is what is important. If **volume** is not called, a default workbox will be calculated.

Source

pgraph.src

See Also

[view](#)

vput

Purpose

Inserts a matrix or string into a data buffer.

Format

```
dbufnew = vput(dbuf, x, xname);
```

Input

<i>dbuf</i>	Nx1 vector, a data buffer containing various strings and matrices. If <i>dbuf</i> is a scalar 0, a new data buffer will be created.
<i>x</i>	LxM matrix or string, item to be inserted into <i>dbuf</i> .
<i>xname</i>	string, the name of <i>x</i> , will be inserted with <i>x</i> into <i>dbuf</i> .

Output

<i>dbufnew</i>	Kx1 vector, the data buffer after <i>x</i> and <i>xname</i> have been inserted.
----------------	---

Remarks

If *dbuf* already contains *x*, the new value of *x* will replace the old one.

Source

`vpack.src`

See Also

[vget](#), [vlist](#), [vread](#)

vread

vread

Purpose

Reads a string or matrix from a data buffer constructed with **vput**.

Format

```
x = vread(dbuf, xname);
```

Input

<i>dbuf</i>	Nx1 vector, a data buffer containing various strings and matrices.
<i>xname</i>	string, the name of the matrix or string to read from <i>dbuf</i> .

Output

<i>x</i>	LxM matrix or string, the item read from <i>dbuf</i> .
----------	--

Remarks

vread, unlike **vget**, does not change the contents of *dbuf*. Reading *x* from *dbuf* does not remove it from *dbuf*.

Source

vpack.src

See Also

[vget](#), [vlist](#), [vput](#)

vtypecv

Purpose

Returns the types of the elements of a data buffer constructed with **vput**.

Format

```
cv = vtypecv(dbuf);
```

Input

<i>dbuf</i>	Nx1 vector, a data buffer containing various strings and matrices.
-------------	--

Output

<i>cv</i>	Kx1 character vector containing the types of the elements of <i>dbuf</i> .
-----------	--

See Also

[vget](#), [vput](#), [vread](#), [vnamecv](#)

wait, waitc

w

wait, waitc

Purpose

Waits until any key is pressed.

Format

```
wait;  
waitc;
```

Remarks

If you are working in terminal mode, these commands do not "see" any keystrokes until ENTER is pressed. **waitc** clears any pending keystrokes before waiting until another key is pressed.

Source

wait.src, waitc.src

See Also

[pause](#)

walkindex

Purpose

Walks the index of an array forward or backward through a specified dimension.

Format

```
ni = walkindex(i, o, dim);
```

Input

<i>i</i>	Mx1 vector of indices into an array, where $M \leq N$.
<i>o</i>	Nx1 vector of orders of an N-dimensional array.
<i>dim</i>	scalar [1-to-M], index into the vector of indices <i>i</i> , corresponding to the dimension to walk through, positive to walk the index forward, or negative to walk backward.

Output

<i>ni</i>	Mx1 vector of indices, the new index.
-----------	---------------------------------------

Remarks

walkindex will return a scalar error code if the index cannot walk further in the specified dimension and direction.

Example

```
orders = (3,4,5,6,7);  
  
//Create a 3x4x5x6x7 dimensional array with each element  
//equal to 1  
a = arrayinit(orders,1);  
  
ind = { 2,3,3 };  
ind = walkindex(ind,orders,-2);
```

window

```
      2  
ind = 2  
      3
```

This example decrements the second value of the index vector *ind*.

```
ind = walkindex(ind,orders,3);
```

```
      2  
ind = 2  
      4
```

Using the **orders** from the example above and the *ind* that was returned, this example increments the third value of the index vector *ind*.

See Also

[nextindex](#), [previousindex](#), [loopnextindex](#)

window

Purpose

Partitions the window into tiled regions (graphic panels) of equal size. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
window(row,col,typ);
```


Input

<i>row</i>	scalar, number of rows of graphic panels.
<i>col</i>	scalar, number of columns of graphic panels.
<i>typ</i>	scalar, graphic panel attribute type. If 1, the graphic panels will be transparent, if 0, the graphic panels will be nontransparent (blanked).

Remarks

The graphic panels will be numbered from 1 to $(row) \times (col)$ starting from the left topmost graphic panel and moving right.

See **makewind** for creating graphic panels of a specific size and position. (For more information, see **Graphic Panels**, Section 1.1.

Source

pwindow.src

See Also

[endwind](#), [begwind](#), [setwind](#), [nextwind](#), [getwind](#), [makewind](#)

writer

Purpose

Writes a matrix to a **GAUSS** data set.

Format

```
y = writer(fh, x);
```

writer

Input

<i>fh</i>	handle of the file that data is to be written to.
<i>x</i>	NxK matrix.

Output

<i>y</i>	scalar specifying the number of rows of data actually written to the data set.
----------	--

Remarks

The file must have been opened with `create`, `open for append`, or `open for update`.

The data in *x* will be written to the data set whose handle is *fh* starting at the current pointer position in the file. The pointer position in the file will be updated, so the next call to **writer** will put the next block of data after the first block. (See `open` and `create` for the initial pointer positions in the file for reading and writing.)

x must have the same number of columns as the data set. `colsf` returns the number of columns in a data set.

writer returns the number of rows actually written to the data set. If *y* does not equal `rows(x)`, the disk is probably full.

If the data set is not double precision, the data will be rounded as it is written out.

If the data contain character elements, the file must be double precision or the character information will be lost.

If the file being written to is the 2-byte integer data type, then missing values will be written out as -32768. These will not automatically be converted to missings on input. They can be converted with the **miss** function:

```
x = miss(x,-32768);
```

Trying to write complex data to a data set that was originally created to store real data will cause a program to abort with an error message. (See [create](#) for details on creating a complex data set.)

Example

```
create fp = data with x,10,8;

if fp == -1;
    errorlog "Can't create output file";
end;
endif;

c = 0;
do until c >= 10000;
    y = randn(100,10);
    k = writer(fp,y);

    if k /= rows(y);
        errorlog "Disk Full";
        fp = close(fp);
    end;
endif;

    c = c+k;
enddo;

fp = close(fp);
```

In this example, a 10000x10 data set of Normal random numbers is written to a data set called `data.dat`. The variable names are `X01 - X10`.

See Also

[open](#), [close](#), [create](#), [readr](#), [saved](#), [seekr](#)

x

xlabel

x

xlabel

Purpose

Sets a label for the X axis. NOTE: This function is for use with the deprecated PQG graphics, use **plotSetXLabel** for equivalent functionality.

Library

pgraph

Format

```
xlabel(str);
```

Input

<i>str</i>	string, the label for the X axis.
------------	-----------------------------------

Source

pgraph.src

See Also

[title](#), [ylabel](#), [zlabel](#)

xlsGetSheetCount

Purpose

Gets the number of sheets in an Excel® spreadsheet.

Format

```
nsheets = xlsGetSheetCount(file);
```

Input

<i>file</i>	string, name of .xls or .xlsx file.
-------------	-------------------------------------

Output

<i>nsheets</i>	scalar, sheet count or an error code.
----------------	---------------------------------------

Portability

Windows, Linux and Mac

Remarks

If **xlsGetSheetCount** fails, it will either terminate with an error message or return a scalar error code, which can be decoded with **scalerr**, depending on the lowest order bit of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code 10.

Example

Example 1

xlsGetSheetSize

If you had an Excel file named 'yarn.xlsx' in the GAUSS home directory, then you could determine the number of sheets in the file with the following code:

```
// File name with full path
fname= getGAUSShome() $+ "examples/yarn.xlsx";
nsheets = xlsGetSheetCount(fname);
```

Example 2

If you do not want your program to terminate in the case of an error in this function, you can set the `trap` state as in the example below.

```
//Turn on trap
trap 1;

nsheets = xlsGetSheetCount(fname);

//Check to see if xlsGetSheetCount returned an error code
if scalmiss(nsheets);
//Code to execute in error case here
endif;
```

See Also

[xlsGetSheetSize](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

xlsGetSheetSize

Purpose

Gets the size (rows and columns) of a specified sheet in an Excel® spreadsheet.

Format

```
{ r, c } = xlsGetSheetSize(file);  
{ r, c } = xlsGetSheetSize(file, sheet);
```

Input

<i>file</i>	string, name of .xls or .xlsx file.
<i>sheet</i>	scalar, sheet index (1-based). Default = 1.

Output

<i>r</i>	scalar, number of rows.
<i>c</i>	scalar, number of columns.

Portability

Windows, Linux and Mac

Remarks

If **xlsGetSheetSize** fails, it will either terminate and print an error message or return a scalar error code, which can be decoded with **scalerr**, depending on the state of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code 10.

If a scalar error code is returned, both return values will be set with the error code.

xlsGetSheetSize

Examples

Example 1

If you had an Excel file named '**yarn.xlsx**' in the GAUSS home directory , then you could determine the number of rows and columns in the first sheet of this file with the following code:

```
// File name with full path
fname= getGAUSShome() $+ "examples/yarn.xlsx";
sheetNum = 1;

// call xlsGetSheetSize function
{ r, c } = xlsGetSheetSize(fname, sheetNum);
```

Example 2

If you do not want your program to terminate in the case of an error in this function, you can set the **trap** state as in the example below.

```
sheetNum = 1;

//Retain the old trap value so it can
//be reset to its previous state
oldtrap = trapchk(1);

//Set trap
trap 1;

{ r, c } = xlsGetSheetSize(fname, sheetNum);

//Check to see if return value is an error code
if scalmiss(r);
    //User error handling code here
endif;
```


See Also

[xlsGetSheetCount](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

xlsGetSheetTypes

Purpose

Gets the cell format types of a row in an Excel® spreadsheet.

Format

```
nsheets = xlsGetSheetTypes(file, sheet, row);
```

Input

<i>file</i>	string, name of .xls or .xlsx file.
<i>sheet</i>	scalar, sheet index (1-based).
<i>row</i>	scalar, the row of cells to be scanned.

Output

<i>types</i>	1xK vector of predefined data types representing the format of each cell in the specified row.
--------------	--

The possible types are:

0	Text
1	Numeric
2	Date

Portability

Windows, Linux and Mac

xlsGetSheetTypes

Example

For example, let us suppose that a file named 'yarn.xlsx' exists in the GAUSS home directory. Let us further suppose that the 'A1' element is a string and the 'B1:C1' elements are numbers. The first row has no other elements. Then the code:

```
// File name with full path
fname= getGAUSShome() $+ "examples/yarn.xlsx";
sheetNum = 1;
rowNum = 2;
ctypes = xlsGetSheetTypes(fname, sheetNum, rowNum);

//Do not print any values after the decimal point
format /rd 6,0;
print ctypes;
```

would produce the following output:

0	0	0	1
---	---	---	---

Remarks

K is the number of columns found in the spreadsheet.

If **xlsGetSheetTypes** fails, it will either terminate and print an error message or return a scalar error code, which can be decoded with **scalerr**, depending on the state of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code 10.

See Also

[xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsMakeRange](#)

xlsMakeRange

Purpose

Builds an Excel® range string from a row/column pair.

Format

```
range = xlsMakeRange(row, col);
```

Input

<i>row</i>	scalar or 2x1 vector.
<i>col</i>	scalar or 2x1 vector.

Output

<i>range</i>	string, an Excel®-formatted range specifier.
--------------	--

Portability

Available on **Windows**, **Linux** and **Mac**.

Remarks

If *row* is a 2x1 vector, it is interpreted as follows

<i>row</i> [1]	starting row
<i>row</i> [2]	ending row

If *col* is a 2x1 vector, it is interpreted as follows:

<i>col</i> [1]	starting column
<i>col</i> [2]	ending column

xlsMakeRange

If **xlsMakeRange** fails, it will either terminate and print an error message or return a scalar error code, which can be decoded with **scalerr**, depending on the state of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code 10.

Example

```
//Scalar inputs
r = 3;
c = 6;
range = xlsMakeRange(r, c);
print range;
```

produces:

F3

```
//2x1 vector inputs
r = { 2, 37 };
c = { 3, 19 };
range = xlsMakeRange(r, c);
print range;
```

produces:

C2:S37

See Also

[xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsGetSheetTypes](#)

xlsReadM

Purpose

Reads from an Excel® spreadsheet into a **GAUSS** matrix.

Format

```
mat = xlsReadM(file);  
mat = xlsReadM(file, range);  
mat = xlsReadM(file, range, sheet);  
mat = xlsReadM(file, range, sheet, vls);
```

Input

<i>file</i>	string, name of .xls or .xlsx file.
<i>range</i>	string, range to read, e.g. "A2:B20", or the starting point of the read, e.g. "A2". Default = "A1".
<i>sheet</i>	scalar, sheet number to read from. Default = 1.
<i>vls</i>	null string or 9x1 matrix, specifies the conversion of Excel® empty cells and special types into GAUSS (see Remarks). A null string results in all empty cells and special types being converted to GAUSS missing values.

Output

<i>mat</i>	matrix or a scalar error code.
------------	--------------------------------

xlsReadM

Examples

Example 1: Basic Example

```
//Create file name with full path
file = getGAUSSHome() %+ "examples/tbill_3mo.xlsx";

//Read in all data below header line
x = xlsReadM(file, "A2");
```

After the code above, the first 10 rows of `x` should be equal to:

19820101000000	12.92
19820201000000	14.28
19820301000000	13.31
19820401000000	13.34
19820501000000	12.71
19820601000000	13.08
19820701000000	11.86
19820801000000	9
19820901000000	8.19
19821001000000	7.97

Example 2: Read From a Range

```
//Create file name with full path
file = getGAUSSHome() %+ "examples/yarn.xlsx";

//Read in data from rows 2-9 of column 'D'
x = xlsReadM(file, "D2:D9");
```

After the code above, `x` should be equal to:

674
370

```
292
338
266
210
170
118
```

Example 3: Reading dates

```
//Create file name with full path
file = getGAUSSHome() $+ "examples/tbill_3mo.xlsx";

//Read the first element below the header from the first
column
date_1 = xlsReadM(file, "A2:A2");
```

If the Excel file has marked a cell as a date, **GAUSS** will read it in DT scalar format. After the code above, `date_1` will be equal to:

```
19820101000000
```

Dates in DT scalar format can be passed in directly to **plotTS** to create time series plots, and also handled by other **GAUSS** date handling functions. For example, we can convert `date_1` to a string with the function **dttostr** (date to string) like this:

```
date_str = dttostr(date_1, "MO-DD-YYYY");
```

After which, `date_str` will be equal to:

```
"01-01-1982"
```

Example 4: Specify Sheet Number

```
//Using the 'file' variable created in the previous example
//Pass in '1' as the third input, to specify the first
```

xlsReadM

```
sheet
x = xlsReadM(file, "A2:A10", 1);
```

Remarks

1. If range is a null string, then by default the read will begin at cell "A1".
2. If **xlsReadM** fails, it will either terminate and print an error message or return a scalar error code, which can be decoded with **scalerr**, depending on the state of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code.

2.1 An error message example

```
//Will end the program and print an error message
x = xlsReadM("nonexistent_file.xlsx");
```

2.2 Turn off error message

```
//Turn error trapping on
trap 1;
x = xlsReadM("nonexistent_file.xlsx");

//Check to see if 'x' is a scalar error code
if scalmiss(x);
    //Code to handle error case here
endif;

//Turn error trapping off
trap 0;
```


3. By default, empty cells are imported as **GAUSS** missing values. The `vls` argument lets users control the import of Excel® empty cells and special types, according to the following table:

Row Number	Excel® Cell
1	empty cell
2	#N/A
3	#VALUE!
4	#DIV/0!
5	#NAME?
6	#REF!
7	#NUM!
8	#NULL!
9	#ERR

Use the following to convert all occurrences of #DIV/0! to +Infinity, and all other empty cells and special types to **GAUSS** missing values:

```
//Create a 9x1 vector of missing values
vls = reshape(miss(0,0),9,1);

//Set the 4th element of 'vls' to +Infinity so that
//Excel #DIV/0! cells will be imported as +Infinity
vls[4] = __INFP;

x = xlsReadM("myfile.xlsx", "A1", 1, vls);
```

Portability

Windows, Linux and Mac

xlsReadSA

The *vls* input is currently ignored on Mac and Linux. Missing values will be returned for all cells that are empty or contain errors.

See Also

[xlsReadSA](#), [xlsWrite](#), [xlsWriteM](#), [xlsWriteSA](#), [xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

xlsReadSA

Purpose

Reads from an Excel® spreadsheet into a **GAUSS** string array or string.

Format

```
s = xlsReadSA(file);
s = xlsReadSA(file, range);
s = xlsReadSA(file, range, sheet);
s = xlsReadSA(file, range, sheet, vls);
```

Input

<i>file</i>	string, name of .xls or .xlsx file.
<i>range</i>	string, range to read, e.g. "A2:B20" or the starting point of the read, e.g. "A2". Default = "A1".
<i>sheet</i>	scalar, sheet number. Default = 1.
<i>vls</i>	null string or 9x1 string array, specifies the conversion of Excel® empty cells and special types into GAUSS (see Remarks). A null string results in all empty cells and special types being converted to null strings. Default = null string.

Output

`s` string, string array or a scalar error code.

Examples

Example 1: Basic Example with Specify Path and Sheet Number

Read all contents from the file "yarn.xlsx" located in GAUSS home working directory as a string array.

```
//Create file name with full path
file = getGAUSSHome() $+ "examples/yarn.xlsx";
// "A1" means start from A1
// 1 = sheet number
// Call xlsReadSA function
s = xlsReadSA(file, "A1", 1);
```

Example 2: Read From a Range

```
data = xlsReadSA(file, "A2:D28");
```

Example 3: Read your own data

Read all contents from the file myfile.xlsx located in your current GAUSS working directory as a string array.

```
s = xlsReadSA("myfile.xlsx");
```

Remarks

1. If range is a null string, then by default the read will begin at cell "A1".
2. If `xlsReadSA` fails, it will either terminate and print an error message or

xlsReadSA

return a scalar error code, which can be decoded with **scalerr**, depending on the state of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code.

```
//Will end the program and print an error message
x = xlsReadSA("nonexistent_file.xlsx");
```

```
//Turn error trapping on
trap 1;
x = xlsReadSA("nonexistent_file.xlsx");

//Check to see if 'x' is a scalar error code
if scalmiss(x);
    //Code to handle error case here
endif;

//Turn error trapping off
trap 0;
```

3. By default, empty cells are imported as empty strings. The `vls` argument lets users control the import of Excel® empty cells and special types, according to the following table:

Row Number	Excel® Cell
1	empty cell
2	#N/A
3	#VALUE!
4	#DIV/0!

5	#NAME?
6	#REF!
7	#NUM!
8	#NULL!
9	#ERR

Use the following to convert all occurrences of #NULL! and empty cells to the string 'NULL', and all other and special types to empty strings:

```
//Create a 9x1 vector of empty strings
vls = reshape("", 9, 1);

//Set the 1st and 8th element of 'vls' to the string
'NULL' so that
//Excel #NULL! and empty cells will be imported as the
string 'NULL'
vls[1] = "NULL";
vls[8] = "NULL";

x = xlsReadSA("myfile.xlsx", "A1", 1, vls);
```

Portability

Windows, Linux and Mac

The `vls` input is currently ignored on Mac and Linux. Missing values will be returned for all cells that are empty or contain errors.

See Also

[xlsReadM](#), [xlsWrite](#), [xlsWriteM](#), [xlsWriteSA](#), [xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

xlsWrite

xlsWrite

Purpose

Writes a **GAUSS** matrix, string, or string array to an Excel® spreadsheet.

Format

```
ret=xlsWrite(data, file);  
ret=xlsWrite(data, file, range);  
ret=xlsWrite(data, file, range, sheet);  
ret=xlsWrite(data, file, range, sheet, vls);
```

Input

<i>data</i>	matrix, string, or string array.
<i>file</i>	string, name of .xls or .xlsx file.
<i>range</i>	string, the starting point of the write, e.g. "A2". Default = "A1".
<i>sheet</i>	scalar, sheet number. Default = 1.
<i>vls</i>	null string or 9x1 matrix or string array, specifies the conversion of GAUSS values or characters into Excel® empty cells and special types (see Remarks). A null string results in all GAUSS missing values and null strings being converted to empty cells. Default = null string.

Output

<i>ret</i>	scalar, 0 if success or a scalar error code.
------------	--

Examples

Example 1: Basic Example

```
x = { 0 1,  
      1 2,  
      3 5 };  
  
//Write contents of 'x' to 'myfile.xlsx'  
//from cell 'A1' to 'B3'  
ret = xlsWrite(x, "myfile.xlsx");
```

'myfile.xlsx' is saved in your current working directory. You can find your current working directory in the main tool bar (in the top of GAUSS).

Example 2: Write To a Range

```
//Create a 1x4 string array of variable names  
head = "Real GDP" $~ "Unemployment" $~ "CPI" $~ "PPI";  
  
//Write the variable names to the cells 'C1:F1'  
ret = xlsWrite(head, "myfile.xlsx", "C1");
```

Example 3: Specify Path and Sheet Number

```
//Create a 10x3 matrix of Bernoulli random variables  
x = rndBernoulli(10, 3, 0.6);  
  
//Write the data from 'x' to cells 'B4:D13' on sheet 2 of  
'myfile.xlsx'  
ret = xlsWrite(x, "C:\\mydata\\myfile.xlsx", "B4", 2);
```

xlsWrite

Remarks

1. If **xlsWrite** fails, it will either terminate and print an error message or return a scalar error code, which can be detected with **scalmiss**, depending on the state of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code.

1.1 An error message example

```
//If this fails, it will end the program and print an
error message
x = xlsWrite("myfile.xlsx");
```

1.2 Turn off error message

