

# GAUSS<sup>TM</sup>

## *Language Reference*

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# Command Reference

# Introduction

# 29

The GAUSS LANGUAGE REFERENCE describes each of the commands, procedures and functions available in the **GAUSS**<sup>TM</sup> 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 **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.

### 29.1 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 ...”
<b>bold text</b>	emphasis	“ <b>...not supported under UNIX.</b> ”
<i>italic text</i>	variables	“... If <i>vnames</i> is a string or has fewer elements than <i>x</i> has columns, it will be ...”

Text Style	Use	Example
monospace	code example	if scalerr(cm); cm = inv(x); endif;
	- or -	
	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 <b>create...</b> ”
SMALL CAPS	reference to section of the manual	“...see OPERATOR PRECEDENCE, Section 11.7...”

## 29.2 Command Components

The following list describes each of the components used in the COMMAND REFERENCE, Chapter 31.

PURPOSE	Describes what the command or function does.
LIBRARY	Lists the library that needs to be activated to access the function.
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.

### 29.3 Using This Manual

Users who are new to **GAUSS** should make sure they have familiarized themselves with **LANGUAGE FUNDAMENTALS**, Chapter 10, before proceeding here. That chapter contains the basics of **GAUSS** programming.

In all, there are over 700 routines described in this **GAUSS LANGUAGE REFERENCE**. We suggest that new **GAUSS** users skim through Chapter 30, and then browse through Chapter 31, the main part of this manual. Here, users can familiarize themselves with the kinds of tasks that **GAUSS** can handle easily.

Chapter 30 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 31.

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 31 under the discussion of that function.

## 29.4 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.

### 29.4.1 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.

### 29.4.2 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.



# Commands by Category 30

## 30.1 Mathematical Functions

### Scientific Functions

<b>abs</b>	Returns absolute value of argument.
<b>arccos</b>	Computes inverse cosine.
<b>arcsin</b>	Computes inverse sine.
<b>atan</b>	Computes inverse tangent.
<b>atan2</b>	Computes angle given a point $x,y$ .
<b>besselj</b>	Computes Bessel function, first kind.
<b>bessely</b>	Computes Bessel function, second kind.
<b>boxcox</b>	Computes the Box-Cox function.
<b>cos</b>	Computes cosine.

<b>cosh</b>	Computes hyperbolic cosine.
<b>curve</b>	Computes a one-dimensional smoothing curve.
<b>digamma</b>	Computes the digamma function.
<b>exp</b>	Computes the exponential function of $x$ .
<b>fmod</b>	Computes the floating-point remainder of $x/y$ .
<b>gamma</b>	Computes gamma function value.
<b>gammai</b>	Compute the inverse incomplete gamma function.
<b>ln</b>	Computes the natural log of each element.
<b>lnfact</b>	Computes natural log of factorial function.
<b>log</b>	Computes the $\log_{10}$ of each element.
<b>mbesseli</b>	Computes modified and exponentially scaled modified Bessels of the first kind of the $n^{th}$ order.
<b>nextn, nextnevn</b>	Returns allowable matrix dimensions for computing FFT's.
<b>optn, optnevn</b>	Returns optimal matrix dimensions for computing FFT's.
<b>pi</b>	Returns $\pi$ .
<b>polar</b>	Graphs data using polar coordinates.
<b>sin</b>	Computes sine.
<b>sinh</b>	Computes the hyperbolic sine.
<b>spline</b>	Computes a two-dimensional interpolatory spline.
<b>sqrt</b>	Computes the square root of each element.
<b>tan</b>	Computes tangent.
<b>tanh</b>	Computes hyperbolic tangent.

---

<b>tocart</b>	Converts from polar to Cartesian coordinates.
<b>topolar</b>	Converts from Cartesian to polar coordinates.
<b>trigamma</b>	Computes trigamma function.

All trigonometric functions take or return values in radian units.

## Differentiation and Integration

<b>gradMT</b>	Computes numerical gradient.
<b>gradMTm</b>	Computes numerical gradient with mask.
<b>gradp</b>	Computes first derivative of a function.
<b>hessMT</b>	Computes numerical Hessian.
<b>hessMTg</b>	Computes numerical Hessian using gradient procedure.
<b>hessMTgw</b>	Computes numerical Hessian using gradient procedure with weights.
<b>hessMTm</b>	Computes numerical Hessian with mask.
<b>hessMTmw</b>	Computes numerical Hessian with mask and weights.
<b>hessMTw</b>	Computes numerical Hessian with weights.
<b>hessp</b>	Computes second derivative of a function.
<b>integrat2</b>	Integrates a 2-dimensional function over a user-defined region.
<b>integrat3</b>	Integrates a 3-dimensional function over a user-defined region.
<b>inthp1</b>	Integrates a user-defined function over an infinite interval.

---

<b>inthp2</b>	Integrates a user-defined function over the $[a, +\infty)$ interval.
<b>inthp3</b>	Integrates a user-defined function over the $[a, +\infty)$ interval that is oscillatory.
<b>inthp4</b>	Integrates a user-defined function over the $[a, b]$ interval.
<b>inthpControlCreate</b>	Creates default <b>inthpControl</b> structure.
<b>intquad1</b>	Integrates a 1-dimensional function.
<b>intquad2</b>	Integrates a 2-dimensional function over a user-defined rectangular region.
<b>intquad3</b>	Integrates a 3-dimensional function over a user-defined rectangular region.
<b>intsimp</b>	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**.

To get a greater degree of accuracy than that provided by **intquad1**, use **intsimp** for 1-dimensional integration.

## Linear Algebra

<b>balance</b>	Balances a matrix.
<b>band</b>	Extracts bands from a symmetric banded matrix.

---

<b>bandchol</b>	Computes the Cholesky decomposition of a positive definite banded matrix.
<b>bandcholsol</b>	Solves the system of equations $Ax = b$ for $x$ , given the lower triangle of the Cholesky decomposition of a positive definite banded matrix $A$ .
<b>bandltsol</b>	Solves the system of equations $Ax = b$ for $x$ , where $A$ is a lower triangular banded matrix.
<b>bandrv</b>	Creates a symmetric banded matrix, given its compact form.
<b>bandsolpd</b>	Solves the system of equations $Ax = b$ for $x$ , where $A$ is a positive definite banded matrix.
<b>chol</b>	Computes Cholesky decomposition, $X = Y'Y$ .
<b>choldn</b>	Performs Cholesky downdate on an upper triangular matrix.
<b>cholsol</b>	Solves a system of equations given the Cholesky factorization of a matrix.
<b>cholup</b>	Performs Cholesky update on an upper triangular matrix.
<b>cond</b>	Computes condition number of a matrix.
<b>crout</b>	Computes Crout decomposition, $X = LU$ (real matrices only).
<b>croutp</b>	Computes Crout decomposition with row pivoting (real matrices only).
<b>det</b>	Computes determinant of square matrix.
<b>detl</b>	Computes determinant of decomposed matrix.
<b>hess</b>	Computes upper Hessenberg form of a matrix (real matrices only).
<b>inv</b>	Inverts a matrix.
<b>invpd</b>	Inverts a positive definite matrix.
<b>invswp</b>	Computes a generalized sweep inverse.

---

<b>lapeighb</b>	Computes eigenvalues only of a real symmetric or complex Hermitian matrix selected by bounds.
<b>lapeighi</b>	Computes eigenvalues only of a real symmetric or complex Hermitian matrix selected by index.
<b>lapeighvb</b>	Computes eigenvalues and eigenvectors of a real symmetric or complex Hermitian matrix selected by bounds.
<b>lapeighvi</b>	Computes selected eigenvalues and eigenvectors of a real symmetric or complex Hermitian matrix.
<b>lapgeig</b>	Computes generalized eigenvalues for a pair of real or complex general matrices.
<b>lapgeigh</b>	Computes generalized eigenvalues for a pair of real symmetric or Hermitian matrices.
<b>lapgeighv</b>	Computes generalized eigenvalues and eigenvectors for a pair of real symmetric or Hermitian matrices.
<b>lapgeigv</b>	Computes generalized eigenvalues, left eigenvectors, and right eigenvectors for a pair of real or complex general matrices.
<b>lapgschur</b>	Computes the generalized Schur form of a pair of real or complex general matrices.
<b>lapgsvdcst</b>	Computes the generalized singular value decomposition of a pair of real or complex general matrices.
<b>lapgsvds</b>	Computes the generalized singular value decomposition of a pair of real or complex general matrices.
<b>lapgsvdst</b>	Computes the generalized singular value decomposition of a pair of real or complex general matrices.
<b>lapsvdcusv</b>	Computes the singular value decomposition a real or complex rectangular matrix, returns compact $u$ and $v$ .
<b>lapsvds</b>	Computes the singular values of a real or complex rectangular matrix.

---

<b>lapsvdusv</b>	Computes the singular value decomposition a real or complex rectangular matrix.
<b>lu</b>	Computes LU decomposition with row pivoting (real and complex matrices).
<b>null</b>	Computes orthonormal basis for right null space.
<b>null1</b>	Computes orthonormal basis for right null space.
<b>orth</b>	Computes orthonormal basis for column space $x$ .
<b>pinv</b>	Generalized pseudo-inverse: Moore-Penrose.
<b>pinvmt</b>	Generalized pseudo-inverse: Moore-Penrose.
<b>qqr</b>	QR decomposition: returns $Q_1$ and $R$ .
<b>qqre</b>	QR decomposition: returns $Q_1$ , $R$ and a permutation vector, $E$ .
<b>qqrep</b>	QR decomposition with pivot control: returns $Q_1$ , $R$ and $E$ .
<b>qr</b>	QR decomposition: returns $R$ .
<b>qre</b>	QR decomposition: returns $R$ and $E$ .
<b>qrep</b>	QR decomposition with pivot control: returns $R$ and $E$ .
<b>qrsol</b>	Solves a system of equations $Rx = b$ given an upper triangular matrix, typically the $R$ matrix from a QR decomposition.
<b>qrtsol</b>	Solves a system of equations $R'x = b$ given an upper triangular matrix, typically the $R$ matrix from a QR decomposition.
<b>qtyr</b>	QR decomposition: returns $Q'Y$ and $R$ .
<b>qtyre</b>	QR decomposition: returns $Q'Y, R$ and $E$ .
<b>qtyrep</b>	QR decomposition with pivot control: returns $Q'Y, R$ and $E$ .
<b>qyr</b>	QR decomposition: returns $QY$ and $R$ .
<b>qyre</b>	QR decomposition: returns $QY, R$ and $E$ .

---

<b>qyrep</b>	QR decomposition with pivot control: returns $QY$ , $R$ and $E$ .
<b>rank</b>	Computes rank of a matrix.
<b>rref</b>	Computes reduced row echelon form of a matrix.
<b>schtoc</b>	Reduces any $2 \times 2$ blocks on the diagonal of the real Schur matrix returned from <b>schur</b> . The transformation matrix is also updated.
<b>schur</b>	Computes Schur decomposition of a matrix (real matrices only).
<b>solpd</b>	Solves a system of positive definite linear equations.
<b>svd</b>	Computes the singular values of a matrix.
<b>svd1</b>	Computes singular value decomposition, $X = USV'$ .
<b>svd2</b>	Computes <b>svd1</b> with compact $U$ .
<b>svdcusv</b>	Computes the singular value decomposition of a matrix so that: $x = u * s * v'$ (compact $u$ ).
<b>svds</b>	Computes the singular values of a matrix.
<b>svdusv</b>	Computes the singular value decomposition of a matrix so that: $x = u * s * v'$ .

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.



---

## Eigenvalues

<b>eig</b>	Computes eigenvalues of general matrix.
<b>eigh</b>	Computes eigenvalues of complex Hermitian or real symmetric matrix.
<b>eighv</b>	Computes eigenvalues and eigenvectors of complex Hermitian or real symmetric matrix.
<b>eigv</b>	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**

## Polynomial Operations

<b>polychar</b>	Computes characteristic polynomial of a square matrix.
<b>polyeval</b>	Evaluates polynomial with given coefficients.
<b>polyint</b>	Calculates $N^{th}$ order polynomial interpolation given known point pairs.
<b>polymake</b>	Computes polynomial coefficients from roots.
<b>polymat</b>	Returns sequence powers of a matrix.
<b>polymult</b>	Multiplies two polynomials together.
<b>polyroot</b>	Computes roots of polynomial from coefficients.

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

### Fourier Transforms

<b>dffft</b>	Computes discrete 1-D FFT.
<b>dfffti</b>	Computes inverse discrete 1-D FFT.
<b>ffft</b>	Computes 1- or 2-D FFT.
<b>fffti</b>	Computes inverse 1- or 2-D FFT.
<b>ffftm</b>	Computes multi-dimensional FFT.
<b>ffftmi</b>	Computes inverse multi-dimensional FFT.
<b>ffftn</b>	Computes 1- or 2-D FFT using prime factor algorithm.
<b>rffft</b>	Computes real 1- or 2-D FFT.
<b>rfffti</b>	Computes inverse real 1- or 2-D FFT.
<b>rffftip</b>	Computes inverse real 1- or 2-D FFT from packed format FFT.
<b>rffftn</b>	Computes real 1- or 2-D FFT using prime factor algorithm.
<b>rffftnp</b>	Computes real 1- or 2-D FFT using prime factor algorithm, returns packed format FFT.
<b>rffftp</b>	Computes real 1- or 2-D FFT, returns packed format FFT.

### Random Numbers

<b>rndbeta</b>	Computes random numbers with beta distribution.
<b>rndcon</b>	Changes constant of the LC random number generator.
<b>rndgam</b>	Computes random numbers with gamma distribution.
<b>rndi</b>	Returns random integers, $0 \leq y < 2^{32}$ .
<b>rndKmbeta</b>	Computes beta pseudo-random numbers.

---

<b>rndKMgam</b>	Computes gamma pseudo-random numbers.
<b>rndKMi</b>	Returns random integers, $0 \leq y < 2^{32}$ .
<b>rndKMn</b>	Computes standard normal pseudo-random numbers.
<b>rndKMnb</b>	Computes negative binomial pseudo-random numbers.
<b>rndKMp</b>	Computes Poisson pseudo-random numbers.
<b>rndKMu</b>	Computes uniform pseudo-random numbers.
<b>rndKMvm</b>	Computes von Mises pseudo-random numbers.
<b>rndLCbeta</b>	Computes beta pseudo-random numbers.
<b>rndLCgam</b>	Computes gamma pseudo-random numbers.
<b>rndLCi</b>	Returns random integers, $0 \leq y < 2^{32}$ .
<b>rndLCn</b>	Computes standard normal pseudo-random numbers.
<b>rndLCnb</b>	Computes negative binomial pseudo-random numbers.
<b>rndLCp</b>	Computes Poisson pseudo-random numbers.
<b>rndLCu</b>	Computes uniform pseudo-random numbers.
<b>rndLCvm</b>	Computes von Mises pseudo-random numbers.
<b>rndmult</b>	Changes multiplier of the LC random number generator.
<b>rndn</b>	Computes random numbers with Normal distribution.
<b>rndnb</b>	Computes random numbers with negative binomial distribution.
<b>rndp</b>	Computes random numbers with Poisson distribution.
<b>rndseed</b>	Changes seed of the LC random number generator.
<b>rndu</b>	Computes random numbers with uniform distribution.