```
//Turn error trapping on
trap 1;
x = xlsWrite("myfile.xlsx");

//Check to see if 'x' is a scalar error code
if scalmiss(x);
    //Code to handle error case here
endif;

//Turn error trapping off
trap 0;
```

2. The **xls** argument lets users control the export to Excel® empty cells and special types, according to the following table:

Row Number	Excel® Cell
1	empty cell
2	#N/A
3	#VALUE!
4	#DIV/0!
5	#NAME?
6	#REF!
7	#NUM!
8	#NULL!
9	#ERR

Use the following to convert all occurrences of 9999.99 to #DIV/0! in Excel® and convert all **GAUSS** missing values to empty cells in Excel®:

```
vls = reshape(error(0),9,1);  
vls[4] = 9999.99;
```

Portability

Windows, Linux and Mac

The `vls` input is currently ignored on Mac and Linux. Missing values will be returned for all cells that are empty or contain errors.

See Also

[xlsReadSA](#), [xlsReadM](#), [xlsWriteM](#), [xlsWriteSA](#), [xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

xlsWriteM

xlsWriteM

Purpose

Writes a **GAUSS** matrix to an Excel® spreadsheet.

Format

```
ret=xlsWriteM(data, file);  
ret=xlsWriteM(data, file, range);  
ret=xlsWriteM(data, file, range, sheet);  
ret=xlsWriteM(data, file, range, sheet, vls);
```

Input

<i>data</i>	matrix.
<i>file</i>	string, name of .xls or .xlsx file.
<i>range</i>	string, the starting point of the write, e.g. "a2". Default = "a1"
<i>sheet</i>	scalar, sheet number. Default = 1.
<i>vls</i>	null string or 9x1 matrix, specifies the conversion of GAUSS values into Excel® empty cells and special types (see Remarks). A null string results in all GAUSS missing values being converted to empty cells. Default = null string.

Output

<i>ret</i>	scalar, 0 if success or a scalar error code.
------------	--

Examples

Example 1: Basic Example

```
x = { 0 1,  
      1 2,  
      3 5 };  
  
//Write contents of 'x' to 'myfile.xlsx'  
//from cell 'A1' to 'B3'  
ret = xlsWriteM(x, "myfile.xlsx");
```

'myfile.xlsx' is saved in your current working directory. You can find your current working directory in the main tool bar (in the top of GAUSS).

Example 2: Write To a Range

```
//Write 'x' from the previous example to the cells 'C2:D4'  
ret = xlsWriteM(x, "myfile.xlsx", "C2");
```

Example 3: Specify Path and Sheet Number

```
//Create a 10x3 matrix of Bernoulli random variables  
x = rndBernoulli(10, 3, 0.6);  
  
//Write the data from 'x' to cells 'B4:D13' on sheet 2 of  
'myfile.xlsx'  
ret = xlsWriteM(x, "C:\\mydata\\myfile.xlsx", "B4", 2);
```

Remarks

1.

The `vls` argument lets users control the export to Excel® empty cells and special types, according to the following table:

xlsWriteM

Row Number	Excel® Cell
1	empty cell
2	#N/A
3	#VALUE!
4	#DIV/0!
5	#NAME?
6	#REF!
7	#NUM!
8	#NULL!
9	#ERR

Use the following to convert all occurrences of 9999.99 to #DIV/0! in Excel® and convert all **GAUSS** missing values to empty cells in Excel®:

```
vls = reshape(error(0),9,1);
vls[4] = 9999.99;
```

- 2. If **xlsWriteM** fails, it will either terminate and print an error message or return a scalar error code, which can be decoded with **scalerr**, depending on the state of the **trap** flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code 10.

Portability

Windows, Linux and Mac

The `vls` input is currently ignored on Mac and Linux. Missing values will be returned for all cells that are empty or contain errors.

See Also

[xlsReadSA](#), [xlsReadM](#), [xlsWrite](#), [xlsWriteSA](#), [xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

xlsWriteSA

Purpose

Writes a **GAUSS** string or string array to an Excel® spreadsheet.

Format

```
ret = xlsWriteSA(data, file);  
ret = xlsWriteSA(data, file, range);  
ret = xlsWriteSA(data, file, range, sheet);  
ret = xlsWriteSA(data, file, range, sheet, vls);
```

Input

<i>data</i>	string or string array.
<i>file</i>	string, name of .xls file.
<i>range</i>	string, the starting point of the write, e.g. "a2". Default = "a1".
<i>sheet</i>	scalar, sheet number. Default = 1.
<i>vls</i>	null string or 9x1 string array, specifies the conversion of GAUSS characters into Excel® empty cells and special types (see Remarks). A null string results in all null strings being converted to empty cells. Default = null string.

xlsWriteSA

Output

<i>ret</i>	scalar, 0 if success or a scalar error code.
------------	--

Examples

Example 1: Basic Example

```
//Create a 1x3 string array of variable names
var_names = "Date" $~ "Price" $~ "Volume";

//Write contents of 'var_names' to 'myfile.xlsx'
//from cell 'A1' to 'C1'
ret = xlsWriteSA(var_names, "myfile.xlsx");
```

'myfile.xlsx' is saved in your current working directory. You can find your current working directory in the main tool bar (in the top of GAUSS).

Example 2: Write To a Range

```
//Create a 1x4 string array of variable names
head = "Real GDP" $~ "Unemployment" $~ "CPI" $~ "PPI";

//Write the variable names to the cells 'C1:F1'
ret = xlsWriteSA(head, "myfile.xlsx", "C1");
```

Example 3: Specify Path and Sheet Number

```
//Create a 3x1 string array
labels = "Normotensive" $| "Hypertensive" $|
"Hypotensive";

//Write the data from 'labels' to cells 'D7:D9' on sheet 2
```

```
of 'myfile.xlsx'  
ret = xlsWriteSA(labels, "C:/mydata/myfile.xlsx", "D7", 2);
```

Remarks

1. The `vls` argument lets users control the export to Excel® empty cells and special types, according to the following table:

Row Number	Excel® Cell
1	empty cell
2	#N/A
3	#VALUE!
4	#DIV/0!
5	#NAME?
6	#REF!
7	#NUM!
8	#NULL!
9	#ERR

Use the following to convert all occurrences of "Division by Zero" to #DIV/0!, and all null strings to empty cells:

```
vls = reshape("", 9, 1);  
vls[4] = "Division by Zero";
```

2. If `xlsWriteSA` fails, it will either terminate and print an error message or return a scalar error code, which can be decoded with `scalerr`, depending on the state of the `trap` flag.

trap 0	Print error message and terminate program.
trap 1	Return scalar error code 10.

xpnd

Portability

Windows, Linux and Mac

The `xls` input is currently ignored on Mac and Linux. Missing values will be returned for all cells that are empty or contain errors.

See Also

[xlsReadM](#), [xlsWrite](#), [xlsWriteM](#), [xlsReadSA](#), [xlsGetSheetCount](#), [xlsGetSheetSize](#), [xlsGetSheetTypes](#), [xlsMakeRange](#)

xpnd

Purpose

Expands a column vector into a symmetric matrix.

Format

```
x = xpnd(v);
```

Input

`v`

Kx1 vector, to be expanded into a symmetric matrix.

Output

`x`

MxM matrix, the results of taking `v` and filling in a symmetric matrix with its elements.

$$M = ((-1 + \text{sqrt}(1 + 8 * K)) / 2)$$

Remarks

If v does not contain the right number of elements, (that is, if $\text{sqrt}(1 + 8 * K)$ is not integral), then an error message is generated.

This function is particularly useful for hard-coding symmetric matrices, because only about half of the matrix needs to be entered.

Example

```
x = { 1,
      2, 3,
      4, 5, 6,
      7, 8, 9, 10 };
y = xpnd(x);
```

After the code above, the variables x and y are equal to:

```

      1
      2
      3
      4      1      2      4      7
x = 5      y = 2      3      5      8
      6      4      5      6      9
      7      7      8      9     10
      8
      9
     10
```

See Also

[vech](#)

x_{tics}

x_{tics}

Purpose

Sets and fixes scaling, axes numbering and tick marks for the X axis. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
xtics(min, max, step, minordiv);
```

Input

<i>min</i>	scalar, the minimum value.
<i>max</i>	scalar, the maximum value.
<i>step</i>	scalar, the value between major tick marks.
<i>minordiv</i>	scalar, the number of minor subdivisions.

Remarks

This routine fixes the scaling for all subsequent graphs until **graphset** is called.

This gives you direct control over the axes endpoints and tick marks. If **x_{tics}** is called after a call to **scale**, it will override **scale**.

X and Y axes numbering may be reversed for **xy**, **logx**, **logy**, and **loglog** graphs. This may be accomplished by using a negative step value in the **x_{tics}** and **y_{tics}** functions.

Source

pscale.src

See Also

[scale](#), [ytics](#), [ztics](#)

xy

Purpose

Graphs X vs. Y using Cartesian coordinates. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
xy(x, y);
```

Input

<i>x</i>	Nx1 or NxM matrix. Each column contains the X values for a particular line.
<i>y</i>	Nx1 or NxM matrix. Each column contains the Y values for a particular line.

Remarks

Missing values are ignored when plotting symbols. If missing values are encountered while plotting a curve, the curve will end and a new curve will begin plotting at the next non-missing value.

Source

`pxy.src`

xyz**See Also**

[xyz](#), [logx](#), [logy](#), [loglog](#)

xyz**Purpose**

Graphs X vs. Y vs. Z using Cartesian coordinates. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
xyz(x, y, z);
```

Input

<i>x</i>	Nx1 or NxK matrix. Each column contains the X values for a particular line.
<i>y</i>	Nx1 or NxK matrix. Each column contains the Y values for a particular line.
<i>z</i>	Nx1 or NxK matrix. Each column contains the Z values for a particular line.

Remarks

Missing values are ignored when plotting symbols. If missing values are encountered while plotting a curve, the curve will end and a new curve will begin plotting at the next non-missing value.

xyz

x

Source

pxyz.src

ylabel

y

ylabel

Purpose

Sets a label for the Y axis. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
ylabel(str);
```

Input

<i>str</i>	string, the label for the Y axis.
------------	-----------------------------------

Source

pgraph.src

See Also

[title](#), [xlabel](#), [ylabel](#)

ytics

Purpose

Sets and fixes scaling, axes numbering and tick marks for the Y axis. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
ytics(min, max, step, minordiv);
```

Input

<i>min</i>	scalar, the minimum value.
<i>max</i>	scalar, the maximum value.
<i>step</i>	scalar, the value between major tick marks.
<i>minordiv</i>	scalar, the number of minor subdivisions.

Remarks

This routine fixes the scaling for all subsequent graphs until **graphset** is called.

This gives you direct control over the axes endpoints and tick marks. If **ytics** is called after a call to **scale**, it will override **scale**.

X and Y axes numbering may be reversed for **xy**, **logx**, **logy** and **loglog** graphs. This may be accomplished by using a negative step value in the **xtics** and **ytics** functions.

Source

pscale.src

yticks

See Also

[scale](#), [xticks](#), [zticks](#)

y

z

zeros

Purpose

Creates a matrix of zeros.

Format

```
y = zeros(r, c);
```

Input

<i>r</i>	scalar, the number of rows.
<i>c</i>	scalar, the number of columns.

Output

<i>y</i>	<i>r</i> x <i>c</i> matrix of zeros.
----------	--------------------------------------

Remarks

This is faster than **ones**.

Noninteger arguments will be truncated to an integer.

zeta

Example

```
y = zeros(3,2);  
print y;
```

The code above produces the following output:

```
0.000    0.000  
0.000    0.000  
0.000    0.000
```

See Also

[ones](#), [eye](#)

zeta

Purpose

Computes the Riemann Zeta function.

Format

```
f = zeta(z);
```

Input

z NxK matrix; *z* may be complex.

Output

f NxK matrix.

Remarks

Euler MacLaurin series.

References

1. Jon Breslaw, 2009

zlabel

Purpose

Sets a label for the Z axis. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
zlabel(str);
```

Input

<i>str</i>	string, the label for the Z axis.
------------	-----------------------------------

Source

pgraph.src

See Also

[title](#), [xlabel](#), [ylabel](#)

ztics

ztics

Purpose

Sets and fixes scaling, axes numbering and tick marks for the Z axis. NOTE: This function is for the deprecated PQG graphics.

Library

pgraph

Format

```
ztics(min, max, step, minordiv);
```

Input

<i>min</i>	scalar, the minimum value.
<i>max</i>	scalar, the maximum value.
<i>step</i>	scalar, the value between major tick marks.
<i>minordiv</i>	scalar, the number of minor subdivisions. If this function is used with contour , contour labels will be placed every <i>minordiv</i> levels. If 0, there will be no labels.

Remarks

This routine fixes the scaling for all subsequent graphs until **graphset** is called.

This gives you direct control over the axes endpoints and tick marks. If **ztics** is called after a call to **scale3d**, it will override **scale3d**.

Source

pscale.src

See Also

[scale3d](#), [xticks](#), [yticks](#), [contour](#)

36 Miscellaneous Topics

36.1 Change Log

The following is a list of changes from the previous version of **GAUSS**. This version is completely backwards compatible with the previous major version, 16.

17.0.2

1. (Windows) Added GUI license management utility for floating network licenses, **rlmservice.exe**.
2. Bug fix: Fixed possible folder duplication in Source Browser.
3. Bug fix for program files not saving on run, in specific cases.
4. Bug fix for possible underflow in **cdftnc**
5. Added ability to remove **tgauss** dependency on Qt libraries.
6. Other minor documentation enhancements and bug fixes.

17.0.1

1. Up to 2-5x speed increase for least squares estimation, using the slash operator '/' for non-square coefficient matrices with few columns (approximately 1-15) and few rows (approximately 2-100).
2. Bug fix: File import dialog now supports empty sheet names for Excel files.
3. Added support for pasting data to the matrix editor from applications that use the legacy carriage return only line ending on Mac.

17.0.0

1. Increased scalability of `threadFor` and `threadBegin`.
2. GAUSS commands that process datasets can now also process `.fmt`, and `.h5` files as well as `.dat`.
3. New support for HDF5 datasets, allows unlimited sized datasets.
4. Added support for an initial subset of Wilkinson-Rogers formula notation for functions such as `dstat`, `dstatmt`, `glm`, `momentd`, `ols`, `olsmt` and more.
5. CSV and Excel (`.xls`, `.xlsx`) files can be used as datasets for functions `quantiled`, `dstatmt`, `glm`, `momentd`, `ols`, `olsmt` and more.
6. New function `cdfEmpirical` for computing the empirical cumulative distribution function, and `plotcdfEmpirical` to graph it.
7. New function `plotAddErrorBar` adds error bars to 2-D plots.
8. New function `plotAddSurface` adds additional surfaces to an existing surface plot.
9. New function `plotSetLegendFont` to control the font family, size and color used in the legend.
10. New function `plotSetZLevels`: user control for the height of levels, rather than just the number of contour levels.
11. New function `plotSetContourLabels`: controls whether numeric label containing contour level height is drawn, as well as the format of the numeric label.
12. New color maps for surface and contour plots.
13. Added option to specify the units and dpi of graphs saved with `plotSave`.
14. Added control for the range of the X and Y axes to the graphics editor.
15. Added option to control units of graph size and DPI to `plotSave`.
16. Added control for viewing angle, lighting, zoom and toggling appearance of the wireframe for surface plots to the graphics editor.
17. New function `sylvester` to compute the solution, X , to the equation $AX + XB = C$.
18. `schur` can now, optionally, return the real or complex Schur form.
19. New function `dot` to compute the dot product of a column or the columns of a matrix.
20. New function `powerM` to raise a matrix to a specified power.
21. `getdims` will now return the number of dimensions of a matrix, string or string array.

22. **getorders** will now return the number of rows and columns for matrices, strings or string arrays.
23. Greatly improved speed and decreased memory usage for **reclassify**.
24. Greatly improved performance of **unique** and **uniquesa** for string arrays.
25. Greatly improved performance of **sortc** for column vectors.
26. Greatly improved performance of linear solve using the slash operator (/) for small matrices and X'X matrix multiplication for large matrices.
27. Greatly improved performance of kronecker product operator (.*.) when one of the matrices is a column vector.
28. Improved performance of **cdffc** when the **d1** parameter is equal to one, by 10-1000x.
29. Improved performance of **crossprd** for the case with fewer than 500 vectors.
30. Added support for complex inputs to **hess** and significant speed up for real matrix inputs larger than approximately 30x30.
31. **tgauss** and the **GAUSS Engine** can now create new 'plot' graphics.
32. New function **rndWishartInv** for taking draws from the Inverse Wishart distribution.
33. New function **pdfWishartInv** computes the probability density function of the Inverse Wishart distribution.
34. New function **ldl** computes the LDL decomposition of a positive semi-definite matrix and returns separate L and D factors.
35. Added support for generalized linear model function, **glm** for inverse-Gaussian distribution and model without an intercept.
36. New function **strtrim** to remove white space from left and right side of elements of a string array.
37. Added support for multi-character delimiters to **strsplit**.
38. New function **strjoin** to combine string array elements into a string separated by a specified delimiter. This function does NOT add a delimiter after the final element as in **strcombine**.
39. Editor now grays out code that is inactive due to a **#define**.
40. Application Install Wizard can install multiple GAUSS application modules at once.
41. Performance improvement: The "forward only" flag (dbQuerySetForwardOnly) now defaults to true.
42. Bug fix for find-and-replace with UTF-8 multibyte characters.

Miscellaneous Topics

43. Bug fix: added support for strings to **selif** and **delif** and fixed memory bug in **delif**.
44. Bug fix: crash when all points sent to **plotLogX**, **plotLogY** or **plotLogLog** were between 10^n and 10^{n+1} .
45. Bug fix: **errorlog** and **errorlogat** now accept 1x1 string arrays as well as strings.
46. Bug fix: **intsimp** would fail with an error when attempting to integrate a function that returned only zeros.
47. Bug fix: ability to scroll to right end in program input/output window with long lines.
48. New example files: **dstatmth5.e**, **glmnormalh5.e**, **plotaddsurf1.e**, **plotadderrorbar1.e**, **plotadderrorbar2.e**, **plotarea_ci_latex.e**, **plotcontour2.e** and **plotxy_latex1.e**.

16.0.5

1. Bug fixes

16.0.4

1. Data Import Wizard now supports GAUSS Data sets (*.dat, *.fmt, *.fst)
2. Debugger now supports loading previous stack frames and viewing frame-specific symbols
3. Improved breakpoint/bookmark behavior
4. Improved file opening behavior from finder on OSX
5. Bug fix: Fix various memory leaks
6. Bug fix: Support plotAddArea/plotAddBar to existing time-series plots
7. Bug fix: Fix following symlinks for file paths.
8. Bug fix: Fix 'Find Usages' for struct members.
9. Bug fix: Support debug tooltips for struct members
10. Bug fix: Fix 3D preview, title rendering and exporting for OSX Retina devices
11. Bug fix: Fix previous document shortcut not activating for Windows/Linux
12. Bug fix: Remove 'Delete' option from context menu for undeleteable items in graphics page
13. Bug fix: Improve autocomplete for structs and struct reference arguments in procs

16.0.3

1. New function **csvWriteM** writes data to a delimited text file from a **GAUSS** matrix.
2. Added ability to toggle bolding of functions in source page.
3. Bug fix: **plotPolar** now supports line symbols.
4. Bug fix: **plotAddAnnotation** did not maintain correct z-order for added annotations.
5. Bug fix: Fix 3D graph export dialog starting size constraints.
6. Bug fix: Fix 'Properties' context menu item on graphics page not coming to top.
7. Bug fix: Fix semi-colons in strings, causing Format Text (code formatting) option to add line break inside string.
8. Bug fix: Fix starting indent keywords in source page being case-sensitive.
9. Bug fix: Properly display long error messages that wrap.
10. Added internal check for, and removal of, completely zero imaginary portion of a complex matrix on input to function **lu**.
11. Bug fix: Add complex support for function **lu** on Mac (already supported on Windows and Linux).
12. Bug fix: Fix memory leak in function 'threadfor' in some specific situations.

16.0.2

1. Added ability for **csvReadM/csvReadSA** to read data from the standard input stream (stdin). Pass **__STDIN** as the filename argument to **csvReadM/csvReadSA**.
2. Added ability for **fgets/fgetsa/fgetsat/fgetst** to read from the standard input stream (stdin) . Pass **__STDIN** as the file handle to **fgets/fgetsa/fgetsat/fgetst**.
3. Added ability for **fputs/fputst** to send data to the standard error stream (stderr) and the standard output stream (stdout). Pass **__STDERR** or **__STDOUT** as the file handle to **fputs/fputst**.
4. Changed R-squared calculation in **ols** for regression through the origin to prevent possibility of negative R-squared.
5. Bug fix: **GAUSS** source path was not searched when a file name was passed to **GAUSS** on the command line at start up.
6. Bug fix: **csvReadM** and **csvReadSA** skipped final line in CSV file if the initial

Miscellaneous Topics

line was blank and **csvRead** was told to skip lines.

7. Bug fix: **cons** would continue to return the final buffer data after hitting EOF.

16.0.1

1. Added Data Import Wizard
2. New functions **csvReadM** and **csvReadSA** read data from a delimited text file into a **GAUSS** matrix or string array.
3. New function **glm** calculates the generalized linear model.
4. New function **rescale** provides for scaling columns of a matrix.
5. New function **sampleData** takes samples with or without replacement from a **GAUSS** matrix.
6. New function **qz** computes the sorted complex QZ decomposition.
7. New function **plotSetAxesPen** sets the color and line thickness of the axes line.
8. New functions added for data recoding/reclassification: **reclassify** and **reclassifyCuts**.
9. Added the following new statistical distribution functions: **pdfBinomial**, **pdfPoisson**, **cdfHyperGeo**, **pdfHyperGeo** and **rndHyperGeo**.
10. New function **integrate1d** uses adaptive quadrature to integrate a user-defined function over a specified range.
11. Added new compiler command **#ifmac** to designate code blocks to be compiled and run only on a Mac.
12. Added additional, optional argument to **rndi** to specify the range of random integers produced.
13. Added option to pass additional data to integration functions **intquad1**, **intquad2** and **intquad3**.
14. Added additional, optional input to **lapgschur** to specify sorting of the eigenvalues.
15. Add additional, optional argument to **strsplit** to specify delimiter.
16. Significant speed up to **svd**, **svd1**, **svd2**, **svds**, **svdcusv** and **svdusv**.
17. Significant speed up to **indnv**.
18. Added ability to pass a variable number of arguments to **GAUSS** procedures.
19. Removed requirement to use a DS structure, added option to directly pass a variable number of matrices and made control structure optional for **eqsolve**, **qnewtonmt**, and **sqpsolve**.

20. Removed requirement to use DS structure and added option to directly pass matrices to all gradient and hessian functions (**gradMT**, **gradMTm**, **gradMTT**, **hessMT**, **hessMTm**, **hessMTg**, **hessMTgw**, **hessMTT**, etc).
21. Removed requirement to pass control structure to *dstatmt*.
22. Made inputs other than file name optional for **xlsReadM**, **xlsReadSA**, **xlsWrite**, **xlsWriteM**, **xlsWriteSA**, **xlsGetSheetSize**, **spreadSheetReadM**, **spreadSheetReadSA**, and **spreadSheetWrite**.
23. F4 hot-key will now run the current statement and then skip to the next in addition to running highlighted text.
24. Improved integration of source editor and debugger. Project view window, 'find usages', editing source and other source editor features are available on debug page.
25. Added multiple new preference options to Tools->Preferences->Debug page to control opening and closing of temporary files and other debug page behavior.
26. New preference option added to activate autocomplete only manually on Source Page (with CTRL+Space).
27. Added autocomplete and tooltips to program input/output window.
28. Added preference to Tools->Preferences->Command to activate autocomplete only manually (with CTRL+Space).
29. Added bolding and separate color control syntax highlighting for all **GAUSS** and user defined procedures (Tools->Preferences->Source Page->Functions).
30. Added support for CTRL+E to open a symbol selected in the program input/output window into a floating symbol editor.
31. Added support for F4 to run highlighted text in program input/output window.
32. Added sysstate cases to assess variable arguments passed in to a **GAUSS** procedure as '...'.
33. Symbol editors remember format preferences until closed instead of using default preferences whenever refreshed.
34. Speed up for load time of **GAUSS** when very large folders are open in project view window.
35. Improved behavior of file associations on Mac.
36. Autocomplete no longer pops up when deleting characters or in the middle of a word.
37. Bug fix for display of gaps between bars of a histogram when using **plotAddHistP** in some cases.

Miscellaneous Topics

38. Bug fix for situation in which a message box could be hidden and unreachable behind a floating symbol editor.
39. Bug fix: autocomplete pop-up window no longer stays visible when page loses focus.
40. Bug fix: 'find usages' did not find instances of variables that were index assigns (i.e. `x[5] = 7;`).
41. Bug fix: Dock widgets incorrectly reset to minimum width in some instances of page change and restart **GAUSS**.
42. New example programs: `glmbinomial1.e`, `glmbinomial2.e`, `glmbinomial3.e`, `glmgamma1.e`, `glmgamma2.e`, `glmgamma3.e`, `glmnormal1.e`, `glmnormal2.e`, `glmpoisson1.e`, `glmpoisson2.e`, `qnewtonmt2.e`, `qnewtonmt3.e`, `qnewtonmt4.e`, `sqpsolvemt1.e`, `sqpsolvemt_nlls.e`, `sqpsolvemt_frontier.e`.

36.2 Common Errors

The following is a list of the some common **GAUSS** programming errors, with a description of their cause and resolution:

1. G0064 Operand missing

An operand is what an operator such as `*`, `+`, `/` acts upon. For example in the statement:

```
x = 2 / 3;
```

both 2 and 3 are operands. “Operand missing”, means that there is an operator that does not have all the of “operands” that it needs to do its job. The statement:

```
x = / 3;
```

will cause the error operand missing, because the slash operator (`/`) needs two operands upon which to operate.

This error commonly occurs with print statements or “implicit print” statements. The GAUSS keyword `print`, takes as input a space separated list of items to print. For example:

```
//Print three separate items 3, 4 and 5
print 3 4 5;
```

gives the print keyword the list (3 4 5) as its input. So the statement:

```
//Print three separate items 2, / and 3
print 2 / 3;
```

also gives the print statement a space separated list with 3 items (2, /, 3). `print` can be successful with the first item, the number 2, but when it tries to print the second item in the list "/", it fails. In this case, "/" is an operator without any operands. You can fix this error by changing:

```
//Returns error 'operand missing' when trying to print
//the operator, '/', as a separate token
print 2 / 3;
```

to either one of the following:

```
//Surround statement with parentheses
print (2 / 3);

//Reduce to one statement by removing spaces
print 2/3;
```

Note that you CAN use spaces in assignments. For example:

```
x = 2 / 3;
```

is valid. It is only the `print` keyword that takes a space separated list of items.

2. G0156: Illegal redefinition of procedure 'a'

As GAUSS is compiling a file, which converts the text in the file into byte code that the can be executed, if it sees a symbol that has not been defined, it assumes that this symbol is a matrix. It makes this assumption because procedures can be defined after they are referenced. For example, this code is perfectly fine in GAUSS:

```
//Define variables
a = 3;
b = 4;

//Call procedure
c = hypotenuse(a, b);

//Legally define procedure after it is called
proc (1) = hypotenuse(a, b);
retp(sqrt(a.^2 + b.^2));
endp;
```

The code below, however, is not:

```
//Clear all variables from GAUSS workspace
new;

//Call procedure with undefined variables
c = hypotenuse(a, b);

//Define variables after they are used,
//causing an error
a = 3;
b = 4;

proc (1) = hypotenuse(a, b);
retp(sqrt(a.^2 + b.^2));
endp;
```


This is because variables, ‘a’ and ‘b’ in this case, must be defined before they are referenced. The error means that some symbol is being referenced before it is used.

3. G0159: Wrong number of parameters

In this case, parameters refer to the inputs (also called arguments) to a GAUSS procedure. For example, if we try to pass 3 inputs to **hypotenuse** procedure defined above, like this:

```
//Call procedure with too many parameters  
c = hypotenuse(3, 4, 5);
```

we will get the error ‘wrong number of parameters’, because we have passed three inputs to a procedure that only takes two inputs.

4. G0527: Fewer returns than targets

The returns from a GAUSS procedure or function are called ‘targets’. In the code:

```
x = ones(2,2);
```

the variable *x* is the target. The **ones** function provides one return, a matrix of ones, which matches our one target. However, if we change the code to:

```
{ x, y } = ones(2,2);
```

we now have two targets (*x* and *y*), but **ones** still provides only one return. This will cause the error ‘fewer returns than targets’.

Miscellaneous Topics

5. G0181 : Illegal assignment - type mismatch

Most types in GAUSS, such as matrices and strings, are weakly typed. This means that you can change a matrix to a string or a string to a matrix with a simple reassignment like this:

```
//Define 'x' to be a 1x1 matrix
x = 5;

//Legally redefine 'x' to be a string
x = "my string";
```

However, sparse matrices and structures are strongly typed. This means that once a variable is declared to be a sparse matrix or a specific structure it may not be reassigned to a different type. For example:

```
//Declare 'a' to be a sparse matrix
sparse matrix a;

//Illegally attempt to define 'a' to be a
//dense matrix with an assignment statement
a = { 1 2,
      3 4 };
```

will return the error “Illegal assignment - type mismatch” as will the code below:

```
//Declare 'myPlot' to be a plotControl structure
struct plotControl myPlot;