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

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

### Fuzzy Conditional Functions

<b>dotfeq</b>	Fuzzy . ==
<b>dotfeqmt</b>	Fuzzy . ==
<b>dotfge</b>	Fuzzy . >=
<b>dotfgemt</b>	Fuzzy . >=
<b>dotfgt</b>	Fuzzy . >
<b>dotfgtmt</b>	Fuzzy . >
<b>dotfle</b>	Fuzzy . <=
<b>dotflemt</b>	Fuzzy . <=
<b>dotflt</b>	Fuzzy . <
<b>dotflmt</b>	Fuzzy . <
<b>dotfne</b>	Fuzzy ./ =
<b>dotfnemt</b>	Fuzzy ./ =
<b>feq</b>	Fuzzy ==
<b>feqmt</b>	Fuzzy ==
<b>fge</b>	Fuzzy >=
<b>fgemt</b>	Fuzzy >=
<b>fgt</b>	Fuzzy >
<b>fgtmt</b>	Fuzzy >
<b>fle</b>	Fuzzy <=

<b>flemt</b>	Fuzzy <=
<b>flt</b>	Fuzzy <
<b>fltgt</b>	Fuzzy <
<b>fne</b>	Fuzzy / =
<b>fnemt</b>	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*.

### Statistical Functions

<b>acf</b>	Computes sample autocorrelations.
<b>astd</b>	Computes the standard deviation of the elements across one dimension of an N-dimensional array.
<b>astds</b>	Computes the ‘sample’standard deviation of the elements across one dimension of an N-dimensional array.
<b>chiBarSquare</b>	computes probability of chi-bar-square statistic
<b>combine</b>	Computes combinations of <i>n</i> things
<b>combine</b>	Computes combinations of <i>n</i> things taken <i>k</i> at a time.
<b>combined</b>	Writes combinations of <i>n</i> things taken <i>k</i> at a time to a <b>GAUSS</b> data set.
<b>ConScore</b>	computes constrained score statistic and its probability

<b>conv</b>	Computes convolution of two vectors.
<b>corrmm</b>	Computes correlation matrix of a moment matrix.
<b>corrms</b>	Computes sample correlation matrix of a moment matrix.
<b>corrvc</b>	Computes correlation matrix from a variance-covariance matrix.
<b>corrxx</b>	Computes correlation matrix.
<b>corrxs</b>	Computes sample correlation matrix.
<b>crossprd</b>	Computes cross product.
<b>design</b>	Creates a design matrix of 0's and 1's.
<b>dstat</b>	Computes descriptive statistics of a data set or matrix.
<b>dstatmt</b>	Computes descriptive statistics of a data set or matrix.
<b>dstatmtControlCreate</b>	Creates default <b>dstatmtControl</b> structure.
<b>gdaDStat</b>	Computes descriptive statistics on multiple N×1 variables in a GDA.
<b>gdaDStatMat</b>	Computes descriptive statistics on a selection of columns in a variable in a GDA.
<b>loess</b>	Computes coefficients of locally weighted regression.
<b>loessmt</b>	Computes coefficients of locally weighted regression.
<b>loessmtControlCreate</b>	Creates default <b>loessmtControl</b> structure.
<b>meanc</b>	Computes mean value of each column of a matrix.
<b>median</b>	Computes medians of the columns of a matrix.
<b>moment</b>	Computes moment matrix ( $x'x$ ) with special handling of missing values.
<b>momentd</b>	Computes moment matrix from a data set.

---

<b>movingave</b>	Computes moving average of a series.
<b>movingaveExpwgt</b>	Computes exponentially weighted moving average of a series.
<b>movingaveWgt</b>	Computes weighted moving average of a series.
<b>numCombinations</b>	Computes number of combinations of $n$ things taken $k$ at a time.
<b>ols</b>	Computes least squares regression of data set or matrix.
<b>olsmt</b>	Computes least squares regression of data set or matrix.
<b>olsmtControlCreate</b>	Creates default <b>olsmtControl</b> structure.
<b>olsqr</b>	Computes OLS coefficients using QR decomposition.
<b>olsqr2</b>	Computes OLS coefficients, residuals, and predicted values using QR decomposition.
<b>olsqrmt</b>	Computes OLS coefficients using QR decomposition.
<b>pacf</b>	Computes sample partial autocorrelations.
<b>princomp</b>	Computes principal components of a data matrix.
<b>quantile</b>	Computes quantiles from data in a matrix, given specified probabilities.
<b>quantiled</b>	Computes quantiles from data in a data set, given specified probabilities.
<b>rndvm</b>	Computes von Mises pseudo-random numbers.
<b>stdc</b>	Computes standard deviation of the columns of a matrix.
<b>stdsc</b>	Computes the ‘sample’ standard deviation of the elements in each column of a matrix.
<b>toeplitz</b>	Computes Toeplitz matrix from column vector.
<b>varmall</b>	Computes the log-likelihood of a Vector ARMA model.

---

<b>varmares</b>	Computes the residuals of a Vector ARMA model.
<b>vcm</b>	Computes a variance-covariance matrix from a moment matrix.
<b>vcms</b>	Computes a sample variance-covariance matrix from a moment matrix.
<b>vcx</b>	Computes a variance-covariance matrix from a data matrix.
<b>vcxs</b>	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.)

## Optimization and Solution

<b>eqsolve</b>	Solves a system of nonlinear equations.
<b>eqSolveMt</b>	Solves a system of nonlinear equations.
<b>eqSolveMtControlCreate</b>	Creates default <b>eqSolveMtControl</b> structure.
<b>eqSolveMtOutCreate</b>	Creates default <b>eqSolveMtOut</b> structure.
<b>eqSolveSet</b>	Sets global input used by <b>eqSolve</b> to default values.
<b>linsolve</b>	Solves $Ax = b$ using the inverse function.
<b>ltrisol</b>	Computes the solution of $Lx = b$ where $L$ is a lower triangular matrix.
<b>lusol</b>	Computes the solution of $LUx = b$ where $L$ is a lower triangular matrix and $U$ is an upper triangular matrix.
<b>QNewton</b>	Optimizes a function using the BFGS descent algorithm.



---

<b>QNewtonmt</b>	Minimizes an arbitrary function.
<b>QNewtonmtControlCreate</b>	Creates default <b>QNewtonmtControl</b> structure.
<b>QNewtonmtOutCreate</b>	Creates default <b>QNewtonmtOut</b> structure.
<b>QProg</b>	Solves the quadratic programming problem.
<b>QProgmt</b>	Solves the quadratic programming problem.
<b>QProgmtInCreate</b>	Creates an instance of a structure of type <b>QProgmtInCreate</b> with the <b>maxit</b> member set to a default value.
<b>sqpSolve</b>	Solves the nonlinear programming problem using a sequential quadratic programming method.
<b>sqpSolveMT</b>	Solves the nonlinear programming problem using a sequential quadratic programming method.
<b>sqpSolveMTControlCreate</b>	Creates an instance of a structure of type <b>sqpSolveMTcontrol</b> set to default values.
<b>sqpSolveMTlagrangeCreate</b>	Creates an instance of a structure of type <b>sqpSolveMTlagrange</b> set to default values.
<b>sqpSolveMToutCreate</b>	Creates an instance of a structure of type <b>sqpSolveMTout</b> set to default values.
<b>sqpSolveSet</b>	Resets global variables used by <b>sqpSolve</b> to default values.
<b>utrisol</b>	Computes the solution of $Ux = b$ where $U$ is an upper triangular matrix.

## Statistical Distributions

<b>cdfbeta</b>	Computes integral of beta function.
<b>cdfbvn</b>	Computes lower tail of bivariate Normal cdf.

<b>cdfbvn2</b>	Returns cdfbvn of a bounded rectangle.
<b>cdfbvn2e</b>	Returns cdfbvn of a bounded rectangle.
<b>cdfchic</b>	Computes complement of cdf of $\chi^2$ .
<b>cdfchii</b>	Computes $\chi^2$ abscissae values given probability and degrees of freedom.
<b>cdfchinc</b>	Computes integral of noncentral $\chi^2$ .
<b>cdffc</b>	Computes complement of cdf of $F$ .
<b>cdffnc</b>	Computes integral of noncentral $F$ .
<b>cdfgam</b>	Computes integral of incomplete $\Gamma$ function.
<b>cdfmvn</b>	Computes multivariate Normal cdf.
<b>cdfmvnce</b>	Computes the complement of the multivariate Normal cumulative distribution function with error management
<b>cdfmvne</b>	Computes multivariate Normal cumulative distribution function with error management
<b>cdfmvn2e</b>	Computes the multivariate Normal cumulative distribution function with error management over the range [a,b]
<b>cdfmvtce</b>	Computes complement of multivariate Student's t cumulative distribution function with error management
<b>cdfmvte</b>	Computes multivariate Student's t cumulative distribution function with error management
<b>cdfmvt2e</b>	Computes multivariate Student's t cumulative distribution function with error management over [a,b]
<b>cdfn</b>	Computes integral of Normal distribution: lower tail, or cdf.
<b>cdfn2</b>	Computes interval of Normal cdf.
<b>cdfnc</b>	Computes complement of cdf of Normal distribution (upper tail).
<b>cdfni</b>	Computes the inverse of the cdf of the Normal distribution.

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<b>cdftc</b>	Computes complement of cdf of $t$ -distribution.
<b>cdftci</b>	Computes the inverse of the complement of the Student's $t$ cdf.
<b>cdftnc</b>	Computes integral of noncentral $t$ -distribution.
<b>cdftvn</b>	Computes lower tail of trivariate Normal cdf.
<b>erf</b>	Computes Gaussian error function.
<b>erfc</b>	Computes complement of Gaussian error function.
<b>lncdfbvn</b>	Computes natural log of bivariate Normal cdf.
<b>lncdfbvn2</b>	Returns log of cdfbvn of a bounded rectangle.
<b>lncdfmvn</b>	Computes natural log of multivariate Normal cdf.
<b>lncdfn</b>	Computes natural log of Normal cdf.
<b>lncdfn2</b>	Computes natural log of interval of Normal cdf.
<b>lncdfnc</b>	Computes natural log of complement of Normal cdf.
<b>lnpdfmvn</b>	Computes multivariate Normal log-probabilities.
<b>lnpdfmvt</b>	Computes multivariate Student's $t$ log-probabilities.
<b>lnpdfn</b>	Computes Normal log-probabilities.
<b>lnpdft</b>	Computes Student's $t$ log-probabilities.
<b>pdfn</b>	Computes standard Normal probability density function.

### Series and Sequence Functions

<b>recserar</b>	Computes autoregressive recursive series.
<b>recsercp</b>	Computes recursive series involving products.
<b>recserrc</b>	Computes recursive series involving division.
<b>seqa</b>	Creates an additive sequence.
<b>seqm</b>	Creates a multiplicative sequence.

---

## Precision Control

<b>base10</b>	Converts number to $x.xxx$ and a power of 10.
<b>ceil</b>	Rounds up towards $+\infty$ .
<b>floor</b>	Rounds down towards $-\infty$ .
<b>machEpsilon</b>	Returns the smallest number such that $1+eps>1$ .
<b>round</b>	Rounds to the nearest integer.
<b>trunc</b>	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.

## 30.2 Finance Functions

<b>AmericanBinomCall</b>	American binomial method Call.
<b>AmericanBinomCall_Greeks</b>	American binomial method call Delta, Gamma, Theta, Vega, and Rho.
<b>AmericanBinomCall_ImpVol</b>	Implied volatilities for American binomial method calls.
<b>AmericanBinomPut</b>	American binomial method Put.
<b>AmericanBinomPut_Greeks</b>	American binomial method put Delta, Gamma, Theta, Vega, and Rho.
<b>AmericanBinomPut_ImpVol</b>	Implied volatilities for American binomial method puts.
<b>AmericanBSCall</b>	American Black and Scholes Call.

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

<b>EuropeanBSPut_Greeks</b>	European Black and Scholes put Delta, Gamma, Omega, Theta, and Vega.
<b>EuropeanBSPut_ImpVol</b>	Implied volatilities for European Black and Scholes puts.
<b>getNextTradingDay</b>	Returns the next trading day.
<b>getNextWeekDay</b>	Returns the next day that is not on a weekend.
<b>getPreviousTradingDay</b>	Returns the previous trading day.
<b>getPreviousWeekDay</b>	Returns the previous day that is not on a weekend.

## 30.3 Matrix Manipulation

### Creating Vectors and Matrices

<b>eye</b>	Creates identity matrix.
<b>let</b>	Creates matrix from list of constants.
<b>matalloc</b>	Allocates a matrix with unspecified contents.
<b>matinit</b>	Allocates a matrix with specified fill value.
<b>ones</b>	Creates a matrix of ones.
<b>zeros</b>	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 FILE I/O, Chapter 20, for more information.)

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**Loading and Storing Matrices**

<b>asciiload</b>	Loads data from a delimited ASCII text file into an $N \times 1$ vector.
<b>dataload</b>	Loads matrices, N-dimensional arrays, strings and string arrays from a disk file.
<b>datasave</b>	Saves matrices, N-dimensional arrays, strings and string arrays to a disk file.
<b>load,</b> <b>loadm</b>	Loads matrix from ASCII or matrix file.
<b>loadd</b>	Loads matrix from data set.
<b>loadf</b>	Loads function from disk file.
<b>loadk</b>	Loads keyword from disk file.
<b>save</b>	Saves symbol to disk file.
<b>saved</b>	Saves matrix to data set.

**Size, Ranking, and Range**

<b>cols</b>	Returns number of columns in a matrix.
<b>colsf</b>	Returns number of columns in an open data set.
<b>counts</b>	Returns number of elements of a vector falling in specified ranges.
<b>countwts</b>	Returns weighted count of elements of a vector falling in specified ranges.
<b>cumprodc</b>	Computes cumulative products of each column of a matrix.
<b>cumsumc</b>	Computes cumulative sums of each column of a matrix.
<b>indexcat</b>	Returns indices of elements falling within a specified range.
<b>maxc</b>	Returns largest element in each column of a matrix.

<b>maxindc</b>	Returns row number of largest element in each column of a matrix.
<b>minc</b>	Returns smallest element in each column of a matrix.
<b>minindc</b>	Returns row number of smallest element in each column of a matrix.
<b>prodc</b>	Computes the product of each column of a matrix.
<b>rankindx</b>	Returns rank index of N×1 vector. (Rank order of elements in vector).
<b>rows</b>	Returns number of rows in a matrix.
<b>rowsf</b>	Returns number of rows in an open data set.
<b>sumc</b>	Computes the sum of each column of a matrix.
<b>sumr</b>	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.