//Illegally attempt to redefine 'myPlot'
//a matrix with an assignment statement
myPlot = 5;
```

To resolve this error, you can: use the `new` command to clear out all variables, delete just that particular variable, or rename the variable. Often times, a variable

declared to be structure or sparse matrix in a previously run program will cause this conflict in a later program. Simply adding a `new` statement to the top of your program will prevent this problem.

36.3 Error Messages

The following is a list of error messages intrinsic to the **GAUSS** programming language. Error messages generated by library functions are not included here.

G0002 File too large

G0003 Indexing a matrix as a vector

A single index can be used only on vectors. Vectors have only one row or only one column.

G0004 Compiler stack overflow - too complex

An expression is too complex. Break it into smaller pieces. Notify Aptech Systems.

G0005 File is already compiled

G0006 Statement too long

Statement longer than 4000 characters.

G0007 End of file encountered

G0008 Syntax error

Compiler	Unrecognizable or incorrect syntax. Semicolon missing on previous statement.
<code>create</code>	Unrecognizable statement in command file, or <code>numvar</code> or <code>outvar</code> statement error.

G0009 Compiler pass out of memory

Compiler pass has run out of memory. Notify Aptech Systems.

G0010 Can't open output file**G0011 Compiled file must have correct extension**

GAUSS requires a `.gcg` extension.

G0012 Invalid drive specifier**G0013 Invalid filename****G0014 File not found****G0015 Directory full****G0016 Too many #include's**

`#include`'d files are nested too deep.

G0017 WARNING: local outside of procedure

A `local` statement has been found outside a procedure definition. The `local` statement will be ignored.

G0018 Read error in program file**G0019 Can't edit .gcg file****G0020 Not implemented yet**

Command not supported in this implementation.

G0021 use must be at the beginning of a program**G0022 User keyword cannot be used in expression**

G0023 Illegal attempt to redefine symbol to an index variable**G0024 Invalid use of ->, probably should be .****G0025 Undefined symbol**

A symbol has been referenced that has not been given a definition.

G0026 Too many symbols

The global symbol table is full. (To set the limit, see [new](#) in the GAUSS LANGUAGE REFERENCE.)

G0027 Invalid directory**G0028 Can't open configuration file**

GAUSS cannot find the configuration file.

G0029 Missing left parenthesis**G0030 Insufficient workspace memory**

The space used to store and manipulate matrices and strings is not large enough for the operations attempted. (To make the main program space smaller and reclaim enough space to continue, see [new](#) in the GAUSS LANGUAGE REFERENCE.)

G0031 Execution stack too deep - expression too complex

An expression is too complex. Break it into smaller pieces. Notify Aptech Systems.

G0032 fn function too large**G0033 Missing right index bracket****G0034 Missing arguments****G0035 Argument too large**

G0036 Matrices are not conformable

For a description of the function or operator being used and conformability rules, see **Matrix Operators**, Section 1.1, or the **GAUSS LANGUAGE REFERENCE MANUAL**.

G0037 Result too large

The size of the result of an expression is greater than the limit for a single matrix.

G0038 Not all the eigenvalues can be computed**G0039 Matrix must be square to invert****G0040 Not all the singular values can be computed****G0041 Argument must be scalar**

A matrix argument was passed to a function that requires a scalar.

G0042 Matrix must be square to compute determinant**G0043 Not implemented for complex matrices****G0044 Matrix must be real****G0045 Attempt to write complex data to real data set**

Data sets, unlike matrices, cannot change from real to complex after they are created. Use `create complex` to create a complex data set.

G0046 Columns don't match

The matrices must have the same number of columns.

G0047 Rows don't match

The matrices must have the same number of rows.

G0048 Matrix singular

The matrix is singular using the current tolerance.

G0049 Target matrix not complex

G0050 Out of memory for program

The main program area is full. (To increase the main program space, see [new](#) in the GAUSS LANGUAGE REFERENCE.)

G0051 Program too large

The main program area is full. (To increase the main program space, see [new](#) in the GAUSS LANGUAGE REFERENCE.)

G0052 No square root - negative element

G0053 Illegal index

An illegal value has been passed in as a matrix index.

G0054 Index overflow

An illegal value has been passed in as a matrix index.

G0055 retp outside of procedure

A [retp](#) statement has been encountered outside a procedure definition.

G0056 Too many active locals

The execution stack is full. There are too many local variables active. Restructure your program. Notify Aptech Systems.

G0057 Procedure stack overflow - expression too complex

The execution stack is full. There are too many nested levels of procedure calls. Restructure your program. Notify Aptech Systems.

G0058 Index out of range

Miscellaneous Topics

You have referenced a matrix element that is out of bounds for the matrix being referenced.

G0059 exec command string too long

G0060 Nonscalar index

G0061 Cholesky downdate failed

G0062 Zero pivot encountered

crout The Crout algorithm has encountered a diagonal element equal to 0. Use **croutp** instead.

G0063 Operator missing

An expression contains two consecutive operands with no intervening operator.

G0064 Operand missing

An expression contains two consecutive operators with no intervening operand.

G0065 Division by zero!

G0066 Must be recompiled under current version

You are attempting to use compiled code from a previous version of **GAUSS**. Recompile the source code under the current version.

G0068 Program compiled under GAUSS-386 real version

G0069 Program compiled under GAUSS-386i complex version

G0070 Procedure calls too deep

You may have a runaway recursive procedure.

G0071 Type mismatch

You are using an argument of the wrong data type (e.g., inputting a matrix when a string is called for).

G0072 Too many files open

The limit on simultaneously open files is 10.

G0073 Redefinition of

`declare` An attempt has been made to initialize a variable that is already initialized. This is an error when `declare :=` is used. `declare !=` or `declare ?=` may be a better choice for your application.

`declare` An attempt has been made to redefine a string as a matrix or procedure, or vice versa. delete the symbol and try again. If this happens in the context of a single program, you have a programming error. If this is a conflict between different programs, use a `new` statement before running the second program.

`let` A string is being forced to type matrix. Use an **`external matrix symbol;`** statement before the `let` statement.

G0074 Can't run program compiled under GAUSS Light

G0075 gscroll input vector the wrong size

G0076 Call Aptech Systems Technical Support

G0077 New size cannot be zero

You cannot **`reshape`** a matrix to a size of zero.

G0078 vargetl outside of procedure

Miscellaneous Topics

G0079 varputl outside of procedure

G0080 File handle must be an integer

G0081 Error renaming file

G0082 Error reading file

G0083 Error creating temporary file

G0084 Too many locals

A procedure has too many local variables.

G0085 Invalid file type

You cannot use this kind of file in this way.

G0086 Error deleting file

G0087 Couldn't open

The auxiliary output file could not be opened. Check the file name and make sure there is room on the disk.

G0088 Not enough memory to convert the whole string

G0089 WARNING: duplicate definition of local

G0090 Label undefined

Label referenced has no definition.

G0091 Symbol too long

Symbols can be no longer than 32 characters.

G0092 Open comment

A comment was never closed.

G0093 Locate off screen**G0094 Argument out of range****G0095 Seed out of range****G0096 Error parsing string**

`parse` encountered a token that was too long.

G0097 String not closed

A string must have double quotes at both ends.

G0098 Invalid character for imaginary part of complex number**G0099 Illegal redefinition of user keyword****G0100 Internal E R R O R ###**

Notify Aptech Systems.

G0101 Argument cannot be zero

The argument to `ln` or `log` cannot be zero.

G0102 Subroutine calls too deep

Too many levels of `gosub`. Restructure your program.

G0103 return without gosub

You have encountered a subroutine without executing a `gosub`.

G0104 Argument must be positive**G0105 Bad expression or missing arguments**

Check the expression in question, or you forgot an argument.

Miscellaneous Topics

G0106 Factorial overflow

G0107 Nesting too deep

Break the expression into smaller statements.

G0108 Missing left bracket [

G0109 Not enough data items

You omitted data in a `let` statement.

G0110 Found) expected] -

G0111 Found] expected) -

G0112 Matrix multiplication overflow

G0113 Unclosed (

G0114 Unclosed [

G0115 Illegal redefinition of function

You are attempting to turn a function into a matrix or string. If this is a name conflict, `delete` the function.

G0116 sysstate: invalid case

G0117 Invalid argument

G0118 Argument must be integer

File handles must be integral.

G0120 Illegal type for save

G0121 Matrix not positive definite

The matrix is either not positive definite, or singular using the current tolerance.

G0122 Bad file handle

The file handle does not refer to an open file or is not in the valid range for file handles.

G0123 File handle not open

The file handle does not refer to an open file.

G0124 readr call too large

You are attempting to read too much in one call.

G0125 Read past end of file

You have already reached the end of the file.

G0126 Error closing file**G0127 File not open for write****G0128 File already open****G0129 File not open for read****G0130 No output variables specified****G0131 Can't create file, too many variables****G0132 Can't write, disk probably full****G0133 Function too long****G0134 Can't seekr in this type of file****G0135 Can't seek to negative row****G0136 Too many arguments or misplaced assignment operator**

Miscellaneous Topics

You have an assignment operator (=) where you want a comparison operator (==), or you have too many arguments.

G0137 Negative argument - erf or erfc

G0138 User keyword must have one argument

G0139 Negative parameter - Incomplete Beta

G0140 Invalid second parameter - Incomplete Beta

G0141 Invalid third parameter - Incomplete Beta

G0142 Nonpositive parameter - gamma

G0143 NaN or missing value - cdfchic

G0144 Negative parameter - cdfchic

G0145 Second parameter < 1.0 - cdfchic

G0146 Parameter too large - Incomplete Beta

G0147 Bad argument to trig function

G0148 Angle too large to trig function

G0149 Matrices not conformable

For a description of the function or operator being used and conformability rules, see **Matrix Operators**, Section 1.1, or the GAUSS LANGUAGE REFERENCE.

G0150 Matrix not square

G0151 Sort failure

G0152 Variable not initialized

You have referenced a variable that has not been initialized to any value.

G0153 Unsuccessful close on auxiliary output

The disk may be full.

G0154 Illegal redefinition of string**G0155 Nested procedure definition**

A `proc` statement was encountered inside a procedure definition.

G0156 Illegal redefinition of procedure

You are attempting to turn a procedure into a matrix or string. If this is a name conflict, delete the procedure.

G0157 Illegal redefinition of matrix**G0158 endp without proc**

You are attempting to end a procedure that you never started.

G0159 Wrong number of parameters

You called a procedure with the wrong number of arguments.

G0160 Expected string variable**G0161 User keywords return nothing****G0162 Can't save proc/keyword/fn with global references**

Remove the global references or leave this in source code form for the autoloader to handle. (See `library` in the GAUSS LANGUAGE REFERENCE.)

G0163 Wrong size format matrix**G0164 Bad mask matrix****G0165 Type mismatch or missing arguments****G0166 Character element too long**

Miscellaneous Topics

The maximum length for character elements is 8 characters.

G0167 Argument must be column vector

G0168 Wrong number of returns

The procedure was defined to return a different number of items.

G0169 Invalid pointer

You are attempting to call a local procedure using an invalid procedure pointer.

G0170 Invalid use of ampersand

G0171 Called symbol is wrong type

You are attempting to call a local procedure using a pointer to something else.

G0172 Can't resize temporary file

G0173 varindx failed during open

The global symbol table is full.

G0174 "." and " " operators must be inside [] brackets

These operators are for indexing matrices.

G0175 String too long to compare

G0176 Argument out of range

G0177 Invalid format string

G0178 Invalid mode for getf

G0179 Insufficient heap space

G0180 Trim too much

You are attempting to trim more rows than the matrix has.

G0181 Illegal assignment - type mismatch**G0182 2nd and 3rd arguments different order****G0274 Invalid parameter for conv****G0275 Parameter is NaN (Not A Number)**

The argument is a NaN (see **Special Data Types**, Section 1.0.1).

G0276 Illegal use of reserved word**G0277 Null string illegal here****G0278 proc without endp**

You must terminate a procedure definition with an `endp` statement.

G0286 Multiple assign out of memory**G0287 Seed not updated**

The seed argument to `rndns` and `rndus` must be a simple local or global variable reference. It cannot be an expression or constant. These functions are obsolete, please use `rndlcn` and `rndlcu`

G0288 Found break not in do loop**G0289 Found continue not in do loop****G0290 Library not found**

The specified library cannot be found on the `lib_path` path. Make sure installation was correct.

G0291 Compiler pass out of memory

Notify Aptech Systems.

G0292 File listed in library not found

Miscellaneous Topics

A file listed in a library could not be opened.

G0293 Procedure has no definition

The procedure was not initialized. Define it.

G0294 Error opening temporary file

One of the temporary files could not be opened. The directory may be full.

G0295 Error writing temporary file

One of the temporary files could not be written to. The disk may be full.

G0296 Can't raise negative number to nonintegral power

G0300 File handle must be a scalar

G0301 Syntax error in library

G0302 File has been truncated or corrupted

getname	File header cannot be read.
load	Cannot read input file, or file header cannot be read.
open	File size does not match header specifications, or file header cannot be read.

G0317 Can't open temp file

G0336 Disk full

G0339 Can't debug compiled program

G0341 File too big

G0347 Can't allocate that many globals

G0351 Warning: Not reinitializing : declare ?=

The symbol is already initialized. It will be left as is.

G0352 Warning: Reinitializing : declare !=

The symbol is already initialized. It will be reset.

G0355 Wrong size line matrix**G0360 Write error****G0364 Paging error****G0365 Unsupported executable file type****G0368 Unable to allocate translation space****G0369 Unable to allocate buffer****G0370 Syntax Error in code statement****G0371 Syntax Error in recode statement****G0372 Token verify error**

Notify Aptech Systems.

G0373 Procedure definition not allowed

A procedure name appears on the left side of an assignment operator.

G0374 Invalid make statement**G0375 make Variable is a Number****G0376 make Variable is Procedure****G0377 Cannot make Existing Variable****G0378 Cannot make External Variable****G0379 Cannot make String Constant****G0380 Invalid vector statement**

Miscellaneous Topics

G0381 vector Variable is a Number

G0382 vector Variable is Procedure

G0383 Cannot vector Existing Variable

G0384 Cannot vector External Variable

G0385 Cannot vector String Constant

G0386 Invalid extern statement

G0387 Cannot extern number

G0388 Procedures always external

A procedure name has been declared in an `extern` statement. This is a warning only.

G0389 extern variable already local

A variable declared in an `extern` statement has already been assigned local status.

G0390 String constant cannot be external

G0391 Invalid code statement

G0392 code Variable is a Number

G0393 code Variable is Procedure

G0394 Cannot code Existing Variable

G0395 Cannot code External Variable

G0396 Cannot code String Constant

G0397 Invalid recode statement

G0398 recode Variable is a Number

G0399 recode Variable is Procedure

G0400 Cannot recode External Variable

G0401 Cannot recode String Constant

G0402 Invalid keep statement

G0403 Invalid drop statement

G0404 Cannot define Number

G0405 Cannot define String

G0406 Invalid select statement

G0407 Invalid delete statement

G0408 Invalid outtyp statement

G0409 outtyp already defaulted to 8

Character data has been found in the output data set before an `outtyp 2` or `outtyp 4` statement. This is a warning only.

G0410 outtyp must equal 2, 4, or 8

G0411 outtyp override...precision set to 8

Character data has been found in the output data set after an `outtyp 2` or `outtyp 4` statement. This is a warning only.

G0412 default not allowed in recode statement

`default` allowed only in `code` statement.

G0413 Missing file name in dataloop statement

Miscellaneous Topics

G0414 Invalid listwise statement

G0415 Invalid lag statement

G0416 lag variable is a number

G0417 lag variable is a procedure

G0418 Cannot lag External Variable

G0419 Cannot lag String Constant

G0421 Command not supported in Run-Time Module

G0428 Cannot use debug command inside program

G0429 Invalid number of subdiagonals

G0431 Error closing dynamic library

G0432 Error opening dynamic library

G0433 Cannot find DLL function

G0434 Error opening default dynamic library

G0435 Invalid mode

G0436 Matrix is empty

G0437 loadexe not supported; use dlibrary instead

G0438 callexe not supported; use dllcall instead

G0439 File has wrong bit order

G0440 File has wrong byte order

G0441 Type vector malloc failed

- G0442 No type vector in gfblock**
 - G0445 Illegal left-hand side reference in procedure**
 - G0446 Argument is the wrong size**
 - G0447 vfor called with illegal loop level**
 - G0454 Failure opening printer for output**
 - G0456 Failure buffering output for printer**
 - G0457 Cannot take log of a negative number**
 - G0458 Attempt to index proc/fn/keyword as a matrix**
 - G0459 Missing right brace**
 - G0460 Unexpected end of statement**
 - G0461 Too many data items**
 - G0462 Negative trim value**
 - G0463 Failure generating graph**
 - G0465 Redefinition of structure, number of elements**
 - G0466 Redefinition of structure, type mismatch**
 - G0467 Redefinition of structure, unrecognized member**
 - G0468 Structure definition inside procedure definition**
 - G0469 Cannot create translator temp file**
 - G0470 Symbol not found**
 - G0472 Invalid name**
-

Miscellaneous Topics

G0473 String not terminated with null byte

G0477 FOR loops nested too deep

G0486 Character argument too long

G0487 License expired

G0490 License manager initialization error

G0491 License manager error

G0492 Licensing failure

G0497 Missing right parenthesis

G0500 Cannot create temporary filename

G0503 Cannot assign matrix to scalar member

G0504 Invalid structure member

G0505 Invalid structure redefinition

G0506 Structure assignment mismatch

G0507 Undefined structure

G0508 Structure argument mismatch

G0509 Too many structure members

G0510 Duplicate name for structure member

G0514 Not supported for structures

G0515 Too many values in locator

G0516 Too many dimensions in result

- G0517 Too many dimensions in argument**
 - G0518 Not implemented for complex**
 - G0519 Illegal dereference of structure array**
 - G0520 Arguments not conformable**
 - G0521 Argument must be real**
 - G0522 Illegal indexing of dereferenced structure**
 - G0523 Numeric argument must be integer**
 - G0524 Found comma, expecting index**
 - G0525 Argument contains NaNs**
 - G0526 Argument must be compact format**
 - G0529 Array orders must be ≥ 1**
 - G0531 Two trailing dimensions of argument must be the same size**
 - G0532 Both dimensions of argument must be the same size**
 - G0533 1-dimensional argument must contain only 1 element**
 - G0534 Cannot create file**
 - G0538 Zero illegal in for loop increment**
 - G0541 Illegal assignment to FOR loop counter**
 - G0542 Object too large for 32-bit version**
 - G0543 Array has too many dimensions for matrix assign**
 - G0547 Array not conformable for indexing**
-

Miscellaneous Topics

G0548 Array not conformable for boolean operation

G0549 Global structure pointer cannot point to local structure

G0550 Invalid use of *

G0551 Feature not authorized

G0553 Path too long

G0554 Unable to create sparse matrix

G0555 Cannot index uninitialized structure

G0556 #IF nesting limit exceeded

G0557 #ELSE without #IF

G0558 #ENDIF without #IF

G0559 Symbol not #DEFINE'd

G0560 Too many #DEFINE's

G0561 Duplicate #DEFINE

G0562 Open `/* */` comment

G0563 Open `@ @` comment

G0564 Illegal redefinition of sparse matrix

G0565 Initializer too large, increase maxdecret in config (.cfg) file

G0566 Can't create profiler data file

G0567 Sparse matrix uninitialized

G0568 Operation not defined for triangular, symmetric, or Hermitian sparse matrix

G0569 Argument must be complex

G0570 Diagonal must be real

G0571 Diagonal must not contain zeros

G0572 Argument must be triangular

G0573 Argument must be symmetric

G0574 Sparse type mismatch

G0575 Unable to load variable

G0576 Threading error

G0577 Expected THREADSTAT, THREADBEGIN, or THREADJOIN

G0578 A THREADJOIN failed

G0579 Cannot call RUN from inside thread

G0580 Unable to converge in allowed number of iterations

G0581 Incorrect Argument: Number of eigenvalues must be positive

G0582 Incorrect Argument: Number of column vectors must be \geq number of eigenvalues +2 and $<$ rows of input matrix

G0583 Could not apply shift during an Arnoldi iteration cycle. Try increasing size of ncv

Miscellaneous Topics

G0584 Invalid Input: 'which' must be 'LM' 'SM' 'LR' 'LI' 'SR' or 'SI' and type string

G0585 Error Return from LAPACK eigenvalue calculation

G0586 dneupd error 1: contact Aptech Systems

G0587 Input matrix must be sparse

G0588 Incorrect Input: Number of eigenvalues must be scalar

G0589 Incorrect Input: Tolerance must be scalar

G0590 No eigenvalues found to specified tolerance in allowed iterations

G0591 Incorrect Input: Max iterations must be scalar

G0592 Incorrect Input: Number of column vectors must be scalar

G0593 Incorrect Input: Third input, probability, must be > 0 and < 1

G0594 Incorrect Input: Number of successes (input 1) must be less than number of trials (input 2)

G0595 Incorrect Input: State vector cannot have more than 1 column

G0596 Incorrect Input: Inputs 1 and 2 (cols and rows) must be scalar or 1x1 matrix

G0597 Incorrect Input: Input must be dense matrix

G0598 Incorrect Input: First input may have 1 column only

G0599 Incorrect Input: Input 2 may not have more columns that input 1 has rows

G0600 Incorrect Input: Input 1 must be square

G0601 Incorrect Input: Input 2 must be square

G0602 Incorrect Input: $1 \leq il < iu$ and $iu \leq \text{rows of } x$

G0603 Failure to converge

36.4 GAUSS Graphics Colors

The following is a chart and list of colors available for use in **GAUSS** graphics with both the name and RGB value listed.

Miscellaneous Topics

	Color Name	RGB Value
	aliceblue	240, 248, 255
	antiquewhite	250, 235, 215
	aqua	0, 255, 255
	aquamarine	127, 255, 212
	azure	240, 255, 255
	beige	245, 245, 220
	bisque	255, 228, 196
	black	0, 0, 0
	blanchedalmond	255, 235, 205
	blue	0, 0, 255
	blueviolet	138, 43, 226
	brown	165, 42, 42
	burlywood	222, 184, 135
	cadetblue	95, 158, 160
	chartreuse	127, 255, 0
	chocolate	210, 105, 30
	coral	255, 127, 80
	cornflowerblue	100, 149, 237
	cornsilk	255, 248, 220
	crimson	220, 20, 60
	cyan	0, 255, 255
	darkblue	0, 0, 139
	darkcyan	0, 139, 139
	darkgoldenrod	184, 134, 11
	darkgray	169, 169, 169
	darkgreen	0, 100, 0
	darkgrey	169, 169, 169
	darkkhaki	189, 183, 107
	darkmagenta	139, 0, 139
	darkolivegreen	85, 107, 47
	darkorange	255, 140, 0
	darkorchid	153, 50, 204
	darkred	139, 0, 0
	darksalmon	233, 150, 122
	darkseagreen	143, 188, 143
	darkslateblue	72, 61, 139
	darkslategray	47, 79, 79
	darkslategrey	47, 79, 79
	darkturquoise	0, 206, 209
	darkviolet	148, 0, 211
	deeppink	255, 20, 147
	deepskyblue	0, 191, 255
	dimgray	105, 105, 105

	Color Name	RGB Value
	dimgrey	105, 105, 105
	dodgerblue	30, 144, 255
	firebrick	178, 34, 34
	floralwhite	255, 250, 240
	forestgreen	34, 139, 34
	fuchsia	255, 0, 255
	gainsboro	220, 220, 220
	ghostwhite	248, 248, 255
	gold	255, 215, 0
	goldenrod	218, 165, 32
	gray	128, 128, 128
	grey	128, 128, 128
	green	0, 128, 0
	greenyellow	173, 255, 47
	honeydew	240, 255, 240
	hotpink	255, 105, 180
	indianred	205, 92, 92
	indigo	75, 0, 130
	ivory	255, 255, 240
	khaki	240, 230, 140
	lavender	230, 230, 250
	lavenderblush	255, 240, 245
	lawngreen	124, 252, 0
	lemonchiffon	255, 250, 205
	lightblue	173, 216, 230
	lightcoral	240, 128, 128
	lightcyan	224, 255, 255
	lightgoldenrodyellow	250, 250, 210
	lightgray	211, 211, 211
	lightgreen	144, 238, 144
	lightgrey	211, 211, 211
	lightpink	255, 182, 193
	lightsalmon	255, 160, 122
	lightseagreen	32, 178, 170
	lightskyblue	135, 206, 250
	lightslategray	119, 136, 153
	lightslategrey	119, 136, 153
	lightsteelblue	176, 196, 222
	lightyellow	255, 255, 224
	lime	0, 255, 0
	limegreen	50, 205, 50
	linen	250, 240, 230
	magenta	255, 0, 255

Miscellaneous Topics

<i>Color Name</i>	<i>RGB Value</i>
maroon	128, 0, 0
mediumaquamarine	102, 205, 170
mediumblue	0, 0, 205
mediumorchid	186, 85, 211
mediumpurple	147, 112, 219
mediumseagreen	60, 179, 113
mediumslateblue	123, 104, 238
mediumspringgreen	0, 250, 154
mediumturquoise	72, 209, 204
mediumvioletred	199, 21, 133
midnightblue	25, 25, 112
mintcream	245, 255, 250
mistyrose	255, 228, 225
moccasin	255, 228, 181
navajowhite	255, 222, 173
navy	0, 0, 128
oldlace	253, 245, 230
olive	128, 128, 0
olivedrab	107, 142, 35
orange	255, 165, 0
orangered	255, 69, 0
orchid	218, 112, 214
palegoldenrod	238, 232, 170
palegreen	152, 251, 152
paleturquoise	175, 238, 238
palevioletred	219, 112, 147
papayawhip	255, 239, 213
peachpuff	255, 218, 185
peru	205, 133, 63
pink	255, 192, 203
plum	221, 160, 221
powderblue	176, 224, 230
purple	128, 0, 128
red	255, 0, 0
rosybrown	188, 143, 143
royalblue	65, 105, 225
saddlebrown	139, 69, 19
salmon	250, 128, 114
sandybrown	244, 164, 96
seagreen	46, 139, 87
seashell	255, 245, 238
sienna	160, 82, 45
silver	192, 192, 192

<i>Color Name</i>	<i>RGB Value</i>
skyblue	135, 206, 235
slateblue	106, 90, 205
slategray	112, 128, 144
slategrey	112, 128, 144
snow	255, 250, 250
springgreen	0, 255, 127
steelblue	70, 130, 180
tan	210, 180, 140
teal	0, 128, 128
thistle	216, 191, 216
tomato	255, 99, 71
turquoise	64, 224, 208
violet	238, 130, 238
wheat	245, 222, 179
white	255, 255, 255
whitesmoke	245, 245, 245
yellow	255, 255, 0
yellowgreen	154, 205, 50

36.5 Obsolete Commands

The following commands will no longer be supported and therefore should not be used when creating new programs.

`color`

`coreleft`

`csrtype`

`denseSubmat`

`dfree`

`disable`

`editm`

`eigcg`

`eigcg2`

`eigch`

`eigch2`

`eigrq`

`eigrq2`

`eigrs`

`eigrs2`

`enable`

`export`

`exportf`

`files`

font
FontLoad
FontUnload
FontUnloadAll
graph
import
importf
isSparse
line
lpos
lprint
lprint on/off
lpwidth
lshow
medit
nametype
ndpchk
ndpclex
ndpcntrl
plot
plotsym

Miscellaneous Topics

`prcsn`

`print on/off`

`rndns`

`rndus`

`scroll`

`setvmode`

`sparseCols`

`sparseEye`

`sparseFD`

`sparseFP`

`sparseHConcat`

`sparseNZE`

`sparseOnes`

`sparseRows`

`sparseScale`

`sparseSet`

`sparseSolve`

`sparseSubmat`

`sparseTD`

`sparseTranspose`

`sparseTrTD`

sparseTScalar

sparseVConcat

spline1d

spline2d

vartype

WinClear

WinClearArea

WinClearTTYlog

WinClose

WinCloseAll

WinGetActive

WinGetAttributes

WinGetColorCells

WinGetCursor

WinMove

WinOpenPQG

WinOpenText

WinOpenTTY

WinPan

WinPrint

WinPrintPQG

Miscellaneous Topics

WinRefresh

WinRefreshArea

WinResize

WinSetActive

WinSetBackground

WinSetColorCells

WinSetColormap

WinSetCursor

WinSetForeground

WinSetRefresh

WinSetTextWrap

WinZoomPQG

36.6 Quick Reference

36.6.1 Basic functions

`new`

remove all
variables
from your
GAUSS
workspace.

`cls`

clear
program
output from
the GAUSS

```
f = filesa(file_pattern);
```

Program
Input/Output
Window.

returns a
string array
containing
all files that
match 'file_
pattern'.

```
chdir new_directory;
```

sets your
GAUSS
current
working
directory to
'new_
directory'.

```
cwd = cdir(0);
```

returns your
GAUSS
current
working
directory.

```
d = getGAUSSHome();
```

returns the
location of
your
GAUSS
installation.

36.6.2 Matrix creation

```
x = { 1 2, 3 4 };
```

create a

Miscellaneous Topics

```
x = seqa(start, step, count);
```

2x2
matrix.
creates a
sequence
of 'count'
numbers
starting at
'start' and
increasing
by 'step'.

```
x = seqm(start, mult, count);
```

creates a
sequence
of 'count'
numbers
starting at
'start' and
increasing
by a
multiple
of 'mult'.

```
x = zeros (m, n);
```

creates an
'm' by 'n'
matrix
with all
elements
set to 0.

```
x = ones (m, n);
```

creates an
'm' by 'n'
matrix
with all
elements

```
x = randn(m, n);
```

set to 1.
creates an
'm' by 'n'
matrix of
random
normal
numbers.

```
x = randu(m, n);
```

creates an
'm' by 'n'
matrix of
uniformly
distributed
random
numbers.

36.6.3 Matrix manipulation

```
a = x[row,col];
```

extract the
element of
'x' located
at 'row:col'.

```
a = x[:,col];
```

extract all
rows of the
specified
column(s)
of 'x'.

```
a = x[row,:];
```

extract all
columns of
the
specified

Miscellaneous Topics

```
a = x[r_start:r_end, .];
```

row(s) of
'x'.

extract all
columns
from the
row range
'row_start'
to 'row_
end'.

```
a = x[a b c, col];
```

extract the
elements
from rows
'a', 'b', and
'c' in
column
'col'.

```
a = x ~ y;
```

horizontally
concatenate
'x' and 'y'.

```
a = x | y;
```

vertically
concatenate
'x' and 'y'.

```
a = reshape(x, m, n);
```

reshape 'x'
to be an 'm'
by 'n'
matrix.

```
a = delrows(x, idx);
```

returns all
rows of 'x'
except
those listed
in 'idx'.


```
a = delif(x, logical);
```

returns all
rows of 'x'
except the
rows that
match a
logical
expression.

```
a = selif(x, logical);
```

returns all
rows of 'x'
that match
a logical
expression.

36.6.4 Operators

Element-by-element (ExE) operators

```
z = x .* y;
```

Element-by-element multiply.

```
z = x ./ y;
```

Element-by-element divide.

```
z = x .^ y;
```

Element-by-element
exponentiation.

```
z = x + y;
```

Element-by-element addition.

```
z = x - y;
```

Element-by-element
subtraction.

Matrix operators

```
z = x * y;
```

Matrix
multiply.

```
b = y / x;
```

Solve a

Miscellaneous Topics

Matrix operators

```
z = x .* y;
```

system of
linear
equations.

```
z = x *~ y;
```

Kronecker
product.

```
z = x';
```

Horizontal
direct
product.

Matrix
transpose.

Scalar logical operators

```
z = x and y;
```

Scalar
logical
AND.

```
z = x or y;
```

Scalar
logical
OR.

Element-by-element (ExE) logical operators

```
z = x .and y;
```

Element-
by-
element
logical
AND.

```
z = x .or y;
```

Element-
by-
element
logical

Element-by-element (ExE) logical operators

`z = x .> y;`

OR.

Element-by-element greater than.

`z = x .< y;`

Element-by-element less than.

`z = x .== y;`

Element-by-element equality test.

`z = x .!= y;`

Element-by-element inequality test.

Matrix logical operators

`z = x > y;`

is every element in 'x' greater than its corresponding element in 'y'.

`z = x < y;`

is every element in 'x' less than its corresponding element in 'y'

`z = x == y;`

does every element of 'x' equal its corresponding element in 'y'.

Miscellaneous Topics

Matrix logical operators

```
z = x != y;
```

is every element in 'x' different than its corresponding element in 'y'.

36.6.5 String creation

```
s = "this is a string";
```

create string containing 'this is a string'.

```
s = "this is " $+ "a string";
```

combine strings.

```
string sa = { "cpi" "ppi", "m1" "m2" };
```

create 2x2 string array.

```
s = ntos(n);
```

convert numeric 'n' to a string.

36.6.6 String array manipulation

```
s = sa[r, c];
```

extract the r, c element of 'sa'.

```
s = sa[., c];
```

extract all rows of the, 'c'th column of 'sa'.

```
s = sa[r, .];
```

extract 'r'th row of 'sa'.

```
sa = "producer" $~  
"prices";
```

horizontal concatenation of strings.

```
sa = "County" $| "State";
```

vertical concatenation of strings.

```
idx = indsav(what,  
where);
```

returns the location of the strings from 'what' in the string array 'where'.

```
su = intrsectsa(s_1, s_2);
```

returns the intersection of the string arrays 's_1' and 's_2'.

36.6.7 Loading and saving data

```
x = xlsReadM(file, range, sheet);
```

reads the specified data into a GAUSS matrix.

```
sa = xlsReadSA(file, range, sheet);
```

reads the specified data into a GAUSS string array.

```
ret = xlsWrite(x, file, range, sheet);
```

writes the data from the GAUSS matrix or string array 'x' into the specified Excel® file.