### Miscellaneous Matrix Manipulation

<b>complex</b>	Creates a complex matrix from two real matrices.
<b>delif</b>	Deletes rows from a matrix using a logical expression.
<b>diag</b>	Extracts the diagonal of a matrix.
<b>diagrv</b>	Puts a column vector into the diagonal of a matrix.
<b>exctsmpl</b>	Creates a random subsample of a data set, with replacement.
<b>imag</b>	Returns the imaginary part of a complex matrix.
<b>indcv</b>	Checks one character vector against another and returns the indices of the elements of the first vector in the second vector.



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<b>indnrv</b>	Checks one numeric vector against another and returns the indices of the elements of the first vector in the second vector.
<b>intersect</b>	Returns the intersection of two vectors.
<b>lowmat</b>	Returns the main diagonal and lower triangle.
<b>lowmat1</b>	Returns a main diagonal of 1's and the lower triangle.
<b>putvals</b>	Inserts values into a matrix or N-dimensional array.
<b>real</b>	Returns the real part of a complex matrix.
<b>reshape</b>	Reshapes a matrix to new dimensions.
<b>rev</b>	Reverses the order of rows of a matrix.
<b>rotater</b>	Rotates the rows of a matrix, wrapping elements as necessary.
<b>selif</b>	Selects rows from a matrix using a logical expression.
<b>setdif</b>	Returns elements of one vector that are not in another.
<b>shiftr</b>	Shifts rows of a matrix, filling in holes with a specified value.
<b>submat</b>	Extracts a submatrix from a matrix.
<b>subvec</b>	Extracts an $N \times 1$ vector of elements from an $N \times K$ matrix.
<b>trimr</b>	Trims rows from top or bottom of a matrix.
<b>union</b>	Returns the union of two vectors.
<b>upmat</b>	Returns the main diagonal and upper triangle.
<b>upmat1</b>	Returns a main diagonal of 1's and the upper triangle.
<b>vec</b>	Stacks columns of a matrix to form a single column.
<b>vech</b>	Reshapes the lower triangular portion of a symmetric matrix into a column vector.
<b>vecr</b>	Stacks rows of a matrix to form a single column.
<b>vget</b>	Extracts a matrix or string from a data buffer constructed with <b>vput</b> .

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<b>vlist</b>	Lists the contents of a data buffer constructed with <b>vput</b> .
<b>vnamecv</b>	Returns the names of the elements of a data buffer constructed with <b>vput</b> .
<b>vput</b>	Inserts a matrix or string into a data buffer.
<b>vread</b>	Reads a string or matrix from a data buffer constructed with <b>vput</b> .
<b>vtypecv</b>	Returns the types of the elements of a data buffer constructed with <b>vput</b> .
<b>xpnd</b>	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**.

## 30.4 Sparse Matrix Handling

<b>denseToSp</b>	Converts a dense matrix to a sparse matrix.
<b>denseToSpRE</b>	Converts a dense matrix to a sparse matrix using a relative epsilon.
<b>packedToSp</b>	Creates a sparse matrix from a packed matrix of non-zero values and row and column indices.
<b>spCreate</b>	Creates a sparse matrix from vectors of non-zero values, row indices, and column indices.
<b>spDenseSubmat</b>	Returns a dense submatrix of a sparse matrix.

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<b>spDiagRvMat</b>	Inserts submatrices along the diagonal of a sparse matrix.
<b>spEye</b>	Creates a sparse identity matrix.
<b>spGetNZE</b>	Returns the non-zero values in a sparse matrix, as well as their corresponding row and column indices.
<b>spGetNumNZE</b>	Returns the number of non-zero elements in a sparse matrix.
<b>spOnes</b>	Generates a sparse matrix containing only ones and zeros
<b>spSubmat</b>	Returns a sparse submatrix of sparse matrix.
<b>spToDense</b>	Converts a sparse matrix to a dense matrix.
<b>spTrTDense</b>	Multiplies a sparse matrix transposed by a dense matrix.
<b>spTScalar</b>	Multiplies a sparse matrix by a scalar.
<b>spZeros</b>	Creates a sparse matrix containing no non-zero values.

## 30.5 N-Dimensional Array Handling

### Creating Arrays

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

### Size, Ranking and Range

<b>amax</b>	Moves across one dimension of an N-dimensional array and finds the largest element.
<b>amin</b>	Moves across one dimension of an N-dimensional array and finds the smallest element.
<b>asum</b>	Computes the sum across one dimension of an N-dimensional array.
<b>getdims</b>	Gets the number of dimensions in an array.
<b>getorders</b>	Gets the vector of orders corresponding to an array.

### Setting and Retrieving Data in an Array

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

---

**Miscellaneous Array Functions**

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

**30.6 Structures**

<b>dsCreate</b>	Creates an instance of a structure of type <b>DS</b> set to default values.
<b>loadstruct</b>	Loads a structure into memory from a file on the disk.
<b>pvCreate</b>	Returns an initialized an instance of structure of type <b>PV</b> .
<b>pvGetIndex</b>	Gets row indices of a matrix in a parameter vector.
<b>pvGetParNames</b>	Generates names for parameter vector stored in structure of type <b>PV</b> .
<b>pvGetParVector</b>	Retrieves parameter vector from structure of type <b>PV</b> .
<b>pvLength</b>	Returns the length of a parameter vector.

<b>pvList</b>	Retrieves names of packed matrices in structure of type <b>PV</b> .
<b>pvPack</b>	Packs general matrix into a structure of type <b>PV</b> with matrix name.
<b>pvPacki</b>	Packs general matrix or array into a <b>PV</b> instance with name and index.
<b>pvPackm</b>	Packs general matrix into a structure of type <b>PV</b> with a mask and matrix name.
<b>pvPackmi</b>	Packs general matrix or array into a <b>PV</b> instance with a mask, name, and index.
<b>pvPacks</b>	Packs symmetric matrix into a structure of type <b>PV</b> .
<b>pvPacksi</b>	Packs symmetric matrix into a <b>PV</b> instance with matrix name and index.
<b>pvPacksm</b>	Packs symmetric matrix into a structure of type <b>PV</b> with a mask.
<b>pvPacksmi</b>	Packs symmetric matrix into a <b>PV</b> instance with a mask, matrix name, and index.
<b>pvPutParVector</b>	Inserts parameter vector into structure of type <b>PV</b> .
<b>pvTest</b>	Tests an instance of structure of type <b>PV</b> to determine if it is a proper structure of type <b>PV</b> .
<b>pvUnpack</b>	Unpacks matrices stored in a structure of type <b>PV</b> .
<b>savestruct</b>	Saves a matrix of structures to a file on the disk.

## 30.7 Data Handling (I/O)

### Spreadsheets

<b>SpreadsheetReadM</b>	Reads and writes Excel files.
<b>SpreadsheetReadSA</b>	Reads and writes Excel files.

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<b>SpreadsheetWrite</b>	Reads and writes Excel files.
<b>xlsGetSheetCount</b>	Gets the number of sheets in an Excel spreadsheet.
<b>xlsGetSheetSize</b>	Gets the size (rows and columns) of a specified sheet in an Excel spreadsheet.
<b>xlsGetSheetTypes</b>	Gets the cell format types of a row in an Excel spreadsheet.
<b>xlsMakeRange</b>	Builds an Excel range string from a row/column pair.
<b>xlsreadm</b>	Reads from an Excel spreadsheet, into a <b>GAUSS</b> matrix.
<b>xlsreadsa</b>	Reads from an Excel spreadsheet, into a <b>GAUSS</b> string array or string.
<b>xlsWrite</b>	Writes a <b>GAUSS</b> matrix, string, or string array to an Excel spreadsheet.
<b>xlswritem</b>	Writes a <b>GAUSS</b> matrix to an Excel spreadsheet.
<b>xlswritesa</b>	Writes a <b>GAUSS</b> string or string array to an Excel spreadsheet.

## Text Files

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

<b>fopen</b>	Opens a file.
<b>fputs</b>	Writes strings to a file.
<b>fputst</b>	Writes strings to a file.
<b>fseek</b>	Positions the file pointer in a file.
<b>fstrerror</b>	Returns an error message explaining the cause of the most recent file I/O error.
<b>ftell</b>	Gets the position of the file pointer in a file.

### GAUSS Data Archives

<b>gdaAppend</b>	Appends data to a variable in a GDA.
<b>gdaCreate</b>	Creates a GDA.
<b>gdaDStat</b>	Computes descriptive statistics on multiple N×1 variables in a GDA.
<b>gdaDStatMat</b>	Computes descriptive statistics on a selection of columns in a variable in a GDA.
<b>gdaGetIndex</b>	Gets the index of a variable in a GDA.
<b>gdaGetName</b>	Gets the name of a variable in a GDA.
<b>gdaGetNames</b>	Gets the names of all the variables in a GDA.
<b>gdaGetOrders</b>	Gets the orders of a variable in a GDA.
<b>gdaGetType</b>	Gets the type of a variable in a GDA.
<b>gdaGetTypes</b>	Gets the types of all the variables in a GDA.
<b>gdaGetVarInfo</b>	Gets information about all of the variables in a GDA.
<b>gdaIsCplx</b>	Checks to see if a variable in a GDA is complex.
<b>gdaLoad</b>	Loads variables in a GDA into the workspace.



<b>gdaPack</b>	Packs the data in a GDA, removing all empty bytes
<b>gdaRead</b>	Gets a variable from a GDA.
<b>gdaReadByIndex</b>	Gets a variable from a GDA, given a variable index.
<b>gdaReadSome</b>	Reads part of a variable from a GDA.
<b>gdaReadSparse</b>	Gets a sparse matrix from a <b>GAUSS</b> Data Archive.
<b>gdaReadStruct</b>	Gets a structure from a <b>GAUSS</b> Data Archive.
<b>gdaReportVarInfo</b>	Gets information about all of the variables in a <b>GAUSS</b> Data Archive and returns it in a string array formatted for printing.
<b>gdaSave</b>	Writes variables in a workspace to a GDA.
<b>gdaUpdate</b>	Updates a variable in a GDA.
<b>gdaUpdateAndPack</b>	Updates a variable in a GDA, leaving no empty bytes if the updated variable is smaller or larger than the variable it is replacing.
<b>gdaWrite</b>	Writes a variable to a GDA.
<b>gdaWrite32</b>	Writes a variable to a GDA using 32-bit system file write commands.
<b>gdaWriteSome</b>	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 [20.3](#).

## Data Sets

<b>close</b>	Closes an open data set (.dat file).
<b>closeall</b>	Closes all open data sets.

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

These functions all operate on **GAUSS** data sets (**.dat** files). For more information, see FILE I/O, Chapter 20.

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 26).

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 **File I/O**, Chapter 20.) See also **create** and **vartypef**.

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

### Data Set Variable Names

<b>getname</b>	Returns column vector of variable names in a data set.
<b>getnamef</b>	Returns string array of variable names in a data set.
<b>indices</b>	Retrieves column numbers and names from a data set.
<b>indices2</b>	Similar to <b>indices</b> , but matches columns with names for dependent and independent variables.
<b>indicesf</b>	Retrieves column numbers and names from a data set.
<b>indicesfn</b>	Retrieves column numbers and names from a data set.
<b>makevars</b>	Decomposes matrix to create column vectors.
<b>setvars</b>	Creates globals using the names in a data set.
<b>vartypef</b>	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

<b>code</b>	Codes the data in a vector by applying a logical set of rules to assign each data value to a category.
<b>code (dataloop)</b>	Creates new variables with different values based on a set of logical expressions.
<b>dataloop (dataloop)</b>	Specifies the beginning of a data loop.
<b>delete (dataloop)</b>	Removes specific rows in a data loop based on a logical expression.
<b>drop (dataloop)</b>	Specifies columns to be dropped from the output data set in a data loop.
<b>dummy</b>	Creates a dummy matrix, expanding values in vector to rows with ones in columns corresponding to true categories and zeros elsewhere.
<b>dummybr</b>	Similar to <b>dummy</b> .
<b>dummydn</b>	Similar to <b>dummy</b> .
<b>extern (dataloop)</b>	Allows access to matrices or strings in memory from inside a data loop.
<b>isinfnanmiss</b>	Returns true if the argument contains an infinity, NaN, or missing value.
<b>issmiss</b>	Returns 1 if matrix has any missing values, 0 otherwise.
<b>keep (dataloop)</b>	Specifies columns (variables) to be saved to the output data set in a data loop.
<b>lag (dataloop)</b>	Lags variables a specified number of periods.
<b>lag1</b>	Lags a matrix by one time period for time series analysis.
<b>lagn</b>	Lags a matrix a specified number of time periods for time series analysis.

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<b>listwise (dataloop)</b>	Controls listwise deletion of missing values.
<b>make (dataloop)</b>	Specifies the creation of a new variable within a data loop.
<b>miss</b>	Changes specified values to missing value code.
<b>missex</b>	Changes elements to missing value using logical expression.
<b>missrv</b>	Changes missing value codes to specified values.
<b>msym</b>	Sets symbol to be interpreted as missing value.
<b>outtyp (dataloop)</b>	Specifies the precision of the output data set.
<b>packr</b>	Delete rows with missing values.
<b>recode</b>	Similar to <b>code</b> , but leaves the original data in place if no condition is met.
<b>recode (dataloop)</b>	Changes the value of a variable with different values based on a set of logical expressions.
<b>scalinfnanmiss</b>	Returns true if the argument is a scalar infinity, NaN, or missing value.
<b>scalmiss</b>	Tests whether a scalar is the missing value code.
<b>select (dataloop)</b>	Selects specific rows (observations) in a data loop based on a logical expression.
<b>subscat</b>	Simpler version of <b>recode</b> , but uses ascending bins instead of logical conditions.
<b>substute</b>	Similar to <b>recode</b> , but operates on matrices.
<b>vector (dataloop)</b>	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

<b>intrleav</b>	Produces one large sorted data file from two smaller sorted files having the same keys.
<b>intrleavsa</b>	Interleaves the rows of two string arrays that have been sorted on a common column.
<b>mergeby</b>	Produces one large sorted data file from two smaller sorted files having a single key column in common.
<b>mergevar</b>	Accepts a list of names of global matrices, and concatenates the corresponding matrices horizontally to form a single matrix.
<b>sortc</b>	Quick-sorts rows of matrix based on numeric key.
<b>sortcc</b>	Quick-sorts rows of matrix based on character key.
<b>sortd</b>	Sorts data set on a key column.
<b>sorthc</b>	Heap-sorts rows of matrix based on numeric key.
<b>sorthcc</b>	Heap-sorts rows of matrix based on character key.
<b>sortind</b>	Returns a sorted index of a numeric vector.
<b>sortindc</b>	Returns a sorted index of a character vector.
<b>sortmc</b>	Sorts rows of matrix on the basis of multiple columns.
<b>sortr</b>	Sorts rows of a matrix of numeric data.
<b>sortrc</b>	Sorts rows of a matrix of character data.
<b>uniqindx</b>	Returns a sorted unique index of a vector.

---

<b>uniqindxsa</b>	Computes the sorted index of a string vector, omitting duplicate elements.
<b>unique</b>	Removes duplicate elements of a vector.
<b>uniquesa</b>	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.