```
x = loadd(dataset);
```

loads the

Miscellaneous Topics

```
ret = saved(x, dataset, v_names);
```

```
x = csvReadM(file, row_range, col_  
range);
```

```
x = csvReadSA(file, row_range, col_  
range);
```

data
from a
GAUSS
dataset
into a
GAUSS
matrix.

saves the
data
from 'x'
to a
GAUSS
dataset.

reads
data
from a
text
delimited
file, such
as CSV,
to a
GAUSS
matrix.

reads
data
from a
text
delimited
file, such
as CSV,
to a

GAUSS
string
array.

36.7 Reserved Words

The following words are used for **GAUSS** functions. You cannot use these names for variables or procedures in your programs:

A

abs

acf

aconcat

acos

aeye

amax

amean

AmericanBinomCall

AmericanBinomCall_Greeks

AmericanBinomCall_ImpVol

AmericanBinomPut

AmericanBinomPut_Greeks

AmericanBinomPut_ImpVol

AmericanBSCall

AmericanBSCall_Greeks

Miscellaneous Topics

`AmericanBSCall_ImpVol`
`AmericanBSPut`
`AmericanBSPut_Greeks`
`AmericanBSPut_ImpVol`
`amin`
`amult`
`and`
`annotationGetDefaults`
`annotationSetBkd`
`annotationSetFont`
`annotationSetLineColor`
`annotationSetLineStyle`
`annotationSetLineThickness`
`annualTradingDays`
`arccos`
`arcsin`
`arctan`
`arctan2`
`areshape`
`arrayalloc`
`arrayindex`

`arrayinit`
`arraytomat`
`asciiload`
`asclabel`
`asin`
`astd`
`astds`
`asum`
`atan`
`atan2`
`atranspose`
`axmargin`

B

`balance`
`band`
`bandchol`
`bandcholsol`
`bandltsol`
`bandrv`
`bandsolpd`
`bar`
`base10`

Miscellaneous Topics

`begwind``besselj``beta``box``boxcox``break`

C

`calcbox``call``callexe``cdfBeta``cdfBetaInv``cdfBinomial``cdfBinomialInv``cdfBvn``cdfBvn2``cdfBvn2e``cdfCauchy``cdfCauchyInv``cdfChic``cdfChii``cdfChinc`

`cdfEmpirical`
`cdfExp`
`cdfExpInv`
`cdfFc`
`cdfFnc`
`cdfFncInv`
`cdfGam`
`cdfGenPareto`
`cdfHyperGeo`
`cdfLaplace`
`cdfLaplaceInv`
`cdfLogistic`
`cdfLogisticInv`
`cdfMvn`
`cdfMvn2e`
`cdfMvnce`
`cdfMvne`
`cdfMvt2e`
`cdfMvtce`
`cdfMvte`
`cdfn`

Miscellaneous Topics

`cdfN2``cdfNc``cdfNcd``cdfNegBinomial``cdfNegBinomialInv``cdfNi``cdfPoisson``cdfPoissonInv``cdfRayleigh``cdfRayleighInv``cdfTc``cdfTci``cdfTnc``cdfTvn``cdfWeibull``cdfWeibullInv``cdir``ceil``cfft``cffti``ChangeDir`

`chdir`
`checkinterrupt`
`chiBarSquare`
`chol`
`choldn`
`cholsol`
`cholup`
`chrs`
`cint`
`clear`
`clearg`
`close`
`closeall`
`cls`
`cmsplit`
`cmsplit2`
`code`
`color`
`cols`
`colsf`
`combinate`

Miscellaneous Topics

`combined`

`comlog`

`commandeerm`

`commandeersa`

`compile`

`complex`

`con`

`cond`

`conformed`

`conj`

`cons`

`conscore`

`continue`

`contour`

`conv`

`convertsatotr`

`convertstrtosa`

`coreleft`

`corrm`

`corrms`

`corrvc`

corr_x

corr_{xs}

cos

cosh

counts

countwts

create

crossprd

crout

croutp

csr_{col}

csr_{lin}

csr_{type}

csvRead_M

csvRead_{SA}

cumprodc

cumsumc

curve

cv_{tos}

cv_{tosa}

D

Miscellaneous Topics

`datacreate`
`datacreatecomplex`
`datalist`
`dataload`
`dataloop`
`dataopen`
`datasave`
`date`
`datestr`
`datestring`
`datestrymd`
`dayinyr`
`dayOfWeek`
`dbAddDatabase`
`dbClose`
`dbCommit`
`dbCreateQuery`
`dbExecQuery`
`dbGetConnectOptions`
`dbGetDatabaseName`
`dbGetDriverName`

dbGetDrivers
dbGetHostName
dbGetLastErrorNum
dbGetLastErrorText
dbGetNumericalPrecPolicy
dbGetPassword
dbGetPort
dbGetPrimaryIndex
dbGetTableHeaders
dbGetTables
dbGetUserName
dbHasFeature
dbIsDriverAvailable
dbIsOpen
dbIsOpenError
dbNumericalPrecPolicy
>dbOpen
dbQueryBindValue
dbQueryClear
dbQueryCols
dbQueryExecPrepared

Miscellaneous Topics

`dbQueryFetchAllM`
`dbQueryFetchAllSA`
`dbQueryFetchNextM`
`dbQueryFetchNextSA`
`dbQueryFinish`
`dbQueryGetBoundValue`
`dbQueryGetBoundValues`
`dbQueryGetField`
`dbQueryGetLastErrorNum`
`dbQueryGetLastErrorText`
`dbQueryGetLastInsertID`
`dbQueryGetLastQuery`
`dbQueryGetPosition`
`dbQueryIsActive`
`dbQueryIsForwardOnly`
`dbQueryIsNull`
`dbQueryIsSelect`
`dbQueryIsValid`
`dbQueryPrepare`
`dbQueryRows`
`dbQuerySeek`

`dbQuerySeekFirst`
`dbQuerySeekLast`
`dbQuerySeekNext`
`dbQuerySeekPrevious`
`dbQuerySetForwardOnly`
`dbRemoveDatabase`
`dbRollback`
`dbSetConnectOptions`
`dbSetDatabaseName`
`dbSetHostName`
`dbSetPassword`
`dbSetPort`
`dbSetUserName`
`dbTransaction`
`debug`
`declare`
`delete`
`deleteFile`
`delif`
`delrows`
`denseSubmat`

Miscellaneous Topics

`denseToSp`

`denseToSpRE`

`denseToZero`

`design`

`det`

`detl`

`dfft`

`dffti`

`dfree`

`diag`

`diagrv`

`digamma`

`disable`

`dlibrary`

`dllcall`

`do`

`dos`

`doswin`

`DOSWinCloseall`

`DOSWinOpen`

`dot`

dotfeq
dotfeqmt
dotfge
dotfgemt
dotfgt
dotfgtmt
dotfle
dotflemt
dotflt
dotfltmt
dotfne
dotfnemt
draw
dsCreate
dstat
dstatmt
dstatmtControlCreate
dtdate
dtday
dttime
dttodtv

Miscellaneous Topics

`dttostr``dttoutc``dtvnormal``dtvtodt``dtvtoutc``dummy``dummybr``dummydn`

E

`ed``edit``editm``eig``eigcg``eigcg2``eigch``eigch2``eigh``eighv``eigr``eigr2``eigrs`

eigrs2
eigv
elapsedTradingDays
else
elseif
enable
end
endfor
endif
endo
endp
endwind
envget
eof
eq
eqSolve
eqSolvemt
eqSolvemtControlCreate
eqSolvemtOutCreate
eqSolveSet
eqv

Miscellaneous Topics

`erf`

`erfc`

`erfcplx`

`erfccplx`

`erfCInv`

`erfInv`

`error`

`errorlog`

`errorlogat`

`etdays`

`ethsec`

`etstr`

`EuropeanBinomCall`

`EuropeanBinomCall_Greeks`

`EuropeanBinomCall_ImpVol`

`EuropeanBinomPut`

`EuropeanBinomPut_Greeks`

`EuropeanBinomPut_ImpVol`

`EuropeanBSCall`

`EuropeanBSCall_Greeks`

`EuropeanBSCall_ImpVol`

EuropeanBSPut
EuropeanBSPut_Greeks
EuropeanBSPut_ImpVol
exctsmpl
exec
execbg
exp
expr
extern
external
eye

F

fcheckerr
fclearerr
feq
feqmt
fflush
fft
ffti
fftm
fftmi
fftn

Miscellaneous Topics

fge

fgemt

fgets

fgetsa

fgetsat

fgetst

fgt

fgtmt

fileinfo

files

filesa

fix

fle

flemt

floor

flt

fltmt

fmod

fn

fne

fnemt

font
fontload
fonts
fontunload
fontunloadall
fopen
for
format
formatcv
formatnv
fputs
fputst
fseek
fstrerror
ftell
ftocv
ftos
ftostrc

G

gamma
gammacplx
gammaii

Miscellaneous Topics

`gausset`
`gdaAppend`
`gdaCreate`
`gdaDStat`
`gdaDStatMat`
`gdaGetIndex`
`gdaGetName`
`gdaGetNames`
`gdaGetOrders`
`gdaGetType`
`gdaGetTypes`
`gdaGetVarInfo`
`gdaIsCplx`
`gdaLoad`
`gdaPack`
`gdaRead`
`gdaReadByIndex`
`gdaReadSome`
`gdaReadSparse`
`gdaReadStruct`
`gdaReportVarInfo`

`gdaSave`
`gdaUpdate`
`gdaUpdateAndPack`
`gdaWrite`
`gdaWriteSome`
`gdtfastcat`
`ge`
`getarray`
`getdims`
`getf`
`getmatrix`
`getmatrix4D`
`getname`
`getnamef`
`getNextTradingDay`
`getNextWeekDay`
`getnr`
`getnrmt`
`getorders`
`getpath`
`getPreviousTradingDay`

Miscellaneous Topics

`getPreviousWeekDay`

`getRow`

`getscalar3D`

`getscalar4D`

`getTrRow`

`getwind`

`glm`

`gosub`

`goto`

`gradMT`

`gradMTm`

`gradMTTm`

`gradMTT`

`gradp`

`gradpcplx`

`graph`

`graphgpg`

`graphinit`

`graphprt`

`graphset`

`graphsev3`

gt

H

h5create

h5open

h5read

h5readAttribute

h5write

h5writeAttribute

hardcopy

hasimag

header

headermt

hess

hessMT

hessMTg

hessMTgw

hessMTm

hessMTmw

hessMTw

hessMTT

hessMTg

hessMTTgw

Miscellaneous Topics

`hessMTTm`

`hessMTTmw`

`hessMTTw`

`hesscplx`

`hessp`

`hist`

`histf`

`histp`

`hsec`

I

`if`

`imag`

`include`

`indcv`

`indexcat`

`indices`

`indices2`

`indicesf`

`indicesfn`

`indnv`

`indsav`

`int`

intgrat2
intgrat3
inthp
inthp1
inthp2
inthp3
inthp4
inthpControlCreate
intquad1
intquad2
intquad3
intrleav
intrleavsa
intrsect
intrsectsa
intsimp
inv
invpd
invswp
iscplx
iscplxsf

Miscellaneous Topics

`isinfnanmiss`

`issmiss`

`isSparse`

`itos`

K

`key`

`keyav`

`keymatchmc`

`keyw`

`keyword`

L

`lag`

`lag1`

`lagn`

`lapeighb`

`lapeighi`

`lapeighvb`

`lapeighvi`

`lapgeig`

`lapgeigh`

`lapgeighv`

`lapgeigv`

lapgschur
lapgsvdcst
lapgsvds
lapgsvdst
lapsvdcusv
lapsvds
lapsvdusv
ldl
ldlp
ldlsol
le
let
lib
library
license_id
line
linsolve
ln
lncdfbvn
lncdfbvn2
lncdfmvn

Miscellaneous Topics

`lncdfn`

`lncdfn2`

`lncdfnc`

`lnfact`

`lngamma`

`lngammacplx`

`lnpdfmvn`

`lnpdfmvt`

`lnpdfn`

`lnpdft`

`load`

`loadarray`

`loadadd`

`loadexe`

`loadf`

`loadk`

`loadm`

`loadp`

`loads`

`loadstruct`

`loadwind`

local
locate
loess
loessmt
loessmtControlCreate
log
loglog
logx
logy
loopnextindex
lower
lowmat
lowmat1
lpos
lprint
lpwidth
lshow
lt
ltrisol
lu
lusol

M

Miscellaneous Topics

`machEpsilon`

`makevars`

`makewind`

`margin`

`matalloc`

`matinit`

`matrix`

`mattoarray`

`maxbytes`

`maxc`

`maxindc`

`maxv`

`maxvec`

`mbesselei`

`mbesselei0`

`mbesselei1`

`mbesseli`

`mbesseli0`

`mbesseli1`

`meanc`

`median`

mergeby

mergebyssa

mergevar

minc

minindc

minv

miss

missex

missrv

moment

momentd

movingave

movingaveExpwgt

movingaveWgt

msym

N

nametype

ndpchk

ndpclex

ndpcntrl

ne

new

Miscellaneous Topics

`nextindex``nextn``nextnevn``nextwind``not``null``null1``numCombinations`**O**`oldfft``oldffti``ols``olsmt``olsmtControlCreate``olsqr``olsqr2``olsqrmt``ones``open``openpgg``optn``optnevn`

or

orth

output

outwidth

P

pacf

packedToSp

packr

parse

pause

pdfBinomial

pdfCauchy

pdfexp

pdfGenPareto

pdfHyperGeo

pdfLaplace

pdfLogistic

pdfn

pdfPoisson

pdfRayleigh

pdfWeibull

pdfWishartInv

Miscellaneous Topics

`pi`
`pinv`
`pinvmt`
`plot`
`plotAddBar`
`plotAddErrorBar`
`plotAddSurface`
`plotAddHist`
`plotAddHistF`
`plotAddHistP`
`plotAddScatter`
`plotAddTS`
`plotAddXY`
`plotAddArrow`
`plotAddArea`
`plotAddShape`
`plotAddTextbox`
`plotArea`
`plotBar`
`plotBox`
`plotCDFEmpirical`

`plotClearLayout`
`plotContour`
`plotCustomLayout`
`plotGetDefaults`
`plotHist`
`plotHistF`
`plotHistP`
`plotLayout`
`plotLogLog`
`plotLogX`
`plotLogY`
`plotOpenWindow`
`plotPolar`
`plotSave`
`plotScatter`
`plotSetBar`
`plotSetBkdColor`
`plotSetColorMap`
`plotSetContourLabels`
`plotSetFill`
`plotSetGrid`

Miscellaneous Topics

`plotSetLegend`
`plotSetLineColor`
`plotSetLineStyle`
`plotSetLineSymbol`
`plotSetLineThickness`
`plotSetNewWindow`
`plotSetTextInterpreter`
`plotSetTitle`
`plotSetWhichYAxis`
`plotSetXLabel`
`plotSetXRange`
`plotSetXTicCount`
`plotSetXTicInterval`
`plotSetXTicLabel`
`plotSetYLabel`
`plotSetYRange`
`plotSetYTicCount`
`plotSetZLabel`
`plotSetZLevels`
`plotSurface`
`plotsym`

plotTS
plotXY
polar
polychar
polyeval
polygamma
polyint
polymake
polymat
polymroot
polymult
polyroot
pop
powerM
pqqwin
prcsn
previousindex
princomp
print
printdos
printfm

Miscellaneous Topics

`printfmt`

`proc`

`prodc`

`psi`

`push`

`putarray`

`putf`

`putvals`

`pvCreate`

`pvGetIndex`

`pvGetParNames`

`pvGetParVector`

`pvLength`

`pvList`

`pvnumoffsets`

`pvoffsets`

`pvPack`

`pvPacki`

`pvPackm`

`pvPackmi`

`pvPacks`

`pvPacksi`

`pvPacksm`

`pvPacksmi`

`pvPutParVector`

`pvTest`

`pvUnpack`

Q

`QNewton`

`QNewtonmt`

`QNewtonmtControlCreate`

`QNewtonmtOutCreate`

`QNewtonSet`

`QProg`

`QProgmt`

`QProgmtInCreate`

`qqr`

`qqre`

`qqrep`

`qr`

`qre`

`qrep`

`qrsol`

Miscellaneous Topics

`qrtsol``qtyr``qtyre``qtyrep``quantile``quantiled``quantilem``quantilemd``qyr``qyre``qyrep`**R**`rank``rankindx``readr``real``reclassify``reclassifyCuts``recode``recserar``recsercp``recserrc`

`register_off`
`register_on`
`register_reset`
`register_show`
`renamefile`
`replay`
`rerun`
`rescale`
`reshape`
`retp`
`return`
`rev`
`rfft`
`rffti`
`rfftip`
`rfftn`
`rfftnp`
`rfftp`
`rndBernoulli`
`rndBeta`
`rndCauchy`

Miscellaneous Topics

`rndChiSquare`

`rndcon`

`rndbeta`

`rndCreateState`

`rndExp`

`rndgam`

`rndGamma`

`rndGeo`

`rndGumbel`

`rndHyperGeo`

`rndi`

`rndKMbeta`

`rndKMgam`

`rndKMi`

`rndKMn`

`rndKMnb`

`rndKMp`

`rndKMu`

`rndKMvm`

`rndLaplace`

`rndLCbeta`

`rndLCgam`
`rndLCi`
`rndLCn`
`rndLCnb`
`rndLCp`
`rndLCu`
`rndLCvm`
`rndLogNorm`
`rndmod`
`rndmult`
`rndMVn`
`rndMVt`
`rndn`
`rndnb`
`rndNegBinomial`
`rndns`
`rndp`
`rndpPoisson`
`rndseed`
`rndStateSkip`
`rndu`

Miscellaneous Topics

`rndus`

`rndvm`

`rndWeibull`

`rndWishart`

`rndWishartInv`

`rotater`

`round`

`rows`

`rowsf`

`rref`

`run`

S

`sampleData`

`satocv`

`satostrC`

`save`

`saveall`

`saved`

`savestruct`

`savewind`

`scale`

`scale3d`

scalerr
scalinfnanmiss
scalmiss
schtoc
schur
screen
scroll
searchsourcepath
seekr
selif
sega
seqm
setarray
setcnvrt
setdif
setdifsa
setvars
setvmode
setvwrmode
setwind
shell

Miscellaneous Topics

`shiftr`

`show`

`showpqq`

`sin`

`singleindex`

`sinh`

`sleep`

`solpd`

`sortc`

`sortcc`

`sortd`

`sorthc`

`sorthcc`

`sortind`

`sortindc`

`sortindmc`

`sortmc`

`sortr`

`sortrc`

`sparseCols`

`sparseEye`

`sparseFD`
`sparseFP`
`sparseHConcat`
`sparseNZE`
`sparseOnes`
`sparseRows`
`sparseScale`
`sparseSet`
`sparseSolve`
`sparseSubmat`
`sparseTD`
`sparseTranspose`
`sparseTrTD`
`sparseTscalar`
`sparseVConcat`
`spChol`
`spConjGradSol`
`spCreate`
`spDenseSubmat`
`spDiagRvMat`
`spEigv`

Miscellaneous Topics

`spEye`

`spGetNZE`

`spLDL`

`spline`

`spline1D`

`spline2D`

`spLU`

`spNumNZE`

`spOnes`

`SpreadsheetReadM`

`SpreadsheetReadSA`

`SpreadsheetWrite`

`spScale`

`spSubmat`

`spToDense`

`spTrTDense`

`spTScalar`

`spZeroes`

`sqpmt_feasible`

`sqpmt_meritFunct`

`sqpSolve`

`sqpSolveMT`
`sqpSolveMTcontrolCreate`
`sqpSolveMTlagrangeCreate`
`sqpSolveMToutCreate`
`sqpSolveset`
`sqrt`
`stdc`
`stdsc`
`stocv`
`stof`
`stop`
`strcombine`
`strindx`
`strjoin`
`string`
`strlen`
`strput`
`strrindx`
`strsect`
`strsplit`
`strsplitPad`

Miscellaneous Topics

`strtodt`

`strtodtd`

`strtof`

`strtofcplx`

`strtrim`

`strtriml`

`strtrimr`

`strtrunc`

`strtrunc1`

`strtruncpad`

`strtruncr`

`struct`

`submat`

`subscat`

`substute`

`subvec`

`sumc`

`sumr`

`surface`

`svd`

`svd1`

svd2

svdcusv

svds

svdusv

sylvester

sysstate

system

T

tab

tan

tanh

tempname

ThreadBegin

ThreadEnd

threadfor

threadendfor

ThreadJoin

ThreadStat

time

timedt

timestr

timeutc

Miscellaneous Topics

`title``tkf2eps``tkf2ps``tkf2ps_margin``tocart``todaydt``toeplitz``token``topolar``trace``trap``trapchk``trigamma``trim``trimr``trunc``type``typecv``typef`

U

`union``unionsa`

uniqindmc
uniqindx
uniqindxsa
unique
uniquemc
uniquesa
until
upmat
upmat1
upper
use
utctodt
utctodtv
utrisol

V

vals
varCovM
varCovMS
varCovX
varCovXS
varget
varget1

Miscellaneous Topics

varmall

varmares

varput

varputl

vartype

vartypef

vcm

vcms

vcx

vcxs

vec

vech

vecr

vfor

vget

view

viewxyz

vlist

vnamecv

volume

vput

vread

vtypecv

W

wait

waitc

walkindex

while

winclear

wincleararea

winclearttylog

winclose

wincloseall

winconvertpgg

window

wingetactive

wingetattributes

wingetcolorcells

wingetcursor

winmove

winopenpgg

winopentext

winopentty

Miscellaneous Topics

winpan

winprint

winprintpgg

winrefresh

winrefresharea

winresize

winsetactive

winsetbackground

winsetcolor

winsetcolorcells

winsetcolormap

winsetcursor

winsetforeground

winsetrefresh

winsettextwrap

winwrite

winzoompgg

writer

X

x_indcv

xlsGetSheetCount

xlsGetSheetSize

`xlsMakeRange`

`xlsReadM`

`xlsReadSA`

`xlsWrite`

`xlsWriteM`

`xlsWriteSA`

`xlabel`

`xor`

`xpnd`

`xtics`

`xy`

`xyz`

Y

`ylabel`

`ytics`

Z

`zeros`

`zeta`

`zlabel`

`ztics`

Index

#define 34-61, 34-62
#definecs 34-61
#else 34-61
#endif 34-61
#iflight 34-62
#ifndef 34-62
#include 34-62, 34-68, 35-878