## 30.8 Compiler Control

<b>#define</b>	Defines a case-insensitive text-replacement or flag variable.
<b>#definecs</b>	Defines a case-sensitive text-replacement or flag variable.
<b>#else</b>	Alternates clause for <b>#if-#else-#endif</b> code block.
<b>#endif</b>	End of <b>#if-#else-#endif</b> code block.
<b>#ifdef</b>	Compiles code block if a variable has been <b>#define</b> 'd.
<b>#iflight</b>	Compiles code block if running <b>GAUSS Light</b> .
<b>#ifndef</b>	Compiles code block if a variable has not been <b>#define</b> 'd.
<b>#ifos2win</b>	Compiles code block if running Windows.
<b>#ifunix</b>	Compiles code block if running UNIX.
<b>#include</b>	Includes code from another file in program.
<b>#linesoff</b>	Compiles program without line number and file name records.

---

<b>#lineson</b>	Compiles program with line number and file name records.
<b>#srcfile</b>	Inserts source file name record at this point (currently used when doing data loop translation).
<b>#srcline</b>	Inserts source file line number record at this point (currently used when doing data loop translation).
<b>#undef</b>	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**, Section [10.4](#), for more information.)

## 30.9 Program Control

### Execution Control

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



## Branching

<b>goto</b>	Unconditional branching.
<b>if...endif</b>	Conditional branching.
<b>pop</b>	Retrieves <b>goto</b> 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;
```

## Looping

<b>break</b>	Jumps out the bottom of a <b>do</b> or <b>for</b> loop.
<b>continue</b>	Jumps to the top of a <b>do</b> or <b>for</b> loop.
<b>do while...endo</b>	Executes a series of statements in a loop as long as a given expression is TRUE (or FALSE).
<b>do until...endo</b>	Loops if FALSE.
<b>for...endfor</b>	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;
    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;
```

### Subroutines

<b>gosub</b>	Branches to subroutine.
<b>pop</b>	Retrieves <b>gosub</b> arguments.
<b>return</b>	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.

## Procedures, Keywords, and Functions

<b>endp</b>	Terminates a procedure definition.
<b>fn</b>	Allows user to create one-line functions.
<b>keyword</b>	Begins the definition of a keyword procedure. Keywords are user-defined functions with local or global variables.
<b>local</b>	Declares variables local to a procedure.
<b>proc</b>	Begins definition of multi-line procedure.
<b>retp</b>	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 12, and **LIBRARIES**, Chapter 18, for details.

### Libraries

<b>declare</b>	Initializes variables at compile time.
<b>external</b>	External symbol definitions.
<b>lib</b>	Builds or updates a <b>GAUSS</b> library.
<b>library</b>	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.

### Compiling

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

### Miscellaneous Program Control

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

## 30.10 OS Functions and File Management

<b>cdir</b>	Returns current directory.
<b>ChangeDir</b>	Changes directory in program.
<b>chdir</b>	Changes directory interactively.
<b>DeleteFile</b>	Deletes files.
<b>dlibrary</b>	Dynamically links and unlinks shared libraries.
<b>dllcall</b>	Calls functions located in dynamic libraries.
<b>dos</b>	Provides access to the operating system from within <b>GAUSS</b> .
<b>envget</b>	Gets an environment string.
<b>exec</b>	Executes an executable program file.

<b>execbg</b>	Provides access to the operating system from within <b>GAUSS</b> .
<b>fileinfo</b>	Takes a file specification, returns names and information of files that match.
<b>filesa</b>	Takes a file specification, returns names of files that match.
<b>getpath</b>	Returns an expanded filename including the drive and path.
<b>searchsourcepath</b>	Searches the source path and (if specified) the <b>src</b> subdirectory of the <b>GAUSS</b> installation directory for a specified file.
<b>shell</b>	Shells to OS.

### 30.11 Workspace Management

<b>clear</b>	Sets matrices equal to 0.
<b>clearg</b>	Sets global symbols to 0.
<b>delete</b>	Deletes specified global symbols.
<b>hasimag</b>	Examines matrix for nonzero imaginary part.
<b>iscplx</b>	Returns whether a matrix is real or complex.
<b>maxbytes</b>	Returns maximum memory to be used.
<b>maxvec</b>	Returns maximum allowed vector size.
<b>new</b>	Clears current workspace.
<b>show</b>	Displays global symbol table.
<b>type</b>	Returns type of argument (matrix or string).

<b>typecv</b>	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.

### 30.12 Error Handling and Debugging

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

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

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

### 30.13 String Handling

<b>chrs</b>	Converts ASCII values to a string.
<b>convertsatostr</b>	Converts a 1×1 string array to a string.

<b>convertstrtosa</b>	Converts a string to a 1×1 string array.
<b>cvtos</b>	Converts a character vector to a string.
<b>ftocv</b>	Converts an N×K matrix to a character matrix.
<b>ftos</b>	Converts a floating point scalar to string.
<b>ftostrC</b>	Converts a matrix to a string array using a C language format specification.
<b>getf</b>	Loads ASCII or binary file into string.
<b>indsav</b>	Checks one string array against another and returns
<b>intrsectsa</b>	Returns the intersection of two string vectors, with duplicates removed. the indices of the first string array in the second string array.
<b>loads</b>	Loads a string file (.fst file).
<b>lower</b>	Converts a string to lowercase.
<b>parse</b>	Parses a string, returning a character vector of tokens.
<b>putf</b>	Writes a string to disk file.
<b>stocv</b>	Converts a string to a character vector.
<b>stof</b>	Converts a string to floating point numbers.
<b>strcombine</b>	Converts an N×M string array to an N×1 string vector by combining each element in a column separated by a user-defined delimiter string.
<b>strindx</b>	Finds starting location of one string in another string.
<b>strlen</b>	Returns length of a string.
<b>strput</b>	Lays a substring over a string.
<b>strrindx</b>	Finds starting location of one string in another string, searching from the end to the start of the string.



---

<b>strsect</b>	Extracts a substring of a string.
<b>strsplit</b>	Splits an $N \times 1$ string vector into an $N \times K$ string array of the individual tokens.
<b>strsplitPad</b>	Splits an $N \times 1$ string vector into an $N \times K$ string array of the individual tokens. Pads on the right with null strings.
<b>strtof</b>	Converts a string array to a numeric matrix.
<b>strtofcp1x</b>	Converts a string array to a complex numeric matrix.
<b>strtriml</b>	Strips all whitespace characters from the left side of each element in a string array.
<b>strtrimr</b>	Strips all whitespace characters from the right side of each element in a string array.
<b>strtrunc</b>	Truncates all elements of a string array to not longer than the specified number of characters.
<b>strtrunc1</b>	Truncates the left side of all elements of a string array by a user-specified number of characters.
<b>strtruncpad</b>	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.
<b>strtruncr</b>	Truncates the right side of all elements of a string array by a user-specified number of characters.
<b>token</b>	Extracts the leading token from a string.
<b>upper</b>	Changes a string to uppercase.
<b>vals</b>	Converts a string to ASCII values.
<b>varget</b>	Accesses the global variable named by a string.
<b>varget1</b>	Accesses the local variable named by a string.
<b>varput</b>	Assigns a global variable named by a string.

---

**varputl** 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**.

## 30.14 Time and Date Functions

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

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

## 30.15 Console I/O

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

## 30.16 Output Functions

### Text Output

<b>cls</b>	Clears the window.
------------	--------------------

---

<b>comlog</b>	Controls interactive command logging.
<b>csrcol</b>	Gets column position of cursor on window.
<b>csrlin</b>	Gets row position of cursor on window.
<b>ed</b>	Accesses an alternate editor.
<b>edit</b>	Edits a file with the <b>GAUSS</b> editor.
<b>format</b>	Defines format of matrix printing.
<b>formatcv</b>	Sets the character data format used by <b>printfmt</b> .
<b>formatnv</b>	Sets the numeric data format used by <b>printfmt</b> .
<b>header</b>	Prints a header for a report.
<b>headermt</b>	Prints a header for a report.
<b>locate</b>	Positions the cursor on the window.
<b>output</b>	Redirects <b>print</b> statements to auxiliary output.
<b>outwidth</b>	Sets line width of auxiliary output.
<b>print</b>	Prints to window.
<b>printdos</b>	Prints a string for special handling by the OS.
<b>printfm</b>	Prints matrices using a different format for each column.
<b>printfmt</b>	Prints character, numeric, or mixed matrix using a default format controlled by the functions <b>formatcv</b> and <b>formatnv</b> .
<b>satostrC</b>	Copies from one string array to another using a C language format specifier string for each element.
<b>screen</b> <b>[[on off]]</b>	Directs/suppresses <b>print</b> statements to window.
<b>tab</b>	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**.

## DOS Compatibility Windows

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

## 30.17 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 **PUBLICATION QUALITY GRAPHICS**, Chapter [24](#).

### Graph Types

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

<b>xy</b>	Graphs X,Y using Cartesian coordinate system.
<b>xyz</b>	Graphs X,Y,Z using 3-D Cartesian coordinate system.

Axes Control and Scaling

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

### Text, Labels, Titles, and Fonts

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

### Main Curve Lines and Symbols

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



---

<b>_plwidth</b>	Controls line thickness for main curves.
<b>_pstype</b>	Controls symbol type for main curves.
<b>_psysiz</b>	Controls symbol size for main curves.
<b>_pzclr</b>	Z level color control for <b>contour</b> and <b>surface</b> .

### Extra Lines and Symbols

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

### Graphic Panel, Page, and Plot Control

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

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

### Output Options

<b>_pscreen</b>	Controls graphics output to window.
<b>_psilent</b>	Controls final beep.
<b>_ptek</b>	Controls creation and name of <code>graphics.tkf</code> file.
<b>_pzoom</b>	Specifies zoom parameters.
<b>graphprt</b>	Generates print, conversion file.
<b>pqgwin</b>	Sets the graphics viewer mode.
<b>setvwrmode</b>	Sets the graphics viewer mode.
<b>tkf2eps</b>	Converts <code>.tkf</code> file to Encapsulated PostScript file.
<b>tkf2ps</b>	Converts <code>.tkf</code> file to PostScript file.

---

**Miscellaneous**

<b>_pbox</b>	Draws a border around graphic panel/window.
<b>_pcrop</b>	Controls cropping of graphics data outside axes area.
<b>_pframe</b>	Draws a frame around 2-D, 3-D plots.
<b>_pmcolor</b>	Controls colors to be used for axes, title, $x$ and $y$ labels, date, box, and background.
<b>graphset</b>	Resets all PQG globals to default values.
<b>rerun</b>	Displays most recently created graph.
<b>view</b>	Sets 3-D observer position in workbox units.
<b>viewxyz</b>	Sets 3-D observer position in plot coordinates.
<b>volume</b>	Sets length, width, and height ratios of 3-D workbox.



# Command Reference 31

## abs

**PURPOSE** Returns the absolute value or complex modulus of  $x$ .

**FORMAT**  $y = \text{abs}(x);$

**INPUT**  $x$   $N \times K$  matrix or sparse matrix or N-dimensional array.

**OUTPUT**  $y$   $N \times K$  matrix or sparse matrix or N-dimensional array containing absolute values of  $x$ .

**EXAMPLE**  $x = \text{rndn}(2,2);$   
 $y = \text{abs}(x);$

$$x = \begin{bmatrix} 0.675243 & 1.053485 \\ -0.190746 & -1.229539 \end{bmatrix}$$

$$y = \begin{matrix} 0.675243 & 1.053485 \\ 0.190746 & 1.229539 \end{matrix}$$

In this example, a 2×2 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**       *y*            N×1 vector, data.  
                 *k*            scalar, maximum number of autocorrelations to compute.  
                 *d*            scalar, order of differencing.

**OUTPUT**      *rk*            K×1 vector, sample autocorrelations.

**EXAMPLE**    *x* = { 20.80,  
          18.58,  
          23.39,  
          20.47,  
          21.78,  
          19.56,  
          19.58,  
          18.91,  
          20.08,  
          21.88 };

*rk* = acf(*x*,4,2);  
print *rk*;

```
-0.74911771
0.48360914
-0.34229330
0.17461180
```

SOURCE    tsutil.src

## aconcat

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

**FORMAT**    `y = aconcat(a,b,dim);`

**INPUT**      *a*            matrix or N-dimensional array.  
               *b*            matrix or K-dimensional array, conformable with *a*.  
               *dim*          scalar, dimension in which to concatenate.

**OUTPUT**    *y*            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**    `a = arrayinit(2|3|4,0);`  
               `b = 3*ones(3,4);`  
               `y = aconcat(a,b,3);`

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

```
a = reshape(seqa(1,1,20),4,5);
```

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

**y** will be a 2×4×5 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.

```
a = arrayinit(2|3|4,0);  
b = seqa(1,1,24);  
b = areshape(b,2|3|4);  
y = aconcat(a,b,5);
```

**y** will be a 2×1×2×3×4 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);
```

**y** will be a 2×3×5 array, such that:

[1,1,1] through [1,3,5] =

```
0 0 0 0 1  
0 0 0 0 2  
0 0 0 0 3
```

[2,1,1] through [2,3,5] =

```
0 0 0 0 4  
0 0 0 0 5  
0 0 0 0 6
```



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**  $a = \text{aeye}(o);$

**INPUT**  $o$  N×1 vector of orders, the sizes of the dimensions of  $a$ .

**OUTPUT**  $a$  N-dimensional array, containing 2-dimensional identity arrays.

**REMARKS** If  $o$  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**  $o = \{ 2, 3, 4 \};$   
 $a = \text{aeye}(o);$

$a$  will be a 2×3×4 array, such that:

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

```

1  0  0  0
0  1  0  0
0  0  1  0

```

## amax

---

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

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

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** *x* N-dimensional array.  
*dim* scalar, number of dimension across which to find the maximum value.

**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 = round(10*rndn(24,1));`  
`x = areshape(x,2|3|4);`  
`dim = 2;`  
`y = amax(x,dim);`

**x** is a 2×3×4 array, such that:

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

```
-14  3   3  -9
-7   21  -4  21
 7   -5  20  -2
```

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

```
10  -12  -9  -4
 1   -6 -10   0
-8   9   8  -6
```

**y** will be a 2×1×4 array, such that:

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

```
7  21  20  21
```

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

```
10  9  8  0
```

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

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

**y** will be a 2×3×1 array, such that:

## amean

---

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

3  
21  
20

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

10  
1  
9

SEE ALSO **amin, maxc**

## amean

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

**FORMAT**  $y = \text{amean}(x, dim);$

**INPUT**  $x$  N-dimensional array.  
 $dim$  scalar, number of dimension to compute the mean across.

**OUTPUT**  $y$  [N-1]-dimensional array.

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

**EXAMPLE**  $x = \text{seqa}(1, 1, 24);$   
 $x = \text{areshape}(x, 2 | 3 | 4);$

---

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

**x** is a 2×3×4 array, such that:

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

1	2	3	4
5	6	7	8
9	10	11	12

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

13	14	15	16
17	18	19	20
21	22	23	24

**y** will be a 1×3×4 array, such that:

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

7	8	9	10
11	12	13	14
15	16	17	18

```
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 2×3×1 array, such that:

## AmericanBinomCall

---

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

2.5  
6.5  
10.5

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

14.5  
18.5  
22.5

SEE ALSO **asum**

## AmericanBinomCall

**PURPOSE** Prices American call options using binomial method.

**FORMAT**  $c = \text{AmericanBinomCall}(S0, K, r, div, tau, sigma, N);$

<b>INPUT</b>	<i>S0</i>	scalar, current price.
	<i>K</i>	M×1 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** *c* M×1 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;

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**

<i>S0</i>	scalar, current price.
<i>K</i>	M×1 vector, strike prices.
<i>r</i>	scalar, risk free rate.

## AmericanBinomCall\_Greeks

---

*div* continuous dividend yield.  
*tau* scalar, elapsed time to exercise in annualized days of trading.  
*sigma* 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*        M×1 vector, delta.  
            *g*        M×1 vector, gamma.  
            *t*        M×1 vector, theta.  
            *v*        M×1 vector, vega.  
            *rh*       M×1 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);`

```
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**  $\sigma = \text{AmericanBinomCall\_ImpVol}(c, S_0, K, r, \text{div}, \tau, N);$

**INPUT**

$c$	M×1 vector, call premiums
$S_0$	scalar, current price.
$K$	M×1 vector, strike prices.
$r$	scalar, risk free rate.
$\text{div}$	continuous dividend yield.
$\tau$	scalar, elapsed time to exercise in annualized days of trading.
$N$	number of time segments.

**OUTPUT**  $\sigma$  M×1 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);

```

## AmericanBinomPut

---

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

```
0.19629517  
0.16991943  
0.12874756
```

SOURCE finprocs.src

## AmericanBinomPut

PURPOSE Prices American put options using binomial method.

FORMAT  $c = \text{AmericanBinomPut}(S0, K, r, div, tau, sigma, N);$

INPUT	$S0$	scalar, current price.
	$K$	$M \times 1$ 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	$c$	$M \times 1$ 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;

```

```

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>	M×1 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** **\_fin\_thetaType** scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.

## AmericanBinomPut\_ImpVol

---

**\_fin\_epsilon** scalar, finite difference stepsize. Default = 1e-8.

OUTPUT    *d*            M×1 vector, delta.  
          *g*            M×1 vector, gamma.  
          *t*            M×1 vector, theta.  
          *v*            M×1 vector, vega.  
          *rh*          M×1 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);  
  
                  -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**  $\sigma = \text{AmericanBinomPut\_ImpVol}(c, S_0, K, r, \text{div}, \tau, N);$

**INPUT**

$c$	M×1 vector, put premiums
$S_0$	scalar, current price.
$K$	M×1 vector, strike prices.
$r$	scalar, risk free rate.
$\text{div}$	continuous dividend yield.
$\tau$	scalar, elapsed time to exercise in annualized days of trading.
$N$	number of time segments.

**OUTPUT**  $\sigma$  M×1 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;
```

## AmericanBSCall

---

```
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    *S0*            scalar, current price.  
          *K*            M×1 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.

OUTPUT    *c*            M×1 vector, call 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* = AmericanBSCall(*S0*,*K*,*r*,0,*tau*,*sigma*);  
            print *c*;

            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** *S0* scalar, current price.  
*K* M×1 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.

**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* M×1 vector, delta.  
*g* M×1 vector, gamma.  
*t* M×1 vector, theta.  
*v* M×1 vector, vega.  
*rh* M×1 vector, rho.

**EXAMPLE** *S0* = 305;

## AmericanBSCall\_ImpVol

---

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

```
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>	M×1 vector, call premiums
	<i>S0</i>	scalar, current price.
	<i>K</i>	M×1 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    *sigma*    M×1 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;

          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>	M×1 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 *c* M×1 vector, put premiums.

EXAMPLE `S0 = 718.46;`

## AmericanBSPut\_Greeks

---

```
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;
```

```
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.

**FORMAT** { *d,g,t,v,rh* } = **AmericanBSPut\_Greeks**(*S0,K,r,div,tau,sigma*);

**INPUT**

<i>S0</i>	scalar, current price.
<i>K</i>	M×1 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**

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

OUTPUT    *d*            M×1 vector, delta.  
            *g*            M×1 vector, gamma.  
            *t*            M×1 vector, theta.  
            *v*            M×1 vector, vega.  
            *rh*          M×1 vector, rho.

```
EXAMPLE    S0 = 305;  
            K = 300;  
            r = .08;  
            sigma = .25;  
            tau = .33;  
            print AmericanBSPut_Greeks (S0,K,r,0,tau,sigma);  
  
                 -0.33296721  
            0.0091658294  
                 -17.556118  
                 77.614237  
                 -40.575963
```

SOURCE    finprocs.src

SEE ALSO    **AmericanBSPut\_Impvol**, **AmericanBSPut**, **AmericanBSCall\_Greeks**,  
              **AmericanBinomPut\_Greeks**

AmericanBSPut\_ImpVol

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

FORMAT    *sigma* = **AmericanBSPut\_ImpVol**(*c*,*S0*,*K*,*r*,*div*,*tau*);

INPUT      *c*            M×1 vector, put premiums

## amin

---

*S0* scalar, current price.  
*K* M×1 vector, strike prices.  
*r* scalar, risk free rate.  
*div* continuous dividend yield.  
*tau* scalar, elapsed time to exercise in annualized days of trading.

OUTPUT *sigma* M×1 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;

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 *x* N-dimensional array.

*dim* scalar, number of dimension across which to find the minimum value.

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 = round(10*randn(24,1));
x = areshape(x,2|3|4);
dim = 2;
y = amin(x,dim);
```

**x** is a 2×3×4 array, such that:

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

```
-14  3  3 -9
-7  21 -4 21
7   -5 20 -2
```

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

```
10 -12 -9 -4
1  -6 -10 0
-8  9  8 -6
```

**y** will be a 2×1×4 array, such that:

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

```
-14 -5 -4 -9
```

## amult

---

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

-8   -12   -10   -6

`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 2×3×1 array, such that:

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

-14  
-7  
-5

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

-12  
-10  
-8

SEE ALSO   **amax, minc**

## 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* 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 `a = areshape(seqa(1,1,12),2|3|2);`  
`b = areshape(seqa(1,1,16),2|2|4);`  
`y = amult(a,b);`

**a** is a 2×3×2 array, such that:

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

```
1  2
3  4
5  6
```

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

```
7  8
9 10
11 12
```

**b** is a 2×2×4 array, such that:

## annualTradingDays

---

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

1	2	3	4
5	6	7	8

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

9	10	11	12
13	14	15	16

**y** will be a 2×3×4 array, such that:

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

11	14	17	20
23	30	37	44
35	46	57	68

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

167	182	197	212
211	230	249	268
255	278	301	324

## annualTradingDays

**PURPOSE**    Compute number of trading days in a given year.

**FORMAT**     $n = \text{annualTradingDays}(a);$



---

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 2006. Holidays are defined in <code>holidays.asc</code> . You may edit that file to modify or add holidays.	
SOURCE	<code>finutils.src</code>	
GLOBALS	<code>_fin_annualTradingDays, _fin_holidays</code>	
SEE ALSO	<code>elapsedTradingDays, getNextTradingDay, getPreviousTradingDay, getNextWeekDay, getPreviousWeekDay</code>	

## arccos

PURPOSE	Computes the inverse cosine.	
FORMAT	$y = \arccos(x);$	
INPUT	$x$	$N \times K$ matrix or $N$ -dimensional array.
OUTPUT	$y$	$N \times K$ 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	$x = \{ -1, -0.5, 0, 0.5, 1 \};$ $y = \arccos(x);$	

## arcsin

---

```
      -1.000000
      -0.500000
x =    0.000000
      0.500000
      1.000000
```

```
      3.141593
      2.094395
y =    1.570796
      1.047198
      0.000000
```

SOURCE trig.src

## arcsin

PURPOSE Computes the inverse sine.

FORMAT  $y = \text{arcsin}(x);$

INPUT  $x$              $N \times K$  matrix or  $N$ -dimensional array.

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  $x = \{ -1, -0.5, 0, 0.5, 1 \};$   
 $y = \text{arcsin}(x);$

```

      -1.000000
      -0.500000
x =    0.000000
      0.500000
      1.000000

```

```

      -1.570796
      -0.523599
y =    0.000000
      0.523599
      1.570796

```

SOURCE    trig.src

## areshape

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

**FORMAT**     $y = \mathbf{areshape}(x, o);$

**INPUT**       $x$             scalar, matrix, or N-dimensional array.  
                $o$             M×1 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**     $x = 3;$   
                $orders = \{ 2, 3, 4 \};$

## arrayalloc

---

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

**y** will be a 2×3×4 array of threes.

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

**y** will be a 2×3×4×5 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 3×2×4 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);**

<b>INPUT</b>	<i>o</i>	N×1 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	<i>y</i>	N-dimensional array.
REMARKS	The contents are unspecified. This function is used to allocate an array that will be written to in sections using <b>setarray</b> .	
EXAMPLE	<pre>orders = { 2,3,4 }; y = arrayalloc(orders, 1);</pre> <p><b>y</b> will be a complex 2×3×4 array with unspecified contents.</p>	
SEE ALSO	<b>arrayinit</b> , <b>setarray</b>	

## arrayindex

PURPOSE	Converts a scalar vector index to a vector of indices for an N-dimensional array.	
FORMAT	<i>i</i> = <b>arrayindex</b> ( <i>si</i> , <i>o</i> );	
INPUT	<i>si</i>	scalar, index into vector or 1-dimensional array.
	<i>o</i>	N×1 vector of orders of an N-dimensional array.
OUTPUT	<i>i</i>	N×1 vector of indices, index of corresponding element in N-dimensional array.
REMARKS	This function and its opposite, <b>singleindex</b> , allow you to easily convert between an N-dimensional index and its corresponding location in a 1-dimensional object of the same size.	
EXAMPLE	<pre>orders = { 2,3,4,5 }; v = rndu(prodc(orders),1); a = areshape(v,orders); vi = 50; ai = arrayindex(vi,orders);</pre>	

## arrayinit

---

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

```
vi = 50
```

```
ai =  
    1  
    3  
    2  
    5
```

```
v[vi] = 0.13220899
```

```
getarray(a,ai) = 0.13220899
```

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

SEE ALSO **singleindex**

## arrayinit

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

**FORMAT**  $y = \text{arrayinit}(o,v);$

<b>INPUT</b>	<i>o</i>	N×1 vector of orders, the sizes of the dimensions of the array.
	<i>v</i>	scalar, value to initialize. If <i>v</i> is complex the result will be complex.

OUTPUT  $y$  N-dimensional array with each element equal to the value of  $v$ .

EXAMPLE `orders = { 2,3,4 };`  
`y = arrayinit(orders, 0);`

$y$  will be a  $2 \times 3 \times 4$  array of zeros.

SEE ALSO **arrayalloc**

## arraytomat

PURPOSE Converts an array to type matrix.

FORMAT  $y = \text{arraytomat}(a);$

INPUT  $a$  N-dimensional array.

OUTPUT  $y$   $K \times L$  or  $1 \times L$  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 `a = arrayinit(3|4,2);`  
`y = arraytomat(a);`

$$y = \begin{matrix} & 2 & 2 & 2 & 2 \\ 2 & 2 & 2 & 2 \\ & 2 & 2 & 2 & 2 \end{matrix}$$

SEE ALSO **mattoarray**

### asciiload

**PURPOSE** Loads data from a delimited ASCII text file into an  $N \times 1$  vector.

**FORMAT**  $y = \text{asciiload}(\text{filename});$

**INPUT** *filename* string, name of data file.

**OUTPUT**  $y$   $N \times 1$  vector.

**REMARKS** 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  $N \times 1$  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  $N \times 1$  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 the command:

```
y = asciiload("myfile.asc");
```



---

```

          2.805
          16.568
          -4.871
y =      3.399
          17.361
          -12.725

```

SEE ALSO **load, dataload**

## asclabel

**PURPOSE** To set up character labels for the X and Y axes.

**LIBRARY** pgraph

**FORMAT** **asclabel**(*xl*,*yl*);

**INPUT** *xl* string or N×1 character vector, labels for the tick marks on the X axis. Set to 0 if no character labels for this axis are desired.

*yl* string or M×1 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:

```
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);
```

## astd

---

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  $\lambda$ . The font Simgrma must be previously loaded in a **fonts** command.

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

Here, the “\2021” produces the “ $\lambda$ ” 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**     $y = \mathbf{astd}(x, dim);$

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

<b>OUTPUT</b>	$y$	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:

$$\text{sqrt}(1/(N-1)*\text{sumc}((x-\text{meanc}(x))^2))$$

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

If **a** =

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

```
-38.357528  2.0560337
-21.331064  32.500431
```

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

```
-6.7540544  16.374989
-15.245137  21.824196
```

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

```
18.636931   35.264181
-21.137995  -33.715808
```

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

```
23.008600  -15.933576
54.852400  -7.8360916
```

then **y** =

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

```
28.229091  21.705020
37.152755  29.944103
```

In this example, 16 standard Normal random variables are generated. They are multiplied by 25 and **areshape**'d into a 4×2×2 array, and the standard deviation is computed across the third dimension of the array.

SEE ALSO **astds**, **stdc**

## **astds**

**PURPOSE** Computes the 'sample' 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.  
 $\text{dim}$  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  $\text{dim}$  will be collapsed to 1.

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 **astd** for the alternate definition.

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

If **a** =

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

```

-38.357528  2.0560337
-21.331064  32.500431

```

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

```

-6.7540544  16.374989
-15.245137  21.824196

```

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

```

18.636931   35.264181
-21.137995  -33.715808

```

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

```

23.008600  -15.933576
54.852400  -7.8360916

```

then **y** =

## asum

---

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

```
24.447110  18.797099
32.175230  25.932354
```

In this example, 24 standard Normal random variables are generated. They are multiplied by 10 and **areshape**'d into a 4×3×2 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** *x* N-dimensional array.  
*dim* 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 = seqa(1,1,24);`  
`x = areshape(x,2|3|4);`  
`y = asum(x,3);`

**x** is a 2×3×4 array, such that:

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

1	2	3	4
5	6	7	8
9	10	11	12

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

13	14	15	16
17	18	19	20
21	22	23	24

**y** will be a 1×3×4 array, such that:

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

14	16	18	20
22	24	26	28
30	32	34	36

**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 2×3×1 array, such that:

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

10
26
42

## atan

---

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

58  
74  
90

SEE ALSO    **amean**

## atan

**PURPOSE**    Returns the arctangent of its argument.

**FORMAT**     $y = \mathbf{atan}(x);$

**INPUT**     $x$             N×K matrix or N-dimensional array.

**OUTPUT**     $y$             N×K 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  $-\frac{\pi}{2}$  to  $+\frac{\pi}{2}$ . To convert radians to degrees, multiply by  $\frac{180}{\pi}$ .

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

**EXAMPLE**     $x = \{ 2, 4, 6, 8 \};$   
                   $z = x/2;$   
                   $y = \mathbf{atan}(z);$



0.785398  
1.107149  
y = 1.249046  
1.325818

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×K matrix or P-dimensional array where the last two dimensions are N×K, the *Y* coordinate.  
  
          *x*            L×M matrix or P-dimensional array where the last two dimensions are L×M, E×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  $\frac{180}{\pi}$ .

**atan2** operates only on the real component of *x*, even if *x* is complex.

EXAMPLE    *x* = 2;  
              *y* = { 2, 4, 6, 8 };  
              *z* = **atan2**(*y,x*);

## atranspose

---