#linesoff 34-62, 34-71, 35-1009
#lineson 34-62, 34-71, 35-1009
#srcfile 34-62
#srcline 34-62
#undef 34-62
__altnam 35-572
__output 35-572, 35-1043, 35-1720
__title 35-572
__Tol 35-572
__vpad 35-572
_eqs_IterInfo 35-572
_eqs_JacobianProc 35-571
_eqs_MaxIters 35-571
_eqs_StepTol 35-572
_eqs_TypicalF 35-572
_eqs_TypicalX 35-572
_loess_Degree 35-1043
_loess_NumEval 35-1043
_loess_Span 35-1043
_loess_WgtType 35-1043
_pageshf 34-88
_pagesiz 34-88
_parrow 34-88
_parrow3 34-88
_paxes 34-86
_paxht 34-87
_pbox 34-90
_pboxctl 34-87
_pboxlim 34-87
_pcolor 34-87
_pcrop 34-90
_pcross 34-86
_pdate 34-87
_perrbar 34-88
_pframe 34-90
_pgrid 34-86
_plctrl 34-87

<code>_plegctl</code> 34-87	<code>_ptitlht</code> 34-87
<code>_plegstr</code> 34-87	<code>_pxpmax</code> 34-86
<code>_pline</code> 34-88	<code>_pxsci</code> 34-86
<code>_pline3d</code> 34-88	<code>_pypmax</code> 34-86
<code>_plotshf</code> 34-88, 34-89	<code>_pysci</code> 34-86
<code>_plotsiz</code> 34-88, 34-89	<code>_pzclr</code> 34-88
<code>_pltype</code> 34-88	<code>_pzoom</code> 34-89
<code>_plwidth</code> 34-88	<code>_pzpmax</code> 34-86
<code>_pmcolor</code> 34-90	<code>_pzsci</code> 34-86
<code>_pmsgctl</code> 34-87	<code>_sqp_A</code> 35-1716
<code>_pmsgstr</code> 34-87	<code>_sqp_B</code> 35-1716
<code>_pnum</code> 34-87	<code>_sqp_Bounds</code> 35-1718
<code>_pnumht</code> 34-87	<code>_sqp_C</code> 35-1717
<code>_protate</code> 34-88	<code>_sqp_D</code> 35-1717
<code>_pscreen</code> 34-89	<code>_sqp_DirTol</code> 35-1719
<code>_psilent</code> 34-89	<code>_sqp_EqProc</code> 35-1717
<code>_pstype</code> 34-88	<code>_sqp_FeasibleTest</code> 35-1720
<code>_psym</code> 34-88	<code>_sqp_GradProc</code> 35-1718
<code>_psym3d</code> 34-88	<code>_sqp_HessProc</code> 35-1719
<code>_psymsiz</code> 34-88	<code>_sqp_IneqProc</code> 35-1717
<code>_ptek</code> 34-89	<code>_sqp_MaxIters</code> 35-1719
<code>_pticout</code> 34-86	<code>_sqp_ParNames</code> 35-1719

- `_sqp_PrintIters` 35-1720
- `_sqp_RandRadius` 35-1720
- `abs` 34-4, 35-52
- absolute value 35-52
- `acf` 34-19, 35-53
- `concat` 34-40, 34-41, 35-58
- additive sequence 35-1639
- `aeye` 34-40, 35-61
- algebra, linear 34-8
- `amax` 34-40, 35-63
- `amean` 34-42, 35-65
- `AmericanBinomCall` 34-30, 35-67
- `AmericanBinomCall_Greeks` 34-30, 35-69
- `AmericanBinomCall_ImpVol` 34-30, 35-71
- `AmericanBinomPut` 34-30, 35-73
- `AmericanBinomPut_Greeks` 34-30, 35-75
- `AmericanBinomPut_ImpVol` 34-31, 35-77
- `AmericanBSCall` 34-31, 35-79
- `AmericanBSCall_Greeks` 34-31, 35-80
- `AmericanBSCall_ImpVol` 34-31, 35-82
- `AmericanBSPut` 34-31, 35-84
- `AmericanBSPut_Greeks` 34-31, 35-85
- `AmericanBSPut_ImpVol` 34-31, 35-87
- `amin` 34-40, 35-89
- `amult` 34-42, 35-92
- `annotationGetDefaults` 34-84, 35-94
- `annotationSetBkd` 34-84, 35-95
- `annotationSetFont` 34-84, 35-97
- `annotationSetLineColor` 34-84, 35-99
- `annotationSetLineStyle` 34-84, 35-101
- `annotationSetLineThickness` 34-84, 35-103
- `annualTradingDays` 34-31, 35-104
- `arcs` 34-4, 35-106
- `arcsin` 34-4, 35-107
- `arctangent` 35-126
- `area plot` 34-80, 35-1200
- `reshape` 34-40, 34-41, 35-108, 35-124
- `arrayalloc` 34-40, 35-110
- `arrayindex` 34-42, 35-112
- `arrayinit` 34-40, 35-114

- arrays 34-40
- arraytomat 34-41, 35-115
- ASCII files 34-56, 34-79, 35-116, 35-751
- ASCII files, reading 35-1030
- asciiload 35-116
- asclabel 34-87, 35-118
- astd 34-19, 35-120
- astds 34-19, 35-122
- asum 34-40, 35-124
- atan 34-4, 35-126
- atan2 34-4, 35-128
- ATOG 34-56, 35-622, 35-623
- atranspose 34-42, 35-130
- autocorrelations 35-53
- auxiliary output 35-1156
- auxiliary output, width 35-1160
- axes 34-86
- axes numbering 34-86
- axes, reversed 35-1960, 35-1965
- axmargin 34-88, 34-89, 35-132
- balance 34-8, 35-135
- band 34-8
- bandchol 34-8, 35-138
- bandcholsol 34-8, 35-140
- bandltsol 34-8, 35-141
- bandrv 34-8, 35-143
- bandsolpd 34-9, 35-145
- bar 34-81, 34-85, 34-85, 35-146
- bar plot 34-80, 35-1234
- base10 34-30, 35-148
- begwind 34-88, 35-149
- Bessel 34-4, 34-4, 35-150, 35-151, 35-1077
- besselj 34-4, 35-150
- bessely 34-4, 35-151
- beta 34-4, 35-153
- beta function 34-4, 35-153
- beta function, incomplete 35-160
- binary file, loading 35-751
- bivariate Normal 34-28, 35-167, 35-169, 35-171
- box 34-85, 34-85, 35-154
- box plot 34-80, 34-81, 35-1236
- boxcox 34-5, 35-156

- branching 34-64
- break 34-65, 35-157
- call 34-64, 34-67, 35-159
- Cartesian coordinates 34-86, 35-1331, 35-1848, 35-1853, 35-1961, 35-1962
- Cauchy 35-173, 35-174, 35-1181, 35-1519
- cdfBeta 34-25, 35-160
- cdfBetaInv 34-25, 35-162
- cdfBinomial 34-25, 35-164
- cdfBinomialInv 34-25, 35-165
- cdfBvn 34-25, 35-167
- cdfBvn2 34-25, 35-169
- cdfBvn2e 34-25, 35-171
- cdfCauchy 34-25, 35-173
- cdfCauchyInv 34-25, 35-174
- cdfChic 34-25, 35-175
- cdfChii 34-25, 35-177
- cdfChinc 34-25, 35-179
- cdfChincInv 35-180
- cdfEmpirical 34-25, 35-182
- cdfExp 34-26, 35-185
- cdfExpInv 34-26, 35-186
- cdfFc 34-26, 35-186
- cdfFnc 34-26, 35-189
- cdfFncInv 34-26, 35-190
- cdfGam 34-26, 35-192
- cdfGenPareto 34-26, 35-194
- cdfHyperGeo 34-26, 35-196
- cdfLaplace 34-26, 35-198
- cdfLaplaceInv 34-26, 35-199
- cdfLogistic 35-199
- cdfLogisticInv 35-201
- cdfm.src 35-209, 35-215, 35-220, 35-226, 35-232, 35-239
- cdfMvn 34-26, 35-201
- cdfMvn2e 34-26, 35-215
- cdfMvnce 34-26, 35-205
- cdfMvne 34-26, 35-210
- cdfMvt2e 34-27, 35-233
- cdfMvtce 34-26, 35-221
- cdfMvte 34-27, 35-227
- cdfN 34-27, 35-239
- cdfN2 34-27, 35-246

- cdfNc 34-27, 35-239
- cdfNegBinomial 34-27, 35-243
- cdfNegBinomialInv 34-27, 35-245
- cdfNi 34-27, 35-248
- cdfPoisson 34-27, 35-250
- cdfPoissonInv 34-27, 35-251
- cdfRayleigh 34-27, 35-252
- cdfRayleighInv 34-27, 35-254
- cdfTc 34-27, 35-255
- cdfTci 34-27, 35-257
- cdfTnc 34-28, 35-258
- cdfTvn 34-28, 35-261
- cdfWeibull 34-28, 35-263
- cdfWeibullInv 34-28, 35-265
- cdir 34-69, 35-266
- ceil 34-30, 34-30, 35-267
- change log 36-1
- change working directory 35-269, 35-269
- changeDir 34-69, 35-269
- characteristic polynomial 35-1333
- chdir 34-69, 35-269
- chi-bar square 35-270
- chi-square 35-175, 35-177
- chi-square, noncentral 35-179, 35-180
- chiBarSquare 34-19, 35-270
- chol 34-9, 34-13, 35-272
- choldn 34-9, 35-274
- Cholesky decomposition 34-8, 34-8, 34-9, 34-13, 35-138, 35-140, 35-272, 35-1677
- cholsol 34-9, 34-13, 35-276
- cholup 34-9, 35-278
- chrs 34-71, 34-79, 35-279
- clear 34-70, 34-70, 35-281
- clear global symbols 35-282
- clear program input/output window 35-289
- clearg 34-70, 35-282
- close 34-55, 34-56, 35-283
- closeall 34-55, 34-56, 35-286
- cls 34-78, 35-289
- code 34-57, 34-59, 35-290
- code (dataloop) 34-57, 35-293
- coefficient of determination 35-1124,

- 35-1135
- coefficients 35-1042, 35-1044, 35-1142
- coefficients, standardized 35-1124, 35-1134
- cols 34-34, 34-35, 35-295
- colsf 34-34, 34-34, 34-35, 35-296
- columns in a matrix 34-34
- columns in a matrix 35-295, 35-296
- combine 34-19, 35-297
- combined 34-19, 35-299
- comlog 34-78, 35-302
- common errors 36-8
- comparison functions 35-509, 35-511, 35-631, 35-633
- compile 34-67, 34-68, 35-303
- compiler control 34-61
- compiling 34-67
- compiling files 35-303
- complex 34-35, 35-305
- complex constants 35-460, 35-998, 35-1741
- complex modulus 35-52
- con 34-77, 34-78, 35-306
- concatenation, matrix 35-58, 35-1084
- cond 34-9, 34-13, 35-309
- condition number 35-309
- conj 35-310
- cons 34-77, 34-78, 35-311
- conScore 34-19
- ConScore 35-312
- console 34-77
- constants, complex 35-460, 35-998, 35-1741
- continue 34-65, 35-316
- contour 34-85, 34-85, 34-88, 35-318
- contour plot 34-80
- conv 34-14, 34-19, 35-319
- conversion, array to type matrix 35-115
- conversion, character to ASCII value 35-1892
- conversion, character vector to string 35-365, 35-366
- conversion, float to ASCII 35-685, 35-686
- conversion, matrix to string array 35-690
- conversion, string array to string 35-320

- conversion, string to floating point 34-72
- conversion, string to string array 35-321
- convertsatostr 34-71, 35-320
- convertstrtosa 34-71, 35-321
- convolution 35-319
- correlation matrix 34-19, 34-19, 34-20, 34-20, 34-20, 35-322, 35-322, 35-324
- corrmm 34-19
- corrms 34-19, 35-324
- corrvc 34-20, 35-322
- corrxx 34-20, 35-322
- corrxs 34-20, 35-324
- cos 34-5, 35-327
- cosh 34-5, 35-328
- cosine 34-5, 34-5, 35-327
- cosine, inverse 34-4, 35-106
- counts 34-34, 35-329
- countwts 34-34, 35-331
- create 34-55, 35-333
- cross-product 34-20, 35-340, 35-1095
- crossprd 34-20, 35-340
- crout 34-9, 34-13, 35-341
- Crout decomposition 34-9, 34-9, 34-13, 35-341, 35-343
- croutp 34-9, 34-13, 35-343
- csrcol 34-78, 35-345
- csrln 34-78, 35-345
- CSV files 34-33
- csvReadM 34-33, 34-45, 35-346
- csvReadSA 34-45, 35-352
- cumprodc 34-34, 35-360
- cumsumc 34-34, 35-362
- cumulative distribution function 35-182, 35-185, 35-186, 35-186, 35-189, 35-190, 35-194, 35-198, 35-199, 35-199, 35-201, 35-201, 35-205, 35-210, 35-215, 35-227, 35-233, 35-239, 35-243, 35-245, 35-246, 35-250, 35-251, 35-252, 35-254, 35-255, 35-257, 35-261, 35-263, 35-265, 35-1238
- cumulative products 35-360
- cumulative sums 35-362
- cursor 34-78, 34-79, 35-345, 35-1041
- curve 34-5, 35-363
- cvtos 34-71, 35-365
- cvtosa 34-71, 35-366

- data coding 34-57
- data file reading 34-44
- data file writing 34-44
- data loop 35-293, 35-466, 35-515, 35-625, 35-950, 35-1061, 35-1159, 35-1486, 35-1636, 35-1914
- data sets 34-55, 34-56, 34-56, 34-61, 35-333, 35-368, 35-370, 35-373, 35-377
- database 34-46
- database connect 34-49
- database errors 34-48
- database information 34-48
- database properties 34-47
- database setup 34-46
- database transaction 34-49
- datacreate 34-55, 35-368
- datacreatecomplex 34-55, 35-370
- datalist 34-55, 35-373
- dataload 34-33, 35-374
- dataloop (dataloop) 34-57, 35-376
- dataopen 34-55, 35-377
- datasave 34-33, 34-33, 35-381
- date 34-75, 35-382, 35-383, 35-384, 35-385
- datestr 34-75, 35-383
- datestring 34-75, 35-384
- datestrymd 34-75, 35-385
- dayinyr 34-75, 35-386
- dayofweek 34-75, 35-387
- dbAddDatabase 34-46
- dbClose 34-49
- dbCommit 34-49
- dbCreateQuery 34-49
- dbExecQuery 34-49
- dbGetConnectOptions 34-47
- dbGetDatabaseName 34-47
- dbGetDriverName 34-47
- dbGetDrivers 34-47
- dbGetHostName 34-47
- dbGetLastErrorNum 34-48
- dbGetLastErrorText 34-49
- dbGetNumericalPrecPolicy 34-47
- dbGetPassword 34-47
- dbGetPort 34-47

dbGetPrimaryIndex 34-48	dbQueryGetLastErrorText 34-49
dbGetTableHeaders 34-48	dbQueryGetLastInsertID 34-50
dbGetTables 34-48	dbQueryGetLastQuery 34-51
dbHasFeature 34-48	dbQueryGetNumRowsAffected 34-51
dbIsDriverAvailable 34-47	dbQueryGetPosition 34-52
dbIsOpen 34-47	dbQueryIsActive 34-51
dbIsOpenError 34-49	dbQueryIsForwardOnly 34-51
dbIsValid 34-47	dbQueryIsNull 34-51
dbOpen 34-49	dbQueryIsSelect 34-51
dbQueryBindValue 34-50	dbQueryIsValid 34-51
dbQueryClear 34-50	dbQueryPrepare 34-50
dbQueryCols 34-50	dbQueryRows 34-51
dbQueryExecPrepared 34-50	dbQuerySeek 34-52
dbQueryFetchAllM 34-52	dbQuerySeekFirst 34-52
dbQueryFetchAllSA 34-52	dbQuerySeekLast 34-52
dbQueryFetchOneM 34-52	dbQuerySeekNext 34-52
dbQueryFetchOneSA 34-52	dbQuerySeekPrevious 34-52
dbQueryFinish 34-50	dbQuerySetForwardOnly 34-51
dbQueryGetBoundValue 34-50	dbRemoveDatabase 34-47
dbQueryGetBoundValues 34-50	dbRollback 34-49
dbQueryGetField 34-53	dbSetConnectOptions 34-48
dbQueryGetLastErrorNum 34-49	dbSetDatabaseName 34-48

- dbSetHostName 34-48
- dbSetNumericalPrecPolicy 34-48
- dbSetPassword 34-48
- dbSetPort 34-48
- dbTransaction 34-50
- debug 34-70, 35-458
- debugger 34-70, 35-458
- debugging 34-70
- declare 34-67, 35-458
- delete 34-70, 35-464
- delete (dataloop) 34-57, 35-466
- deleteFile 34-69, 35-467
- deletion 35-468, 35-473
- delif 34-35, 34-38, 34-38, 35-468
- delrows 34-35, 35-473
- denseToSp 34-38, 35-475
- denseToSpRE 34-38, 35-476
- denToZero 35-478
- derivatives 35-820
- derivatives, second partial 35-869
- descriptive statistics 35-516, 35-523
- design 34-20, 35-480
- design matrix 34-20, 35-480
- det 34-9, 34-13, 35-482
- determinant 35-482, 35-484
- detl 34-9, 34-13, 35-484
- dfft 34-14, 35-485
- dffti 34-14, 35-486
- diag 34-35, 35-487
- diagonal 35-487, 35-490
- diagrv 34-35, 35-490
- differentiation 34-6
- digamma 34-5, 35-492
- directory 35-266
- dlibrary 34-69, 35-492
- dllcall 34-69, 35-495
- do loop 34-65, 34-65, 35-157, 35-316, 35-497
- do until 34-65
- do while 34-65, 35-497, 35-497
- dos 34-69, 35-500
- doswin 34-79, 35-503
- DOSWinCloseall 34-79, 35-503
- DOSWinOpen 34-79, 35-504

- dot 34-17, 34-20, 35-507
- dotfeq 34-17, 35-509
- dotfeqmt 34-17, 35-511
- dotfge 34-17, 35-509
- dotfgemt 34-18, 35-511
- dotfgt 34-18, 35-509
- dotfgtmt 34-18, 35-511
- dotfle 34-18, 35-509
- dotflemt 34-18, 35-511
- dotflt 34-18, 35-509
- dotfltmt 34-18, 35-511
- dotfne 34-18, 35-509
- dotfnemt 34-18, 35-511
- draw 34-85, 35-513
- drop (dataloop) 34-57, 35-515
- DS structure 35-515
- dsCreate 34-42, 35-515
- dstat 34-67, 35-516
- dstatmt 34-20, 35-523
- dstatmtControlCreate 34-20, 35-531
- dtdate 34-75, 35-532
- dtday 34-75, 35-534
- dttime 34-75, 35-535
- dttodtv 34-75, 35-536
- dttostr 34-75, 35-537
- dttoutc 34-75, 35-540
- dtv vector 35-542, 35-545
- dtvnormal 34-75, 35-542
- dtvtodt 34-75, 35-543
- dtvtoutc 34-76, 35-545
- dummy 34-57, 34-59, 35-546
- dummy variables 35-546, 35-548, 35-550
- dummybr 34-57, 34-59, 35-548
- dummydn 34-57, 34-59, 35-550
- Durbin-Watson statistic 35-1123, 35-1135
- dynamic libraries 34-69, 34-69
- ed 34-78, 35-553
- edit 34-78, 35-554
- editor 35-554
- editor, alternate 35-553
- eig 34-13, 34-13, 35-556
- eigenvalues 34-13, 34-13, 34-13, 35-556, 35-558, 35-960, 35-962, 35-

- 968, 35-969
- eigenvalues and eigenvectors 34-9, 34-10, 34-10, 34-13, 34-13, 34-39, 35-560, 35-561, 35-964, 35-966, 35-971, 35-973, 35-1687
- eigh 34-13, 34-13, 35-558
- eighv 34-13, 34-13, 35-560
- eigv 34-13, 34-13, 35-561
- elapsedTradingDays 34-31, 35-563
- else 35-876
- elseif 35-876
- empty matrix 35-296, 35-999, 35-1030, 35-1601, 35-1624
- end 34-64, 35-565
- end of file 34-55, 34-56, 35-569
- endfor 34-65
- endif 34-64
- endo 34-65, 34-65, 34-65
- endp 34-66, 34-67, 35-566
- endwind 34-88, 35-567
- envget 34-69, 35-568
- environment, search 35-568
- eof 34-55, 34-56, 35-569
- eqSolve 34-23, 35-571
- eqSolveMt 34-23, 35-576
- EqSolveMtControlCreate 34-23, 35-584
- eqSolveMtOutCreate 34-23, 35-585
- eqSolveSet 34-23, 35-587
- erf 34-28, 35-587
- erfc 34-28, 35-587
- erfcplx 34-28, 35-589
- erfCInv 35-555
- erfcplx 34-28, 35-589
- erfInv 35-555
- error 34-71, 34-71, 35-590
- error bar 34-81
- error code 35-590
- error function 35-555, 35-587, 35-589
- error handling 34-70
- error messages 36-13
- error trapping 35-1855
- errorlog 34-71, 35-593
- errorlogat 35-593
- etdays 34-76, 35-594
- ethsec 34-76, 35-595

- etstr 34-76, 35-597
- EuropeanBinomCall 34-31, 35-598
- EuropeanBinomCall_Greeks 34-31, 35-600
- EuropeanBinomCall_ImpVol 34-31, 35-602
- EuropeanBinomPut 34-31, 35-604
- EuropeanBinomPut_Greeks 34-31, 35-605