```
          0.785398
          1.107149
z =       1.249046
          1.325818
```

SEE ALSO    **atan, sin, cos, pi, tan, arcsin, arccos**

## atranspose

**PURPOSE**    Transposes an N-dimensional array.

**FORMAT**     $y = \text{atranspose}(x, nd);$

**INPUT**     $x$             N-dimensional array.

$nd$             N×1 vector of dimension indices, the new order of dimensions.

**OUTPUT**     $y$             N-dimensional array, transposed according to  $nd$ .

**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**     $x = \text{seqa}(1, 1, 24);$   
               $x = \text{areshape}(x, 2 | 3 | 4);$   
               $nd = \{ 2, 1, 3 \};$   
               $y = \text{atranspose}(x, nd);$

This example transposes the dimensions of  $\mathbf{x}$  that correspond to the first and second elements of the vector of orders.  $\mathbf{x}$  is a 2×3×4 array, such that:

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

1	2	3	4
5	6	7	8
9	10	11	12

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

13	14	15	16
17	18	19	20
21	22	23	24

**y** will be a 3×2×4 array such that:

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

1	2	3	4
13	14	15	16

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

5	6	7	8
17	18	19	20

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

9	10	11	12
21	22	23	24

```
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 3×4×2 array **y**, such that:

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

```
1  13
2  14
3  15
4  16
```

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

```
5  17
6  18
7  19
8  20
```

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

```
9  21
10 22
11 23
12 24
```

SEE ALSO    **areshape**

## axmargin

**PURPOSE**    Sets absolute margins for the plot axes which control placement and size of plot.

LIBRARY    `pgraph`

FORMAT    `axmargin(l,r,t,b);`

INPUT    *l*            scalar, the left margin in inches.  
           *r*            scalar, the right margin in inches.  
           *t*            scalar, the top margin in inches.  
           *b*            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 9×6.855 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(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`

## balance

---

### balance

PURPOSE    Balances a square matrix.

FORMAT     $\{ b, z \} = \mathbf{balance}(x)$

INPUT     $x$              $K \times K$  matrix or N-dimensional array where the last two dimensions are  $K \times K$ .

OUTPUT    $b$              $K \times K$  matrix or N-dimensional array where the last two dimensions are  $K \times K$ , balanced matrix.

$z$              $K \times K$  matrix or N-dimensional array where the last two dimensions are  $K \times K$ , 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.

**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,

$$b = z^{-1} x z$$

**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);
```

```
      100.0  100.0  37.5
b =   80.0   50.0  15.0
      56.0   32.0   9.0
```

```
      4.0  0.0  0.0
z =   0.0  2.0  0.0
      0.0  0.0  0.5
```

band

PURPOSE    Extracts bands from a symmetric banded matrix.

FORMAT     $a = \text{band}(y,n);$

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    $x = \{ 1 \ 2 \ 0 \ 0,$

## bandchol

---

```
2 8 1 0,  
0 1 5 2,  
0 0 2 3 };
```

```
bx = band(x,1);
```

```
bx =  
0.0000000 1.0000000  
2.0000000 8.0000000  
1.0000000 5.0000000  
2.0000000 3.0000000
```

SEE ALSO **bandchol**, **bandcholsol**, **bandltsol**, **bandrv**, **bandsolpd**

## bandchol

**PURPOSE** Computes the Cholesky decomposition of a positive definite banded matrix.

**FORMAT**  $l = \mathbf{bandchol}(a);$

**INPUT**  $a$   $K \times N$  compact form matrix.

**OUTPUT**  $l$   $K \times N$  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  $A = L \times L'$ .  $a$  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);  
  
      0.0000000  1.0000000  
bx =  2.0000000  8.0000000  
      1.0000000  5.0000000  
      2.0000000  3.0000000
```

```
cx = bandchol(bx);  
  
      0.0000000  1.0000000  
cx =  2.0000000  2.0000000  
      0.5000000  2.1794495  
      0.91766294 1.4689774
```

SEE ALSO **band, bandcholsol, bandltsol, bandrv, bandsolpd**

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**  $x = \text{bandcholsol}(b,l);$

**INPUT**  $b$   $K \times M$  matrix.  
 $l$   $K \times N$  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  $A = L \times L'$ .  $l$  is the compact form of  $L$ ; see **band** for a description of the format of  $l$ .

$b$  can have more than one column. If so,  $Ax = b$  is solved for each column. That is,

$$A * x[:, i] = b[:, i]$$

**EXAMPLE**  $x = \{ \begin{array}{cccc} 1 & 2 & 0 & 0, \\ 2 & 8 & 1 & 0, \\ 0 & 1 & 5 & 2, \\ 0 & 0 & 2 & 3 \end{array} \};$

$bx = \text{band}(x, 1);$

```
bx =
0.0000000  1.0000000
2.0000000  8.0000000
1.0000000  5.0000000
2.0000000  3.0000000
```

$cx = \text{bandchol}(bx);$

```
cx =
0.0000000  1.0000000
2.0000000  2.0000000
0.5000000  2.1794495
0.91766294 1.4689774
```

$xi = \text{bandcholsol}(\text{eye}(4), cx);$

---

```

                2.0731707   -0.53658537   0.14634146  -0.097560976
      xi =      -0.53658537   0.26829268  -0.073170732   0.048780488
                0.14634146  -0.073170732   0.29268293  -0.19512195
                -0.097560976  0.048780488  -0.19512195   0.46341463

```

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**  $x = \text{bandltsol}(b, A);$

**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**  $x = \{ \begin{array}{cccc} 1 & 2 & 0 & 0, \\ 2 & 8 & 1 & 0, \\ 0 & 1 & 5 & 2, \\ 0 & 0 & 2 & 3 \end{array} \};$

## bandrv

---

```
bx = band(x,1);
```

```
      0.0000000  1.0000000
bx =  2.0000000  8.0000000
      1.0000000  5.0000000
      2.0000000  3.0000000
```

```
cx = bandchol(bx);
```

```
      0.0000000  1.0000000
cx =  2.0000000  2.0000000
      0.5000000  2.1794495
      0.91766294 1.4689774
```

```
xci = bandltsol(eye(4),cx);
```

```
      1.0000000  0.00000000  0.00000000  0.00000000
xci = -1.0000000  0.50000000  0.00000000  0.00000000
      0.22941573 -0.11470787  0.45883147  0.00000000
      -0.14331487 0.071657436 -0.28662975 0.68074565
```

SEE ALSO **band**, **bandchol**, **bandcholsol**, **bandrv**, **bandsolpd**

## bandrv

**PURPOSE** Creates a symmetric banded matrix, given its compact form.

**FORMAT**  $y = \text{bandrv}(a);$

INPUT  $a$   $K \times N$  compact form matrix.

OUTPUT  $y$   $K \times K$  symmetric banded matrix.

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 ( $N^{th}$ ) 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  $K \times K$  matrix with  $N-1$  subdiagonals.

EXAMPLE  $x = \{ \begin{array}{l} 1 \ 2 \ 0 \ 0, \\ 2 \ 8 \ 1 \ 0, \\ 0 \ 1 \ 5 \ 2, \\ 0 \ 0 \ 2 \ 3 \end{array} \};$

$bx = \text{band}(x, 1);$

$bx = \begin{array}{cc} 0.0000000 & 1.0000000 \\ 2.0000000 & 8.0000000 \\ 1.0000000 & 5.0000000 \\ 2.0000000 & 3.0000000 \end{array}$

$x = \text{bandrv}(bx);$

$x = \begin{array}{cccc} 1.0000000 & 2.0000000 & 0.00000000 & 0.00000000 \\ 2.0000000 & 8.0000000 & 1.0000000 & 0.00000000 \\ 0.00000000 & 1.0000000 & 5.0000000 & 2.0000000 \\ 0.00000000 & 0.0000000 & 2.0000000 & 3.0000000 \end{array}$

SEE ALSO **band**, **bandchol**, **bandcholsol**, **bandltsol**, **bandsolpd**

## bar

---

### 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**  $b$   $K \times M$  matrix.  
 $A$   $K \times N$  compact form matrix.

**OUTPUT**  $x$   $K \times M$  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

**PURPOSE** Generates a bar graph.

**LIBRARY** pgraph

**FORMAT** **bar**( $val, ht$ );

INPUT	<i>val</i>	N×1 numeric vector, bar labels. If scalar 0, a sequence from 1 to <b>rows</b> ( <i>ht</i> ) will be created.
	<i>ht</i>	N×K numeric vector, bar heights. K overlapping or side-by-side sets of N bars will be graphed. For overlapping bars, the first column should contain the set of bars with the greatest height and the last column should contain the set of bars with the least height. Otherwise the bars which are drawn first may be obscured by the bars drawn last. This is not a problem if the bars are plotted side-by-side.
GLOBAL INPUT	<b>_pbarwid</b>	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. If this value is positive, the bars will overlap. If negative, the bars will be plotted side-by-side. The default is 0.5.
	<b>_pbartyp</b>	K×2 matrix. The first column controls the bar shading: <div><b>0</b> no shading. <b>1</b> dots. <b>2</b> vertical cross-hatch. <b>3</b> diagonal lines with positive slope. <b>4</b> diagonal lines with negative slope. <b>5</b> diagonal cross-hatch. <b>6</b> solid.</div> The second column controls the bar color.
REMARKS	Use <b>scale</b> or <b>ytics</b> 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;	

## base10

---

```
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 to */
                           /* 10 (magenta) */
_pnum = 2;
bar(t,x);                  /* Use t vector to label X axis. */
```

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  $\{ M, P \} = \mathbf{base10}(x);$

INPUT  $x$  scalar, number to break down.

OUTPUT  $M$  scalar, in the range  $-10 < M < 10$ .

$P$  scalar, integer power such that:

$$M * 10^P = x$$

EXAMPLE  $\{ b, e \} = \mathbf{base10}(4500);$

$$b = 4.5000000$$



$e = 3.0000000$

SOURCE    `base10.src`

## begwind

PURPOSE    Initializes global graphic panel variables.

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, setwind, nextwind, getwind**

## besselj

PURPOSE    Computes a Bessel function of the first kind,  $J_n(x)$ .

FORMAT     $y = \text{besselj}(n, x);$

INPUT	$n$	N×K matrix or P-dimensional array where the last two dimensions are N×K, the order of the Bessel function. Nonintegers will be truncated to an integer.
	$x$	L×M matrix or P-dimensional array where the last two dimensions are L×M, E×E 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     $n = \{ 0, 1 \};$   
               $x = \{ 0.1 \ 1.2, \ 2.3 \ 3.4 \};$   
               $y = \text{besselj}(n,x);$

$$y = \begin{array}{cc} 0.99750156 & 0.67113274 \\ 0.53987253 & 0.17922585 \end{array}$$

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       $n$              $N \times K$  matrix or P-dimensional array where the last two dimensions are  $N \times K$ , the order of the Bessel function. Nonintegers will be truncated to an integer.

$x$              $L \times M$  matrix or P-dimensional array where the last two dimensions are  $L \times M$ ,  $E \times E$  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     $n = \{ 0, 1 \};$   
               $x = \{ 0.1 \ 1.2, \ 2.3 \ 3.4 \};$   
               $y = \text{bessely}(n,x);$

---

$$y = \begin{array}{cc} -1.5342387 & 0.22808351 \\ 0.052277316 & 0.40101529 \end{array}$$

SEE ALSO **besselj**, **mbesseli**

box

PURPOSE     Graphs data using the box graph percentile method.

LIBRARY     pgraph

FORMAT     **box**(*grp*,*y*);

INPUT     *grp*        1×M vector. This contains the group numbers corresponding to each column of *y* data. If scalar 0, a sequence from 1 to **cols**(*y*) will be generated automatically for the X axis.

*y*         N×M matrix. Each column represents the set of *y* values for an individual percentiles box symbol.

GLOBAL     **\_pboxctl**        5×1 vector, controls box style, width, and color.

INPUT

- [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 50<sup>th</sup> 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.

## boxcox

---

- 2 Statistical standard with the minimum and maximum calculated according to interquartile range as follows:

$$intqrang = 75^{th} - 25^{th}$$

$$min = 25^{th} - 1.5intqrang$$

$$max = 75^{th} + 1.5intqrang$$

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**

1×M 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**.

**\_pcolor**

1×M vector or scalar for symbol colors. If scalar, all symbols will be one color.

REMARKS If missing values are encountered in the y data, they will be ignored during calculations and will not be plotted.

SOURCE `pbox.src`

## boxcox

PURPOSE Computes the Box-Cox function.

FORMAT `y = boxcox (x,lambda);`

INPUT `x`            `M`×`N` matrix or `P`-dimensional array where the last two dimensions are `M`×`N`.  
  
`lambda`           `K`×`L` matrix or `P`-dimensional array where the last two dimensions are `K`×`L`, `E`×`E` conformable to `x`.

OUTPUT `y`            `max(M,L) × max(N,K)` or `P`-dimensional array where the last two dimensions are `max(M,L) × max(N,K)`.

REMARKS    Allowable range for `x` is:

$$x > 0$$

The **boxcox** function computes

$$\text{boxcox}(x) = \frac{x^\lambda - 1}{\lambda}$$

EXAMPLE    `x = { .2 .3, 1.5 2.5 };`  
              `lambda = { .4, 2 };`  
              `y = boxcox(x,lambda)`

$$y = \begin{matrix} -1.1867361 & -0.95549787 \\ 0.62500000 & 2.62500000 \end{matrix}$$

PURPOSE    Breaks out of a **do** or **for** loop.

## call

---

FORMAT    **break;**

EXAMPLE    `x = rndn(4,4);`  
            `r = 0;`  
            `do while r < rows(x);`  
                `r = r + 1;`  
                `c = 0;`  
                `do while c < cols(x);`  
                    `c = c + 1;`  
                    `if c = \, = r;`  
                        `x[r,c] = 1;`  
                    `elseif c > r;`  
                        `break;    /* terminate inner do loop */`  
                    `else;`  
                        `x[r,c] = 0;`  
                    `endif;`  
                `endo; /* break jumps to the statement after this endo */`  
            `endo;`

```
x =
    1.000  0.326 -2.682 -0.594
    0.000  1.000 -0.879  0.056
    0.000  0.000  1.000 -0.688
    0.000  0.000  0.000  1.000
```

REMARKS    This command works just like in C.

SEE ALSO    **continue, do, for**

## 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 **call chol**(**x**);  
**y** = **detl**;

The above example is the fastest way to compute the determinant of a positive definite matrix. The result of **chol** is discarded and **detl** 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 *x* N×K matrix.  
*a* L×M matrix, E×E conformable with *x*.  
*b* P×Q matrix, E×E conformable with *x* and *a*.

OUTPUT **y** max(N,L,P) by max(K,M,Q) matrix.

## cdfbeta

---

**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**  $x = \{ .1, .2, .3, .4 \};$   
 $a = 0.5;$   
 $b = 0.3;$   
 $y = \text{cdfbeta}(x,a,b);$

$y =$   
0.142285  
0.206629  
0.260575  
0.310875

**SEE ALSO** **cdfchic, cdfffc, cdfn, cdfnc, cdftc, gamma**

**TECHNICAL NOTES** **cdfbeta** has the following approximate accuracy:

	$\max(a,b) \leq 500$	absolute error is approx. $\pm 5\text{e-}13$
500 <	$\max(a,b) \leq 10,000$	absolute error is approx. $\pm 5\text{e-}11$
10,000 <	$\max(a,b) \leq 200,000$	absolute error is approx. $\pm 1\text{e-}9$
200,000 <	$\max(a,b)$	Normal approximations are used; absolute error is approx. $\pm 2\text{e-}9$

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cdfbvn

PURPOSE	Computes the cumulative distribution function of the standardized bivariate Normal density (lower tail).		
FORMAT	$c = \text{cdfbvn}(h,k,r);$		
INPUT	$h$	$N \times K$ matrix, the upper limits of integration for variable 1.	
	$k$	$L \times M$ matrix, $E \times E$ conformable with $h$ , the upper limits of integration for variable 2.	
	$r$	$P \times Q$ matrix, $E \times E$ conformable with $h$ and $k$ , the correlation coefficients between the two variables.	
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:

$$-\infty < h < +\infty$$

$$-\infty < k < +\infty$$

$$-1 \leq r \leq 1$$

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:

$$h = (ht - ux) ./ sx;$$

$$k = (kt - uy) ./ sy;$$

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  $\pm 5.0\text{e-}9$  for the entire range of arguments.

c

- 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  $y = \text{cdfbvn2}(h, dh, k, dk, r);$

INPUT

$h$	N×1 vector, starting points of integration for variable 1.
$dh$	N×1 vector, increments for variable 1.
$k$	N×1 vector, starting points of integration for variable 2.
$dk$	N×1 vector, increments for variable 2.
$r$	N×1 vector, correlation coefficients between the two variables.

OUTPUT

$y$	N×1 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 N×1 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`

c

## cdfbvn2e

**PURPOSE** Returns the bivariate Normal cumulative distribution function of a bounded rectangle.

**FORMAT**  $\{ y, e \} = \text{cdfbvn2e}(h, dh, k, dk, r);$

**INPUT**

- $h$   $N \times 1$  vector, starting points of integration for variable 1.
- $dh$   $N \times 1$  vector, increments for variable 1.
- $k$   $N \times 1$  vector, starting points of integration for variable 2.
- $dk$   $N \times 1$  vector, increments for variable 2.
- $r$   $N \times 1$  vector, correlation coefficients between the two variables.

**OUTPUT**

- $y$   $N \times 1$  vector, the integral over the rectangle bounded by  $h$ ,  $h + dh$ ,  $k$ , and  $k + dk$  of the standardized bivariate Normal distribution.
- $e$   $N \times 1$  vector, an error estimate.

**REMARKS** Scalar input arguments are okay; they will be expanded to  $N \times 1$  vectors. **cdfbvn2e** computes:

$$\text{cdfbvn}(h+dh, k+dk, r) + \text{cdfbvn}(h, k, r) - \text{cdfbvn}(h, k+dk, r) - \text{cdfbvn}(h+dh, k, r)$$

The real answer is  $y \pm e$ . The size of the error depends on the input arguments.

**EXAMPLE** Example 1

## **cdfchic**

---

```
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**

## **cdfchic**

**PURPOSE** Computes the complement of the cdf of the chi-square distribution.

**FORMAT**  $y = \text{cdfchic}(x, n)$

<b>INPUT</b>	$x$	$N \times K$ 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  $\infty$  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:

$$x \geq 0$$

$$n > 0$$

A -1 is returned for those elements with invalid inputs.

This equals  $1 - \chi_n^2(x)$ . 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  $x = \{ .1, .2, .3, .4 \};$   
 $n = 3;$   
 $y = \text{cdfchic}(x,n);$

$y =$   
0.991837  
0.977589  
0.960028  
0.940242

SEE ALSO **cdfbeta**, **cdfffc**, **cdfn**, **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:  $1 - \text{cdfgam}(0.5*x, 0.5*n);$

- 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**

<i>p</i>	M×N matrix, probabilities.
<i>n</i>	L×K matrix, E×E conformable with <i>p</i> , degrees of freedom.

**OUTPUT**

<i>c</i>	max(M,L) by max(N,K) matrix, abscissae values for chi-squared distribution.
----------	---

**EXAMPLE** The following generates a 3×3 matrix of pseudo-random numbers with a chi-squared distribution with expected value of 4:

```
rndseed 464578;  
x = cdfchii(rndu(3,3),4+zeros(3,3));
```

```
          2.1096456  1.9354989  1.7549182  
x =  4.4971008  9.2643386  4.3639694  
      4.5737473  1.3706243  2.5653688
```

**SOURCE** `cdfchii.src`



SEE ALSO `gammaii`

c

cdfchinc

**PURPOSE** Computes the cumulative distribution function for the noncentral chi-square distribution.

**FORMAT** `y = cdfchinc(x,v,d);`

**INPUT**

<code>x</code>	N×1 vector, values of upper limits of integrals, must be greater than 0.
<code>v</code>	scalar, degrees of freedom, $v > 0$ .
<code>d</code>	scalar, noncentrality parameter, $d > 0$ . This is the square root of the noncentrality parameter that sometimes goes under the symbol lambda. (See Scheffe, <i>The Analysis of Variance</i> , App. IV, 1959.)

**OUTPUT** `y` N×1 vector.

**REMARKS** `y` is the integral from 0 to `x` of the noncentral chi-square distribution with `v` 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`.

- Relation to `cdfchic`:  
`cdfchic(x,v) = 1 - cdfchinc(x,v,0);`
- The formula used is taken from Abramowitz and Stegun, *Handbook of Mathematical Functions*, 1970, 942, formula 26.4.25.

**EXAMPLE** `x = { .5, 1, 5, 25 };  
print cdfchinc(x,4,2);`

```
0.0042086234
0.016608592
0.30954232
0.99441140
```

SOURCE `cdfnonc.src`

SEE ALSO **`cdffnc`**, **`cdftnc`**

## **cdffc**

**PURPOSE** Computes the complement of the cumulative distribution function of the  $F$  distribution.

**FORMAT** `y = cdffc(x,n1,n2);`

**INPUT** `x`             $N \times K$  matrix.  
`n1`             $L \times M$  matrix,  $E \times E$  conformable with `x`.  
`n2`             $P \times Q$  matrix,  $E \times E$  conformable with `x` and `n1`.

**OUTPUT** `y`             $\max(N,L,P)$  by  $\max(K,M,Q)$  matrix

**REMARKS** `y` is the integral from `x` to  $\infty$  of the  $F$  distribution with `n1` and `n2` degrees of freedom.

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, subtract **`cdffc(x,n1,n2)`** 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.

Allowable ranges for the arguments are:

$$\begin{aligned}x &\geq 0 \\n1 &> 0 \\n2 &> 0\end{aligned}$$

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 **cdfbeta**( $n2/(n2+n1*x)$ ,  $n2/2$ ,  $n1/2$ );

EXAMPLE    `x = { .1, .2, .3, .4 };`  
               `n1 = 0.5;`  
               `n2 = 0.3;`  
               `print cdffc(x,n1,n2);`

```
0.751772
0.708152
0.680365
0.659816
```

SEE ALSO    **cdfbeta**, **cdfchic**, **cdfn**, **cdfnc**, **cdftc**, **gamma**

- REFERENCES
1. Bol'shev, L.N. "Asymptotically Perason's Transformations." *Teor. Veroyat. Primen. 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**      $y = \text{cdffnc}(x, n1, n2, d);$

**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 sometimes goes under the symbol lambda. (See Scheffe, <i>The Analysis of Variance</i> , App. IV, 1959.)

**OUTPUT**      $y$              $N \times 1$  vector.

**TECHNICAL NOTES**

- Relation to cdffc:  
 $\text{cdffnc}(x, n1, n2) = 1 - \text{cdffnc}(x, n1, n2, 0);$
- The formula used is taken from Abramowitz and Stegun, *Handbook of Mathematical Functions*, 1970, 947, formula 26.6.20.

**SOURCE**     `cdfnonc.src`

SEE ALSO **cdftnc**, **cdfchinc**

c

cdfgam

PURPOSE Computes the incomplete gamma function.

FORMAT  $g = \text{cdfgam}(x, \text{intlim});$

INPUT  $x$   $N \times K$  matrix of data.  
 $\text{intlim}$   $L \times M$  matrix,  $E \times E$  compatible with  $x$ , containing the integration limit.

OUTPUT  $g$   $\max(N, L)$  by  $\max(K, M)$  matrix.

REMARKS The incomplete gamma function returns the integral

$$\int_0^{\text{intlim}} \frac{e^{-t} t^{(x-1)}}{\text{gamma}(x)} dt$$

The allowable ranges for the arguments are:

$$\begin{aligned} x &> 0 \\ \text{intlim} &\geq 0 \end{aligned}$$

A -1 is returned for those elements with invalid inputs.

EXAMPLE  $x = \{ 0.5 \quad 1 \quad 3 \quad 10 \};$