- EuropeanBinomPut_ImpVol 34-31, 35-608
- EuropeanBSCall 34-31, 35-609
- EuropeanBSCall_Greeks 34-32, 35-611
- EuropeanBSCall_ImpVol 34-32, 35-613
- EuropeanBSPut 34-32, 35-614
- EuropeanBSPut_Greeks 34-32, 35-616
- EuropeanBSPut_ImpVol 34-32, 35-618
- Excel files 34-34
- exctsmpl 34-35, 35-619
- exec 34-69, 35-621
- execbg 34-69, 35-622
- execution control 34-64
- exp 34-5, 35-623
- exponential 35-185, 35-186, 35-1182
- exponential function 35-623
- extern (dataloop) 34-57, 35-625
- external 34-67, 35-626
- eye 34-32, 35-627
- F distribution 35-186
- fcheckerr 34-45, 35-629
- fclearerr 34-45, 35-630
- feq 34-18, 35-631
- feqmt 34-18, 35-633
- fflush 34-45, 35-635
- fft 34-14, 35-635
- ffti 34-14, 35-636
- fftm 34-14, 35-637
- fftimi 34-14, 35-640
- fftn 34-14, 35-643
- fge 34-18, 35-631
- fgemt 34-18, 35-633
- fgets 34-45, 35-645
- fgetsa 34-45, 35-646
- fgetsat 34-45, 35-647
- fgetst 34-46, 35-648

- fgt 34-18, 35-631
- fgtmt 34-18, 35-633
- file handle 35-283, 35-337, 35-1149
- file management 34-69
- fileinfo 34-69, 35-649
- filesa 34-69, 35-651
- finance functions 34-30
- fle 34-18, 35-631
- flemt 34-18, 35-633
- floor 34-30, 35-652
- flt 34-18, 35-631
- fltmt 34-19, 35-633
- fmod 34-5, 35-654
- fn 34-66, 35-656
- fne 34-19, 35-631
- fnemt 34-19, 35-633
- fonts 34-87, 35-657
- fopen 34-46, 35-658
- for 34-65, 35-660
- for loop 34-65, 34-65, 35-157, 35-316, 35-660
- format 34-78, 35-663
- formatcv 34-78, 34-79, 35-673
- formatnv 34-78, 34-79, 35-674
- Fourier transform 35-635, 35-636, 35-1506, 35-1507
- Fourier transform, discrete 35-485, 35-486
- Fourier transforms 34-14
- fputs 34-46, 35-675
- fputst 34-46, 35-678
- fseek 34-46, 35-681
- fstrerror 34-46, 35-683
- ftell 34-46, 35-684
- ftocv 34-71, 34-75, 35-685
- ftos 34-72, 34-74, 35-686
- ftostrC 34-72, 35-690
- functions 34-66
- fuzzy comparison functions 35-509, 35-511, 35-631, 35-633
- fuzzy conditional functions 34-17
- gamma 34-5, 35-693
- gamma function 35-693, 35-694, 35-695, 35-1023
- gamma, incomplete 35-192

- gamma, log 35-1021
- gammacplx 34-5, 35-694
- gammair 34-5, 35-695
- Gauss-Legendre quadrature 35-919, 35-922, 35-926
- gauss colors 36-39
- GAUSS Data Archives 34-53, 34-54, 34-54, 34-54, 35-697, 35-699, 35-700, 35-704, 35-709, 35-710, 35-711, 35-712, 35-714, 35-715, 35-717, 35-719, 35-720, 35-724, 35-726, 35-727, 35-728, 35-731, 35-732, 35-734, 35-735, 35-738, 35-740, 35-742, 35-743, 35-744, 35-746
- GAUSS Graphics Colors 36-39
- gausset 33-7, 34-68, 35-696
- gdaAppend 34-53, 35-697
- gdaCreate 34-53, 35-699
- gdaDStat 34-20, 34-53, 35-700
- gdaDStatMat 34-20, 34-53, 35-704
- gdaGetIndex 34-53, 35-709
- gdaGetName 34-53, 35-710
- gdaGetNames 34-53, 34-53, 35-711
- gdaGetOrders 35-712
- gdaGetType 34-53, 35-714
- gdaGetTypes 34-53, 35-715
- gdaGetVarInfo 34-53, 35-717
- gdaIsCplx 34-53, 35-719
- gdaLoad 34-53, 35-720
- gdaPack 34-54, 35-724
- gdaRead 34-54, 35-726
- gdaReadByIndex 34-54, 35-727
- gdaReadSome 34-54, 35-728
- gdaReadSparse 34-54, 35-731
- gdaReadStruct 34-54, 35-732
- gdaReportVarInfo 34-54, 35-734
- gdaSave 34-54, 35-735
- gdaUpdate 34-54, 35-738
- gdaUpdateAndPack 34-54, 35-740
- gdaVars 35-742
- gdaWrite 34-54, 35-743
- gdaWrite32 34-54, 35-744
- gdaWriteSome 34-54, 35-746
- generalized inverse 35-1197, 35-1199
- Generalized Pareto 34-26, 34-29, 35-194, 35-1183

- getarray 34-41, 35-749
- getdims 34-41, 35-750
- getf 34-72, 35-751
- getGAUSSHome 34-69, 35-754
- getmatrix 34-41, 35-755
- getmatrix4D 34-41, 35-757
- getname 34-56, 35-759
- getnamef 34-56, 34-57, 35-760
- getNextTradingDay 34-32, 35-762
- getNextWeekDay 34-32, 35-763
- getnr 34-55, 35-764
- getnrmt 34-55, 35-765
- getorders 34-41, 35-766
- getpath 34-69, 35-767
- getPreviousTradingDay 34-32, 35-769
- getPreviousWeekDay 34-32, 35-770
- getRow 35-770
- getscalar3D 34-41, 35-772
- getscalar4D 34-41, 35-773
- getTrRow 35-775
- getwind 34-89, 35-776
- glm 34-20, 35-777
- global control variables 35-696
- global variable 35-458
- Goertzel algorithm 35-486, 35-487
- gosub 34-66, 34-66, 34-66, 35-807
- goto 34-64, 34-64, 35-810
- gradcplx 34-6, 34-6, 35-818
- gradient 35-818
- gradMT 34-6, 35-811
- gradMTm 34-6, 35-813
- gradMTT 34-6, 35-815
- gradMTTm 34-6, 35-817
- gradp 34-6, 34-8, 35-818
- graphic panels 35-1064, 35-1066, 35-1112
- Graphics 34-80, 34-80, 34-85
- graphprt 34-89, 35-820
- graphs, saving 35-1257
- graphset 34-90, 35-824
- h5create 34-46, 35-825
- h5open 34-46, 35-828
- h5read 34-46, 35-831
- h5readAttribute 34-46, 35-838

- h5write 34-46, 35-840
- h5writeAttribute 34-46, 35-842
- hasimag 34-70, 35-845
- HDF 5 files 34-46
- header 34-78, 35-846
- headermt 34-78, 35-847
- help facility 35-1007
- Hermitian matrix 34-9, 34-9, 34-9, 34-9,
34-10, 34-10, 34-13, 34-13, 35-
560, 35-960, 35-962, 35-964, 35-
966, 35-969, 35-971
- hess 34-9, 35-849
- hesscplx 34-7, 34-7, 35-868
- Hessian 34-6, 34-6, 34-6, 34-6, 34-6,
34-6, 34-7, 34-7, 34-7, 34-7, 35-
850, 35-852, 35-854, 35-855, 35-
857, 35-859, 35-861, 35-863, 35-
864, 35-866, 35-868
- hessMT 34-6, 35-850
- hessMTg 34-6, 35-852
- hessMTgw 34-6, 35-854
- hessMTm 34-6, 35-855
- hessMTmw 34-6, 35-857
- hessMTT 34-6, 35-859
- hessMTTg 34-7, 35-861
- hessMTTgw 34-7, 35-863
- hessMTTm 34-7, 35-864
- hessMTw 34-7, 35-866
- hessp 34-7, 34-8, 35-868
- hist 34-85, 35-870
- histf 34-85, 35-872
- histogram 34-80, 34-80, 34-80, 34-81,
34-81, 34-81, 34-85, 34-85, 34-85,
35-870, 35-872, 35-873, 35-1216,
35-1217, 35-1218, 35-1249, 35-
1250, 35-1250
- histp 34-85, 35-873
- hsec 34-76, 34-76, 35-874
- hyperbolic cosine 35-328
- hyperbolic sine 35-1659
- hyperbolic tangent 35-1832
- identity matrix 35-627, 35-1690
- if 34-64, 35-876
- imag 34-35, 35-877
- imaginary matrix 35-877
- incomplete beta function 35-160
- incomplete gamma function 35-192
- indcv 34-36, 35-880

index variables	35-1147	intquad2	34-8, 34-8, 35-922
indexcat	34-34, 35-882	intquad3	34-8, 34-8, 35-926
indices	34-56, 34-57, 35-883	intrleav	34-60, 34-61, 35-930
indices2	34-56, 34-57, 35-885	intrleavsa	34-60, 35-932
indicesf	34-56, 35-886	intrsect	34-36, 34-72, 35-933
indicesfn	34-56, 35-888	intrsectsa	35-936
indnv	34-36, 35-889	intsimp	34-8, 34-8, 35-938
indsav	34-72, 35-891	inv	34-9, 34-13, 35-939
infinity	35-945, 35-1626	inverse cosine	35-106
input, console	35-306	inverse sine	35-107
input, keyboard	35-306	inverse, generalized	35-1197, 35-1199
integrate1d	34-7, 35-892	inverse, matrix	35-939
integration	34-6	inverse, sweep	35-941
intersection	35-933	invpd	34-9, 34-13, 35-939
intgrat2	34-7, 34-8, 35-897	invswp	34-9, 35-941
intgrat3	34-7, 34-8, 35-899	iscplx	34-55, 34-70, 35-942
inthp1	34-7, 35-902	iscplxsf	35-943
inthp2	34-7, 35-906	isden	35-944
inthp3	34-7, 35-910	isinfnanmiss	34-58, 35-945
inthp4	34-7, 35-914	ismiss	35-946
inthpControlCreate	34-7, 35-918	itos	35-948
intquad1	34-7, 34-8, 34-8, 35-919	Jacobian	35-818

- keep (dataloop) 34-58, 35-950
- key 34-77, 34-77, 34-77, 35-951
- keyav 34-77, 35-952
- keyboard input 35-306, 35-311
- keyboard, reading 35-951
- keyw 34-77, 34-78, 35-953
- keyword 34-66, 35-953
- keyword procedure 35-953
- keywords 34-66
- lag (dataloop) 34-58
- lag1 34-58, 35-957
- lagn 34-58, 35-958
- lambda 35-179
- lapeighb 34-9, 35-960
- lapeighi 34-9, 35-962
- lapeighvi 34-9, 35-966
- lapeigvb 34-9, 35-964
- lapgeig 34-10, 35-968
- lapgeigh 34-10, 35-969
- lapgeighv 34-10, 35-971
- lapgeigv 34-10, 35-973
- lapgschur 34-10
- lapgsvdest 34-10, 35-974
- lapgsvds 34-10, 35-977
- lapgsvdst 34-10, 35-979
- Laplace 34-26, 34-26, 34-29, 35-198, 35-199, 35-1187
- lapsvdcusv 34-10, 35-987
- lapsvds 34-10, 35-989
- lapsvdusv 34-10, 35-990
- ldl 34-10, 35-992
- LDL decomposition 34-10, 34-10, 34-11
- ldlp 34-10, 35-994
- ldlsol 34-11, 34-23, 35-996
- least squares regression 35-1120, 35-1129
- legend 34-82, 35-1276
- let 35-997
- lib 35-1001
- libraries 34-67
- libraries, active 34-67, 35-1005
- libraries, shared 35-492
- library 34-67, 35-1001, 35-1005
- line numbers 35-1009
- line plot 34-80, 34-80, 34-80, 34-81

- line thickness 35-1285
- line type 35-1281
- linear algebra 34-8
- linear equation 35-1661, 35-1678
- linsolve 34-23, 35-1010
- listwise (dataloop) 34-58, 35-1012
- ln 34-5, 35-1012
- lncdfbvn 34-28, 35-1013
- lncdfbvn2 34-28, 35-1014
- lncdfmvn 34-28, 35-1016
- lncdfn 34-28, 35-1017
- lncdfn.src 35-1014, 35-1017, 35-1018,
35-1020, 35-1021
- lncdfn2 34-28, 35-1018
- lncdfnc 34-28, 35-1020
- lnfact 34-5, 35-1021
- lngammacplx 34-5, 35-1023
- lnpdfmvn 34-28, 35-1024
- lnpdfmvt 34-28, 35-1025
- lnpdfn 34-29, 35-1026
- lnpdft 34-29, 35-1027
- load 34-33, 34-33, 35-1028
- loadarray 35-1033
- loadadd 34-33, 34-33, 34-55, 35-1035
- loadf 34-33, 34-33, 35-1028
- loadk 34-33, 34-33, 35-1028
- loadm 34-33, 34-33, 35-1028
- loadp 34-68, 34-68, 35-1028
- loads 34-72, 35-1028
- loadstruct 34-42, 35-1039
- loadwind 34-89, 35-1040
- local 34-66, 34-67, 35-1041
- local variable declaration 35-1041
- locate 34-78, 35-1041
- loess 34-20, 35-1042
- loessmt 34-20, 35-1044
- loessmtControlCreate 34-20, 35-1046
- log 34-5, 35-1047
- log coordinates 35-1048, 35-1253
- log factorial 35-1021
- log gamma 35-1021
- log, base 10 35-1047
- log, natural 35-1012
- logging commands 35-302

- logistic 35-199, 35-201
- loglog 34-85, 35-1048
- logx 34-85, 35-1049
- logy 34-85, 35-1050
- looping 34-65
- loopnextindex 34-42, 35-1051
- lower 34-72, 35-1054
- lower triangular matrix 35-1057, 35-1060
- lowmat 34-36, 34-38, 35-1056
- lowmat1 34-36, 34-38, 35-1056
- ltrisol 34-23, 35-1057
- lu 34-11, 35-1058
- LU decomposition 34-11, 34-11, 34-39, 35-1058
- lusol 34-11, 34-23, 35-1060
- machEpsilon 34-30, 35-1061
- machine epsilon 35-1061
- make (dataloop) 34-58, 35-1061
- makevars 34-56, 34-57, 35-1062
- makewind 34-89, 35-1064
- margin 34-89, 35-1066
- matalloc 34-32, 35-1067
- matinit 34-32, 34-33, 35-1068
- matrix manipulation 34-32
- matrix, creation 35-997
- matrix, empty 35-296, 35-999, 35-1030, 35-1601, 35-1624
- matrix, ones 35-1145
- matrix, zeros 35-1967
- mattoarray 34-40, 35-1069
- maxbytes 34-70, 35-1076
- maxc 34-34, 35-1070
- maximum element 35-1070
- maximum element index 35-1071
- maxindc 34-34, 35-1071
- maxv 35-1073
- maxvec 34-70, 35-1075
- mbesseli 34-5, 35-1077
- mean 34-20
- meanc 34-20, 35-1080
- median 34-20, 34-20, 35-1081
- memory, clear all 35-1107
- mergeby 34-60, 34-61, 35-1083
- mergevar 34-60, 35-1084

- merging 34-60, 35-1083
- minc 34-34, 35-1086
- minimum element 35-1086
- minimum element index 35-1087
- minindc 34-34, 35-1087
- minv 35-1089
- miscellaneous topics 36-1
- miss 34-58, 34-59, 35-1090
- missex 34-58, 34-59, 35-1093
- missing character 35-1104
- missing values 34-58, 35-518, 35-524, 35-945, 35-1090, 35-1093, 35-1104, 35-1173, 35-1626, 35-1627
- missrv 34-58, 34-59, 35-1090
- moment 34-21, 35-1095
- moment matrix 34-21, 34-21, 35-1097
- momentd 34-21, 35-1097
- Moore-Penrose pseudo-inverse 34-11, 34-11, 35-1197, 35-1199
- movingave 34-21, 35-1101
- movingaveExpwgt 34-21, 35-1102
- movingaveWgt 34-21, 35-1103
- msym 34-58, 35-1104
- multi-threading 34-63, 35-1834, 35-1835, 35-1839, 35-1840
- multiplicative sequence 35-1639
- N-dimensional arrays 34-36, 34-40, 35-61, 35-63, 35-65, 35-89, 35-92, 35-110, 35-112, 35-114, 35-120, 35-122, 35-124, 35-130
- NaN 35-945, 35-1626
- new 34-70, 35-1107
- nextindex 34-42, 35-1108
- nextn 34-5, 35-1110
- nextnevn 34-5, 35-1110
- nextwind 34-89, 35-1112
- nonlinear equations 34-23, 34-23, 35-571, 35-576
- Normal distribution 34-27, 34-27, 35-248, 35-1017, 35-1018, 35-1020
- Normal distribution, bivariate 35-1013, 35-1014
- Normal distribution, multivariate 35-1016
- Normal distribution, trivariate 35-261
- ntos 35-1112
- null 34-11, 34-13, 35-1115
- null space 35-1115, 35-1117

- null1 34-11, 34-13, 35-1117
- numCombinations 34-21, 35-1118
- obsolete commands 36-42
- ols 34-21, 34-67, 35-1120
- olsmt 34-21, 35-1129
- olsmtControlCreate 34-21, 35-1140
- olsqr 34-21, 35-1141
- olsqr2 34-21, 35-1142
- olsqrmt 34-21, 35-1143
- ones 34-32, 34-33, 35-1145
- open 34-55, 34-56, 35-1146
- optimization 34-23
- optn 34-5, 35-1152
- optnevn 34-5, 35-1152
- orth 34-11, 34-13, 35-1154
- orthogonal complement 35-1116
- orthonormal 34-11, 34-11, 34-11, 35-1115, 35-1117, 35-1154
- OS functions 34-69
- output 34-78, 34-78, 34-79, 35-1156
- output functions 34-78
- outtyp (dataloop) 34-58, 35-1159
- outwidth 34-78, 35-1160
- pacf 34-21, 35-1162
- packedToSp 34-38, 35-1171
- packr 34-58, 35-1173
- pairwise deletion 35-518, 35-524, 35-1096
- parse 34-72, 35-1175
- pause 34-64, 35-1179
- pdfBinomial 34-29, 35-1179
- pdfCauchy 34-29, 35-1181
- pdfexp 34-29, 35-1182
- pdfGenPareto 34-29, 35-1183
- pdfHyperGeo 34-29, 35-1185
- pdfLaplace 34-29, 35-1187
- pdflogistic 34-29, 35-1188
- pdfn 34-29, 35-1189
- pdfPoisson 34-29, 35-1190
- pdfRayleigh 34-29, 35-1192
- pdfWeibull 34-29, 35-1193
- pdfWishartInv 34-29, 35-1194
- pi 34-5, 35-1196
- pinv 34-11, 35-1197

- pinvmt 34-11, 35-1199
- plot customization 35-1246
- plotAddArea 34-81, 35-1200
- plotAddArrow 35-1204
- plotAddBar 34-81, 34-81, 35-1206
- plotAddBox 34-81, 35-1207
- plotAddErrorBar 35-1208
- plotAddHist 34-81, 35-1216
- plotAddHistF 34-81, 35-1217
- plotAddHistP 34-81, 35-1218
- plotAddPolar 34-81, 35-1219
- plotAddScatter 34-81, 35-1220
- plotAddShape 35-1221
- plotAddSurface 34-81, 35-1223
- plotAddTextbox 35-1228
- plotAddTS 34-81, 35-1231
- plotAddXY 34-81, 35-1232
- plotArea 34-80, 35-1233
- plotBar 34-80, 35-1234
- plotBox 34-80, 35-1236
- plotCDFEmpirical 34-80, 35-1238
- plotClearLayout 34-82, 35-1241
- plotContour 34-80, 35-1243
- plotCustomLayout 34-82, 35-1246
- plotGetDefaults 34-82, 35-1247
- plotHist 34-80, 35-1249
- plotHistF 34-80, 35-1250
- plotHistP 34-80, 35-1250
- plotLayout 34-82, 35-1251
- plotLogLog 34-80, 35-1253
- plotLogX 34-80, 35-1253
- plotLogY 34-80, 35-1254
- plotOpenWindow 34-82, 35-1255
- plotPolar 34-80, 35-1256
- plotSave 34-82, 35-1257
- plotScatter 34-80, 35-1260
- plotSetAxesPen 34-82, 35-1261
- plotSetBar 34-82, 35-1262
- plotSetBkdColor 34-82, 35-1265
- plotSetColorMap 34-82, 35-1267
- plotSetContourLabels 34-82, 35-1270
- plotSetFill 34-82, 35-1273
- plotSetGrid 34-82, 35-1275
- plotSetLegend 34-82, 35-1276

- plotSetLegendFont 34-82, 35-1278
- plotSetLineColor 34-82, 35-1280
- plotSetLineStyle 34-83, 35-1281
- plotSetLineSymbol 34-83, 35-1282
- plotSetLineThickness 34-83, 35-1285
- plotSetNewWindow 34-83, 35-1286
- plotSetTextInterpreter 34-83, 35-1287
- plotSetTitle 34-83, 35-1292
- plotSetWhichYAxis 34-83, 35-1295
- plotSetXLabel 34-83, 35-1296
- plotSetXRange 34-83, 35-1299
- plotSetXTicCount 34-83, 35-1300
- plotSetXTicInterval 34-83, 35-1303
- plotSetXTicLabel 34-83, 35-1306
- plotSetYLabel 34-83, 35-1311
- plotSetYRange 34-83, 35-1314
- plotSetYTicCount 34-83, 35-1315
- plotSetZLabel 34-84, 35-1319
- plotSetZLevels 34-84, 35-1321
- plotSurface 34-80, 35-1325
- plotTS 34-81, 35-1327
- plotXY 34-81, 35-1331
- pointer 35-1041
- Poisson 35-250, 35-251, 35-1190, 35-1585, 35-1587
- polar 34-5, 35-1332, 35-1848, 35-1853
- polar plot 34-80, 34-81, 35-1219, 35-1256
- polychar 34-13, 35-1333
- polyeval 34-13, 35-1334
- polygamma 34-5, 35-1336
- polyint 34-13, 35-1339
- polymake 34-14, 35-1340
- polymat 34-14, 35-1342
- polymroot 35-1343
- polymult 34-14, 35-1345
- polynomial 34-13, 34-13, 34-14, 34-14, 34-14
- polynomial interpolation 34-13, 35-1339
- polynomial operations 34-13
- polynomial regression 35-1343
- polynomial, characteristic 35-1333
- polynomial, evaluation 35-1334
- polynomial, roots 35-1340, 35-1347
- polyroot 34-14, 35-1347

- pop 34-64, 34-66, 35-1348
- powerM 34-11, 35-1349
- PQG fonts 35-657
- pqgwin 34-89, 35-1351
- precision control 34-30
- predicted values 35-1142
- previousindex 34-42, 35-1352
- princomp 34-21, 35-1353
- print 34-78, 34-78, 34-79, 35-1355
- printdos 34-78, 35-1366
- printfm 34-79, 35-1367
- printfmt 34-78, 34-78, 34-79, 35-1370
- probability density function, Normal 34-29
- probabilty mass function 35-1190
- proc 34-66, 35-1373
- procedure 34-66, 35-1373
- procedures 34-66
- prodc 34-34, 35-1375
- products 35-1375
- program control 34-64
- program space 35-1654
- program, run 35-1605
- pseudo-inverse 35-1197, 35-1199
- psi 34-5, 35-1376
- Publication Quality Graphics 35-513, 35-567, 35-1620, 35-1621, 35-1622
- putarray 34-41, 35-1377
- putf 34-72, 35-1379
- putvals 34-36, 35-1380
- PV structure 35-1382, 35-1383, 35-1383, 35-1385, 35-1387, 35-1388, 35-1389, 35-1390, 35-1392, 35-1394, 35-1396, 35-1398, 35-1400, 35-1402, 35-1405, 35-1407, 35-1407
- pvCreate 34-43, 35-1382
- pvGetIndex 34-43, 35-1383
- pvGetParNames 34-43, 35-1383
- pvGetParVector 34-43, 35-1385
- pvLength 34-43, 35-1387
- pvList 34-43, 35-1388
- pvPack 34-43, 35-1389
- pvPacki 34-43, 35-1390
- pvPackm 34-43, 35-1392
- pvPackmi 34-43, 35-1394
- pvPacks 34-43, 35-1396

- pvPacksi 34-43, 35-1398
- pvPacksm 34-43, 35-1400
- pvPacksmi 34-44, 35-1402
- pvPutParVector 34-44, 35-1405
- pvTest 34-44, 35-1407
- pvUnpack 34-44, 35-1407
- QNewton 34-24, 35-1409
- QNewtonmt 34-24, 35-1412
- QNewtonmtControlCreate 34-24, 35-1418
- QNewtonmtOutCreate 34-24, 35-1419
- QNewtonSet 35-1420
- QProg 34-24, 35-1420
- QProgmt 34-24, 35-1422
- QProgmtInCreate 34-24, 35-1425
- qqr 34-11, 34-13, 35-1426
- qqre 34-11, 35-1428
- qqrep 34-11, 35-1431
- qr 34-11, 35-1433
- QR decomposition 34-11, 34-11, 34-11, 34-11, 34-11, 34-11, 34-11, 34-11, 34-12, 34-12, 34-12, 34-12, 34-12, 34-13, 34-21, 34-21, 34-21, 35-1141, 35-1142, 35-1143, 35-1426, 35-1428, 35-1431, 35-1433, 35-1435, 35-1438, 35-1443, 35-1446, 35-1449, 35-1458, 35-1460, 35-1462
- qre 34-11, 35-1435
- qrep 34-11, 35-1438
- qrsol 34-11, 34-13, 35-1440
- qrtsol 34-11, 35-1441
- qtyr 34-11, 35-1443
- qtyre 34-11, 35-1446
- qtyrep 34-12, 35-1449
- quadrature 35-919, 35-922, 35-926
- quantile 34-21, 35-162, 35-165, 35-1451
- quantiled 34-22, 35-1454
- query building 34-50
- query data retrieval 34-52
- query information 34-50
- query iteration 34-52
- query manipulation 34-50
- qyr 34-12, 35-1458
- qyre 34-12, 35-1460
- qyrep 34-12, 34-13, 35-1462

- qz 34-12
- random numbers 34-14, 34-16, 34-16,
34-16, 34-16, 34-16, 34-16, 34-16,
34-16, 34-16, 34-16, 34-16, 34-16,
34-17, 34-17, 34-17, 34-22, 35-
1520, 35-1522, 35-1524, 35-1527,
35-1529, 35-1530, 35-1532, 35-
1534, 35-1541, 35-1543, 35-1544,
35-1546, 35-1548, 35-1550, 35-
1551, 35-1553, 35-1555, 35-1574,
35-1575, 35-1577, 35-1579, 35-
1581, 35-1583, 35-1585, 35-1587,
35-1590, 35-1593, 35-1594
- rank 34-12, 34-13, 35-1470
- rankindx 34-34, 35-1471
- Rayleigh 35-252, 35-254, 35-1192
- readr 34-55, 34-56, 35-1472
- real 34-36, 35-1474
- reclassify 34-58, 35-1475
- reclassifyCuts 34-59, 35-1478
- recode 34-59, 34-59, 34-59, 35-1483
- recode (dataloop) 34-59, 35-1486
- recode) 34-59
- recserar 34-30, 35-1488
- recsercp 34-14, 34-30, 35-1490
- recserrc 34-14, 34-30, 35-1492
- reduced row echelon form 35-1603
- regression 35-1042, 35-1044
- relative error 35-240, 35-247, 35-256
- renamefile 35-1494
- rerun 34-90, 35-1495
- rescale 35-1495
- reserved words 36-57
- reshape 34-36, 35-1501
- residuals 35-1123, 35-1135, 35-1142
- retp 35-1503
- return 34-66, 34-66, 35-1503
- rev 34-36, 35-1504
- rfft 34-14, 35-1506
- rffti 34-14, 35-1507
- rfftip 34-14, 35-1507
- rfftn 34-14, 35-1509
- rfftnp 34-14, 35-1511
- rfftp 34-14, 35-1513
- rndBernoulli 34-14, 35-1514, 35-1516
- rndBeta 34-14, 34-15
- rndCauchy 34-15, 35-1519
- rndChiSquare 35-1520

<code>rndcon</code> 35-1522	<code>rndLCnb</code> 35-1565
<code>rndCreateState</code> 34-15, 35-1524	<code>rndLCp</code> 35-1567
<code>rndExp</code> 34-15, 35-1527	<code>rndLCu</code> 35-1569
<code>rndgam</code> 35-1529	<code>rndLCvm</code> 35-1572
<code>rndGamma</code> 34-15, 35-1530	<code>rndLogNorm</code> 34-16, 35-1574
<code>rndGeo</code> 34-15, 35-1532	<code>rndmult</code> 35-1522
<code>rndGumbel</code> 34-15, 35-1534	<code>rndMVn</code> 34-16, 35-1575
<code>rndHyperGeo</code> 34-15, 35-1535	<code>rndMVt</code> 34-16, 35-1577
<code>rndi</code> 34-15, 35-1538	<code>rndn</code> 34-16, 35-1579
<code>rndKMbeta</code> 34-15, 35-1541	<code>rndnb</code> 35-1581
<code>rndKMgam</code> 34-15, 35-1543	<code>rndNegBinomial</code> 34-16, 35-1583
<code>rndKMi</code> 34-15, 35-1544	<code>rndp</code> 35-1585
<code>rndKMn</code> 34-15, 35-1546	<code>rndPoisson</code> 34-16, 35-1587
<code>rndKMnb</code> 34-16, 35-1548	<code>rndseed</code> 34-16, 34-17, 35-1522
<code>rndKMp</code> 34-16, 35-1550	<code>rndStateSkip</code> 34-16, 35-1588
<code>rndKMu</code> 34-16, 35-1551	<code>rndu</code> 34-16, 35-1590
<code>rndKMvm</code> 34-16, 35-1553	<code>rndvm</code> 34-22, 35-1593
<code>rndLaplace</code> 34-16, 35-1555	<code>rndWeibull</code> 34-17, 35-1594
<code>rndLCbeta</code> 35-1557	<code>rndWishart</code> 34-17, 35-1595
<code>rndLCgam</code> 35-1559	<code>rndWishartInv</code> 34-17, 35-1597
<code>rndLCi</code> 35-1560	<code>rotater</code> 34-36, 35-1598
<code>rndLCn</code> 35-1563	<code>round</code> 34-30, 34-30, 35-1600

- round down 35-652
- round up 35-267
- rows 34-35, 34-35, 35-1601
- rowsf 34-35, 34-35, 35-1602
- rref 34-12, 34-13, 35-1603
- run 34-64, 35-1605
- sampleData 35-1608
- satostrC 34-79, 35-1610
- save 34-33, 34-68, 34-68, 35-1613
- saveall 34-68, 35-1616
- saved 34-33, 34-55, 35-1617
- savestruct 34-44, 35-1619
- savewind 34-89, 35-1620
- scalar error code 34-71
- scalar expression 35-876
- scale 34-86, 35-1621
- scale3d 34-86, 35-1622
- scalerr 34-71, 35-1623
- scalinfnanmiss 34-59, 35-1626
- scaling 35-1621, 35-1622
- scalmiss 34-58, 34-59, 35-1627
- scatter plot 34-80, 35-1220, 35-1260
- schtoc 34-12, 35-1628
- schur 34-10, 34-12, 35-1630
- Schur 34-12, 35-1628, 35-1630
- scientific functions 34-4
- screen 34-79, 35-1633
- searchsourcepath 34-69, 35-1634
- seekr 34-55, 34-56, 35-1635
- select (dataloop) 34-59, 35-1636
- selif 34-36, 34-38, 34-38, 35-1637
- seqa 34-30, 35-1639
- seqm 34-30, 35-1639
- sequence function 35-1639
- sequence functions 34-29
- series functions 34-29
- set difference function 35-1642
- setarray 34-41, 35-1640
- setdif 34-36, 35-1642
- setdifsa 35-1645
- setvars 34-57, 34-57, 35-1646
- setvwrmode 34-89, 35-1647
- setwind 34-89, 35-1648
- shell 34-70, 35-1649

- shiftr 34-36, 35-1651
- show 34-70, 35-1653
- Simpson's method 35-938
- sin 34-5, 35-1656
- sine, inverse 34-4, 35-107
- singleindex 34-42, 35-1657
- singular value decomposition 34-12, 34-12, 35-309, 35-1470, 35-1790, 35-1792, 35-1795, 35-1801
- singular values 34-12, 35-1788, 35-1798
- sinh 34-5, 35-1659
- sleep 34-64, 35-1660
- solpd 34-12, 34-13, 35-1661
- sort data file 35-1666
- sort index 35-1669
- sort, heap sort 35-1667
- sort, multiple columns 34-60, 35-1670
- sort, quicksort 35-1664
- sortc 34-60, 34-61, 35-1664
- sortcc 34-60, 34-61, 34-61, 35-1664
- sortd 34-60, 34-61, 34-61, 35-1666
- sorthc 34-60, 34-61, 35-1667
- sorthcc 34-60, 35-1667
- sortind 34-60, 34-61, 35-1669
- sortindc 34-60, 34-61, 35-1669
- sorting 34-60
- sortmc 34-60, 34-61, 35-1670
- sortr 34-61, 35-1672
- sortrc 34-61, 35-1672
- sparse matrices 34-38, 35-1674, 35-1677, 35-1678, 35-1681, 35-1682, 35-1684, 35-1687, 35-1690, 35-1691, 35-1694, 35-1695, 35-1697, 35-1698, 35-1707, 35-1709, 35-1711, 35-1712, 35-1713, 35-1714
- spBiconjGradSol 34-38, 35-1674
- spChol 34-38, 35-1677
- spConjGradSol 34-38, 35-1678
- spCreate 35-1681
- spCreatee 34-38
- spDenseSubmat 34-39, 35-1682
- spDiagRvMat 34-39, 35-1684
- spEigv 34-39, 35-1687
- spEye 34-39, 35-1690
- spGetNumNZE 34-39, 35-1691
- spGetNZE 34-39
- spLDL 34-39, 35-1694

- spline 34-5, 35-1693
- spLU 34-39
- spNumNZE 35-1697
- spOnes 34-39, 35-1698
- spreadSheetReadM 34-44, 35-1700
- spreadSheetReadSA 35-1702
- SpreadsheetReadSA 34-44
- spreadsheets 34-44, 34-45, 35-1700, 35-1702, 35-1704, 35-1931, 35-1932, 35-1935, 35-1937, 35-1939, 35-1944, 35-1948, 35-1952, 35-1955
- spreadSheetWrite 34-44, 35-1704
- spScale 35-1707
- spSubmat 34-39, 35-1709
- spToDense 34-39, 35-1711
- spTrTDense 34-39, 35-1712
- spTScalar 34-39, 35-1713
- spZeros 34-39, 35-1714
- sqpSolve 34-24, 35-1716
- sqpSolveMT 34-24, 35-1723
- sqpSolveMTControl structure 34-24, 35-1731
- sqpSolveMTControlCreate 35-1731
- sqpSolveMTlagrangeCreate 34-24, 35-1732
- sqpSolveMToutCreate 34-24, 35-1733
- sqpSolveSet 34-25, 35-1734
- sqrt 34-6, 35-1735
- square root 35-1735
- standard deviation 35-120, 35-122, 35-522, 35-525, 35-1736, 35-1738
- standard deviation of residual 35-1124, 35-1134
- standard errors 35-1124, 35-1134
- statistical distributions 34-25
- statistical functions 34-19
- statistics, descriptive 35-516, 35-523
- stdc 34-22, 35-1736
- stdsc 34-22, 35-1738
- Stirling's formula 35-1022
- stocv 34-72, 35-1739
- stof 34-72, 35-1740
- stop 34-64, 35-1741
- stop program 35-1741
- strcombine 34-72, 35-1742
- strindx 34-72, 34-74, 35-1744

- string arrays 35-1760, 35-1762
- string handling 34-71
- string index 35-1744, 35-1748
- string length 35-1745
- string, substring 35-1751
- strjoin 34-72, 35-1749
- strlen 34-73, 34-74
- strput 34-73, 35-1746
- strrindx 34-73, 34-74, 35-1748
- strsect 34-73, 34-74, 35-1751
- strsplit 34-73, 35-1753
- strsplitPad 34-73, 35-1757
- strtodt 34-76, 35-1758
- strtodf 34-73, 35-1760
- strtofcplx 34-73, 35-1762
- strtrim 34-73, 35-1763
- strtriml 34-73, 35-1765
- strtrimr 34-73, 35-1766
- strtrunc 34-73, 35-1767
- strtruncl 34-74, 35-1768
- strtruncpad 34-74, 35-1769
- strtruncr 34-74, 35-1770
- structure, DS 34-42
- structure, PV 34-43, 34-43, 34-43, 34-43, 34-43, 34-43, 34-44, 34-44, 34-44
- structures 34-42
- structures, matrices of 35-1619
- submat 34-37, 35-1770
- submatrix 35-1684, 35-1709, 35-1770
- subroutine 35-1503
- subroutines 34-66
- subsample 35-619
- subscat 34-59, 34-59, 35-1772
- substitution 35-1776
- substute 34-59, 34-59, 35-1776
- subvec 34-37, 35-1779
- sum 35-1781, 35-1783
- sumc 35-1781
- sumr 34-35, 35-1783
- sunc 34-35
- surface 34-85, 34-88, 35-1785
- surface plot 34-80
- svd 34-12, 34-13, 35-1788
- svd1 34-12, 35-1790

- svd2 34-12, 35-1792
- svdcusv 34-12, 35-1795
- svds 34-12, 35-1798
- svdusv 34-12, 34-13, 35-1801
- sweep inverse 35-941
- sylvester 34-12, 35-1804
- Sylvester 35-1804
- symbol table 35-1107
- symbol table type 35-1865, 35-1867
- symbols, allocate maximum number 35-1107
- sysstate 34-68, 35-1806
- system 34-64, 35-1806, 35-1828
- t distribution, Student's 34-29, 35-221, 35-227, 35-233, 35-255, 35-257, 35-258, 35-1577
- tab 34-79, 35-1830
- tan 34-6, 35-1831
- tanh 34-6, 35-1832
- tempname 34-55, 35-1833
- text files 34-45
- thickness, line 35-1285
- threadBegin 34-63, 34-63
- ThreadBegin 35-1834
- threadEnd 34-63, 34-63
- ThreadEnd 35-1835
- threadEndFor 34-63
- threadfor 35-1836
- threadFor 34-63
- threadfor loop 35-1836
- threadJoin 34-63, 34-63
- ThreadJoin 35-1839
- threads 34-63, 35-1834, 35-1835, 35-1839, 35-1840
- threadStat 34-63, 34-63, 34-63
- ThreadStat 35-1840
- time 34-76, 35-1841
- time and date functions 34-75
- time series 35-1231
- time, elapsed 35-594, 35-595, 35-597
- timedt 34-76, 35-1841
- timestr 34-76, 35-1842
- timeutc 34-76, 35-1843
- timing functions 35-874
- title 34-87, 35-1844

- tkf2eps 34-89, 35-1846
- tkf2ps 34-90, 35-1847
- tocart 34-6, 35-1848
- todaydt 34-76, 35-1848
- toeplitz 34-22, 34-22, 35-1849
- Toeplitz matrix 35-1849
- token 34-74, 35-1850
- topolar 34-6, 35-1853
- trace 34-71, 34-71, 35-1853
- trace program execution 35-1853
- transpose 35-130
- trap 34-71, 35-1855
- trap flag 35-1855, 35-1859
- trap state 35-1625
- trapchk 34-71, 35-1859
- triangular matrix, lower 35-1057, 35-1060
- triangular matrix, upper 35-274, 35-278, 35-1060, 35-1440, 35-1441
- trigamma 34-6, 35-1861
- trimr 34-37, 35-1862
- trivariate Normal 35-261
- trunc 34-30, 34-30, 35-1863
- truncating 35-1767, 35-1768, 35-1769, 35-1770, 35-1863
- type 34-70, 35-1865
- typecv 34-70, 35-1867
- typef 34-55, 35-1869
- underdetermined 35-1124, 35-1134
- union 34-37, 35-1871
- unionsa 35-1872
- uniqindx 34-61, 34-61, 35-1873
- uniqindxsa 34-61, 35-1875
- unique 34-61, 34-61, 35-1876
- uniquesa 34-61, 35-1882
- until 34-65, 35-497
- upmat 34-37, 34-38, 35-1883
- upmat1 34-37, 34-38, 35-1883
- upper 34-74, 35-1885
- upper triangular matrix 35-1060, 35-1440, 35-1441
- use 34-68, 35-1886
- user-defined function 35-953, 35-1373
- utctodt 34-76, 35-1888
- utctodtv 34-76, 35-1889
- utrisol 34-25, 35-1891

- vals 34-74, 35-1892
- varCovM 34-22, 35-1894
- varCovMS 34-22, 35-1896
- varCovX 34-22, 35-1894
- varCovXS 34-22, 35-1896
- varget 34-74, 35-1898, 35-1900
- vargetl 34-74
- variable names 34-56, 35-1646
- variance 35-518, 35-525
- variance-covariance matrix 34-22, 34-22, 34-22, 34-22, 34-22, 34-23, 34-23, 35-1894, 35-1896, 35-1909, 35-1910
- varindxi 35-1146
- varmall 34-22, 35-1902
- varmares 34-22, 35-1903
- varput 34-74, 35-1905
- varputl 34-74, 35-1906
- vartypef 34-57, 34-57, 35-1908
- vcm 34-22, 35-1909
- vcms 34-22, 35-1910
- vcx 34-23, 35-1909
- vcxs 34-23, 35-1910
- vec 34-37, 35-1911
- vech 34-37, 34-38, 35-1913
- vecr 34-37, 35-1911
- vector (dataloop) 34-59, 35-1914
- vectors 34-32
- vget 34-37, 35-1915
- view 34-90, 35-1916
- viewxyz 34-90, 35-1917
- vlist 34-37, 35-1918
- vnamecv 34-37, 35-1919
- volume 34-90, 35-1920
- vput 34-37, 34-37, 35-1921
- vread 34-37, 35-1922
- vtypecv 34-37, 35-1923
- wait 34-77, 34-78, 35-1924
- waitc 34-77, 34-78, 35-1924
- walkindex 34-42, 35-1924
- Weibull 35-263, 35-265, 35-1193, 35-1594
- weighted count 35-331
- while 34-65, 35-497
- window 34-89, 35-1926

window, clear 35-289
workbox 35-1916, 35-1918
workspace 34-70
writer 34-56, 35-1927
X-axis 35-1303, 35-1306
xlabel 34-87, 35-1930
xlsGetSheetCount 34-44, 35-1931
xlsGetSheetSize 34-44, 35-1932
xlsGetSheetTypes 34-44, 35-1935
xlsMakeRange 34-45, 35-1937
xlsReadM 34-34, 34-45, 35-1939
xlsReadSA 34-45, 35-1944
xlsWrite 34-45, 35-1948
xlsWriteM 34-45, 35-1952
xlsWriteSA 34-45, 35-1955
xpnd 34-38, 34-38, 34-38, 35-1958
xtics 34-86, 35-1960
xy 34-86, 35-1232, 35-1961
xyz 35-1962
ylabel 34-87, 35-1964
ytics 34-86, 35-1965
zeroes 34-32, 35-1967
zeros 34-33
zeta 34-6, 35-1968
zlabel 34-87, 35-1969
ztics 34-86, 35-1970