```
intlim = seqa(0,.2,6);
g = cdfgam(x,intlim);
```

```
x = 0.500000 1.00000 3.00000 10.0000
```

```
intlim =
0.000000
0.200000
0.400000
0.600000
0.800000
1.000000
```

```
g =
0.000000 0.000000 0.000000 0.000000
0.472911 0.181269 0.00114848 2.35307E - 014
0.628907 0.329680 0.00792633 2.00981E - 011
0.726678 0.451188 0.0231153 9.66972E - 010
0.794097 0.550671 0.0474226 1.43310E - 008
0.842701 0.632120 0.0803014 1.11425E - 007
```

This computes the integrals over the range from 0 to 1, in increments of .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. $\pm 6e-13$
$500 \leq x \leq 10,000$	the absolute error is approx. $\pm 3e-11$
$10,000 < x$	a Normal approximation is used and the absolute error is approx. $\pm 3e-10$

## 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.

## cdfmvn

**PURPOSE** Computes multivariate Normal cumulative distribution function.

**FORMAT**  $y = \text{cdfmvn}(x, r);$

**INPUT**  $x$   $K \times L$  matrix, abscissae.  
 $r$   $K \times K$  matrix, correlation matrix.

**OUTPUT**  $y$   $L \times 1$  vector,  $Pr(X < x|r)$ .

**SEE ALSO** **cdfbvn**, **cdfn**, **lncdfmvn**

**SOURCE** `lncdfn.src`

## cdfmvnce

**PURPOSE** Computes the complement of the multivariate Normal cumulative distribution function with error management

**FORMAT**  $\{y, err, retcode\} = \text{cdfmvnce}(ctl, x, r, m);$

**INPUT**  $ctl$  instance of a **cdfmControl** structure with members  
 $ctl.maxEvaluations$  scalar, maximum number of evaluations  
 $ctl.absErrorTolerance$  scalar absolute error tolerance

		<code>ctl.relative</code>	error tolerance
	$x$	$N \times K$ matrix, abscissae.	
	$r$	$K \times K$ matrix, correlation matrix.	
	$m$	$K \times 1$ vector, means	
OUTPUT	$y$	$L \times 1$ vector, $Pr(X > x r, m)$ .	
	$err$	$L \times 1$ vector, estimates of absolute error	
	$retcode$	$L \times 1$ vector, return codes,	
		<b>0</b> normal completion with $err < ctl.absErrorTolerance$ .	
		<b>1</b> $err > ctl.absErrorTolerance$ and $ctl.maxEvaluations$ exceeded; increase $ctl.maxEvaluations$ to decrease error.	
		<b>2</b> $K > 100$ or $K < 1$ .	
		<b>3</b> $R$ not positive semi-definite.	
		<b>missing</b> $R$ not properly defined.	

REMARKS **cdfmvne** evaluates the following integral

$$\Phi(x_i, R, m) = \frac{1}{\sqrt{|R|(2\pi)^m}} \int_{x_{i1}}^{\infty} \int_{x_{i2}}^{\infty} \dots \int_{x_{iK}}^{\infty} e^{-\frac{1}{2}(z-m')'R^{-1}(z-m')} dz$$

SEE ALSO **cdfmvn2e**, **cdfmvnce**, **cdfmvte**

SOURCE **cdfm.src**

- 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.  
Genz, A., “Numerical computation of multivariate normal probabilities”, *Journal of Computational and Graphical Statistics*, 1:141-149, 1992.



**PURPOSE** Computes multivariate Normal cumulative distribution function with error management

**FORMAT**  $\{y, err, retcode\} = \text{cdfmvne}(ctl, x, r, m);$

**INPUT** *ctl* instance of a **cdfmControl** structure with members  
           *ctl.maxEvaluations* scalar, maximum number of evaluations  
           *ctl.absErrorTolerance* scalar absolute error tolerance  
           *ctl.relative* error tolerance

*x*  $N \times K$  matrix, abscissae.

*r*  $K \times K$  matrix, correlation matrix.

*m*  $K \times 1$  vector, means

**OUTPUT** *y*  $L \times 1$  vector,  $Pr(X < x|r, m)$ .

*err*  $L \times 1$  vector, estimates of absolute error

*retcode*  $L \times 1$  vector, return codes,

**0** normal completion with  $err < ctl.absErrorTolerance$ .

**1**  $err > ctl.absErrorTolerance$  and  $ctl.maxEvaluations$  exceeded;  
increase  $ctl.maxEvaluations$  to decrease error.

**2**  $K > 100$  or  $K < 1$ .

**3**  $R$  not positive semi-definite.

**missing**  $R$  not properly defined.

**REMARKS** **cdfmvne** evaluates the following integral

$$\Phi(x_i, R, m) = \frac{1}{\sqrt{|R|(2\pi)^m}} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \dots \int_{-\infty}^{x_{iK}} e^{-\frac{1}{2}(z-m')'R^{-1}(z-m')} dz$$

**SEE ALSO** **cdfmvne**, **cdfmvn2e**, **cdfmvtce**

## cdfmvn2e

---

SOURCE    `cdfm.src`

- 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.
- 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    *ctl*            instance of a **cdfmControl** structure with members

*ctl.maxEvaluations*    scalar, maximum number of evaluations

*ctl.absErrorTolerance*    scalar absolute error tolerance

*ctl.relative*            error tolerance

*a*             $N \times K$  matrix, lower limits.

*b*             $N \times K$  matrix, upper limits.

*r*             $K \times K$  matrix, correlation matrix.

*m*             $K \times 1$  vector, means

OUTPUT    *y*             $L \times 1$  vector,  $Pr(X > a \text{ and } X < b | r, m)$ .

*err*             $L \times 1$  vector, estimates of absolute error

*retcode*     $L \times 1$  vector, return codes,

**0**    normal completion with  $err < ctl.absErrorTolerance$ .

**1**     $err > ctl.absErrorTolerance$  and *ctl.maxEvaluations* exceeded;  
                    increase *ctl.maxEvaluations* to decrease error.

**2**     $K > 100$  or  $K < 1$ .

**3**  $R$  not positive semi-definite.

**missing**  $R$  not properly defined.

REMARKS **cdfmvne** evaluates the following integral

$$\Phi(a_i, b_i, R, m) = \frac{1}{\sqrt{|R|(2\pi)^m}} \int_{a_{i1}}^{b_{i1}} \int_{a_{i2}}^{b_{i2}} \dots \int_{a_{iK}}^{b_{iK}} e^{-\frac{1}{2}(z-m')'R^{-1}(z-m')} dz$$

SEE ALSO **cdfmvne**, **cdfmvnce**, **cdfmvt2e**

SOURCE **cdfm.src**

- 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.  
Genz, A., “Numerical computation of multivariate normal probabilities”, *Journal of Computational and Graphical Statistics*, 1:141-149, 1992.

## cdfmvtce

PURPOSE Computes complement of multivariate Student's t cumulative distribution function with error management

FORMAT  $\{y, err, retcode\} = \text{cdfmvtce}(ctl, x, R, m);$

INPUT *ctl* instance of a **cdfmControl** structure with members  
     *ctl.maxEvaluations* scalar, maximum number of evaluations  
     *ctl.absErrorTolerance* scalar absolute error tolerance  
     *ctl.relative error* tolerance  
*x*  $N \times K$  matrix, abscissae.  
*R*  $K \times K$  matrix, correlation matrix.

	$m$	$K \times 1$ vector, noncentralities
	$n$	scalar, degrees of freedom
OUTPUT	$y$	$L \times 1$ vector, $Pr(X > x r, m)$ .
	$err$	$L \times 1$ vector, estimates of absolute error
	$retcode$	$L \times 1$ vector, return codes, <b>0</b> normal completion with $err < ctl.absErrorTolerance$ . <b>1</b> $err > ctl.absErrorTolerance$ and $ctl.maxEvaluations$ exceeded; increase $ctl.maxEvaluations$ to decrease error. <b>2</b> $K > 100$ or $K < 1$ . <b>3</b> $R$ not positive semi-definite. <b>missing</b> $R$ not properly defined.

REMARKS The central multivariate Student's t cdf for the i-th row of  $x$  is defined by

$$\begin{aligned}
 T(x_i, R, n) &= \frac{\Gamma(\frac{n+K}{2})}{\Gamma(\frac{n}{2}) \sqrt{|R|} (n\pi)^K} \int_{x_{i1}}^{\infty} \int_{x_{i2}}^{\infty} \cdots \int_{x_{iK}}^{\infty} \left(1 + \frac{z' \Sigma^{-1} z}{n}\right)^{-\frac{n+K}{2}} dz \\
 &\equiv \frac{2^{1-\frac{n}{2}}}{\Gamma(\frac{n}{2})} \int_0^{\infty} s^{n-1} e^{-\frac{s^2}{2}} \Phi\left(-\infty, \frac{sx_i}{\sqrt{n}}, R\right) ds
 \end{aligned}$$

where

$$\Phi(x_i, R) = \frac{1}{\sqrt{|R|} (2\pi)^m} \int_{x_{i1}}^{\infty} \int_{x_{i2}}^{\infty} \cdots \int_{x_{iK}}^{\infty} e^{-\frac{1}{2} z' R^{-1} z} dz$$

For the noncentral cdf we have

$$T(x_i, R, n, m) = \frac{2^{1-\frac{n}{2}}}{\Gamma(\frac{n}{2})} \int_0^{\infty} s^{n-1} e^{-\frac{s^2}{2}} \Phi\left(\frac{sx_i}{\sqrt{n}} - m', \infty, R\right) ds$$

SEE ALSO **cdfmvt2e**, **cdfmvtce**, **cdfmvne**

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.  
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**  $\{y, err, retcode\} = \text{cdfmvte}(ctl, x, R, m);$

**INPUT** *ctl* instance of a **cdfmControl** structure with members

- ctl.maxEvaluations* scalar, maximum number of evaluations
- ctl.absErrorTolerance* scalar absolute error tolerance
- ctl.relative error* tolerance

*x*  $N \times K$  matrix, abscissae.

*R*  $K \times K$  matrix, correlation matrix.

*m*  $K \times 1$  vector, noncentralities

*n* scalar, degrees of freedom

**OUTPUT** *y*  $L \times 1$  vector,  $Pr(X < x | r, m)$ .

*err*  $L \times 1$  vector, estimates of absolute error

*retcode*  $L \times 1$  vector, return codes,

**0** normal completion with  $err < ctl.absErrorTolerance$ .

- 1** *err* > *ctl.absErrorTolerance* and *ctl.maxEvaluations* exceeded;  
increase *ctl.maxEvaluations* to decrease error.
- 2**  $K > 100$  or  $K < 1$ .
- 3** *R* not positive semi-definite.
- missing** *R* not properly defined.

**REMARKS** The central multivariate Student's t cdf for the i-th row of *x* is defined by

$$\begin{aligned}
 T(x_i, R, n) &= \frac{\Gamma(\frac{n+K}{2})}{\Gamma(\frac{n}{2}) \sqrt{|R|} (n\pi)^K} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \cdots \int_{-\infty}^{x_{iK}} \left(1 + \frac{z' \Sigma^{-1} z}{n}\right)^{-\frac{n+K}{2}} dz \\
 &\equiv \frac{2^{1-\frac{n}{2}}}{\Gamma(\frac{n}{2})} \int_0^\infty s^{n-1} e^{-\frac{s^2}{2}} \Phi\left(-\infty, \frac{sx_i}{\sqrt{n}}, R\right) ds
 \end{aligned}$$

where

$$\Phi(x_i, R) = \frac{1}{\sqrt{|R|} (2\pi)^m} \int_{-\infty}^{x_{i1}} \int_{-\infty}^{x_{i2}} \cdots \int_{-\infty}^{x_{iK}} e^{-\frac{1}{2} z' R^{-1} z} dz$$

For the noncentral cdf we have

$$T(x_i, R, n, m) = \frac{2^{1-\frac{n}{2}}}{\Gamma(\frac{n}{2})} \int_0^\infty s^{n-1} e^{-\frac{s^2}{2}} \Phi\left(-\infty, \frac{sx_i}{\sqrt{n}} - m', R\right) ds$$

**SEE ALSO** `cdfmvte`, `cdfmvt2e`, `cdfmvnce`

**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.  
 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} = <b>cdfmvt2e</b> (ctl,a,b,R,m);		
INPUT	ctl	instance of a <b>cdfmControl</b> structure with members	
		ctl.maxEvaluations	scalar, maximum number of evaluations
		ctl.absErrorTolerance	scalar absolute error tolerance
		ctl.relative error	tolerance
	a	N × K matrix, lower limits.	
	b	N × K matrix, upper limits.	
	R	K × K matrix, correlation matrix.	
	m	K × 1 vector, noncentralities	
	n	scalar, degrees of freedom	
OUTPUT	y	L × 1 vector, $Pr(X > a \text{ and } X < b r,m)$ .	
	err	L × 1 vector, estimates of absolute error	
	retcode	L × 1 vector, return codes,	
		0	normal completion with $err < \text{ctl.absErrorTolerance}$ .
		1	$err > \text{ctl.absErrorTolerance}$ and $\text{ctl.maxEvaluations}$ exceeded; increase $\text{ctl.maxEvaluations}$ to decrease error.
		2	$K > 100$ or $K < 1$ .
		3	R not positive semi-definite.

**missing**  $R$  not properly defined.

**REMARKS** The central multivariate Student's  $t$  cdf for the  $i$ -th row of  $x$  is defined by

$$\begin{aligned} T(x_i, R, n) &= \frac{\Gamma(\frac{n+K}{2})}{\Gamma(\frac{n}{2}) \sqrt{|R|} (n\pi)^K} \int_{a_{i1}}^{b_{i1}} \int_{a_{i2}}^{b_{i2}} \dots \int_{a_{iK}}^{b_{iK}} \left(1 + \frac{z' \Sigma^{-1} z}{n}\right)^{-\frac{n+K}{2}} dz \\ &\equiv \frac{2^{1-\frac{n}{2}}}{\Gamma(\frac{n}{2})} \int_0^\infty s^{n-1} e^{-\frac{s^2}{2}} \Phi\left(-\infty, \frac{sx_i}{\sqrt{n}}, R\right) ds \end{aligned}$$

where

$$\Phi(x_i, R) = \frac{1}{\sqrt{|R|} (2\pi)^m} \int_{a_{i1}}^{b_{i1}} \int_{a_{i2}}^{b_{i2}} \dots \int_{a_{iK}}^{b_{iK}} e^{-\frac{1}{2} z' R^{-1} z} dz$$

For the noncentral cdf we have

$$T(x_i, R, n, m) = \frac{2^{1-\frac{n}{2}}}{\Gamma(\frac{n}{2})} \int_0^\infty s^{n-1} e^{-\frac{s^2}{2}} \Phi\left(\frac{sa_i}{\sqrt{n}} - m', \frac{sb_i}{\sqrt{n}} - m', R\right) ds$$

**SEE ALSO** `cdfmvtc`, `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.  
Genz, A., "Numerical computation of multivariate normal probabilities", *Journal of Computational and Graphical Statistics*, 1:141-149, 1992.



**PURPOSE** **cdfn** computes the cumulative distribution function (cdf) of the Normal distribution. **cdfnc** computes 1 minus the cdf of the Normal distribution.

**FORMAT**  $n = \text{cdfn}(x);$   
 $nc = \text{cdfnc}(x);$

**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: **cdfn**( $x$ ) + **cdfnc**( $x$ ) = 1. However, many applications expect **cdfn**( $x$ ) to approach 1, but never actually reach it. Because of this, we have capped the return value of **cdfn** at 1 - machine epsilon, or approximately 1 - 1.11e-16. As the relative error of **cdfn** is about  $\pm 5e-15$  for **cdfn**( $x$ ) around 1, this does not invalidate the result. What it does mean is that for **abs**( $x$ ) > (approx.) 8.2924, the identity does not hold true. If you have a need for the uncapped value of **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     $x = \{-2 \ -1 \ 0 \ 1 \ 2\};$   
               $n = \text{cdfn}(x);$   
               $nc = \text{cdfnc}(x);$

$x$	$=$	-2.00000	-1.00000	0.00000	1.00000	2.00000
$n$	$=$	0.02275	0.15866	0.50000	0.84134	0.97725
$nc$	$=$	0.97725	0.84134	0.50000	0.15866	0.02275

SEE ALSO    **erf, erfc, cdfbeta, cdfchic, cdftc, cdffc, gamma**

TECHNICAL    For the integral from  $-\infty$  to  $x$ :  
NOTES

	$x \leq -37$	<b>cdfn</b> underflows and 0.0 is returned
-36	$< x < -10$	<b>cdfn</b> has a relative error of approx. $\pm 5\text{e-}12$
-10	$< x < 0$	<b>cdfn</b> has a relative error of approx. $\pm 1\text{e-}13$
0	$< x$	<b>cdfn</b> has a relative error of approx. $\pm 5\text{e-}15$

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.

## cdfn2

**PURPOSE** Computes interval of Normal cumulative distribution function.

**FORMAT** `y = cdfn2(x,dx);`

**INPUT** `x` M×N matrix, abscissae.  
`dx` K×L matrix, E×E 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.,  $Pr(x \leq X \leq x + dx)$ .

**REMARKS** The relative error is:

$$\begin{array}{llll}
 |x| \leq 1 & \text{and} & dx \leq 1 & \pm 1e - 14 \\
 1 < |x| < 37 & \text{and} & |dx| < 1/|x| & \pm 1e - 13 \\
 \min(x, x + dx) > -37 & \text{and} & y > 1e - 300 & \pm 1e - 11 \text{ or better}
 \end{array}$$

A relative error of  $\pm 1e-14$  implies that the answer is accurate to better than  $\pm 1$  in the 14<sup>th</sup> digit.

**EXAMPLE** `print cdfn2(1,0.5);`  
  
`9.1848052662599017e-02`  
  
`print cdfn2(20,0.5);`

## **cdfni**

---

```
2.7535164718736454e-89
```

```
print cdfn2(20,1e-2);
```

```
5.0038115018684521e-90
```

```
print cdfn2(-5,2);
```

```
1.3496113800582164e-03
```

```
print cdfn2(-5,0.15);
```

```
3.3065580013000255e-07
```

SEE ALSO **lncdfn2**

SOURCE **lncdfn.src**

## **cdfni**

PURPOSE Computes the inverse of the cdf of the Normal distribution.

FORMAT  $x = \mathbf{cdfni}(p);$

INPUT  $p$  N×K real matrix, Normal probability levels,  $0 \leq p \leq 1$ .

OUTPUT  $x$  N×K real matrix, Normal deviates, such that  $\mathbf{cdfn}(x) = p$

REMARKS  $\mathbf{cdfn}(\mathbf{cdfni}(p)) = p$  to within the errors given below:

	$p \leq 4.6\text{e-}308$	-37.5 is returned
$4.6\text{e-}308 < p < 5\text{e-}24$		accurate to $\pm 5$ in 12 <sup>th</sup> digit
$5\text{e-}24 < p < 0.5$		accurate to $\pm 1$ in 13 <sup>th</sup> digit
$0.5 < p < 1 - 2.22045\text{e-}16$		accurate to $\pm 5$ in 15 <sup>th</sup> digit
$p \geq 1 - 2.22045\text{e-}16$		8.12589... is returned

SEE ALSO **cdfn**

cdftc

- PURPOSE** Computes the complement of the cdf of the Student's *t* distribution.
- FORMAT**  $y = \text{cdftc}(x,n);$
- INPUT** *x* N×K matrix.  
*n* L×M matrix, E×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:

$$\begin{aligned} -\infty &< x < +\infty \\ n &> 0 \end{aligned}$$

A -1 is returned for those elements with invalid inputs.

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     $x = \{ .1, .2, .3, .4 \};$   
               $n = 0.5;$   
               $y = \text{cdftc}(x,n);$

                  0.473165  
               $y =$    0.447100  
                  0.422428  
                  0.399555

SEE ALSO    **cdftci**

TECHNICAL    For results greater than 0.5e-30, the absolute error is approx.  $\pm 1\text{e-}14$  and the  
NOTES        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.

**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  $t$  probability levels,  $0 \leq p \leq 1$ .  
                   $n$               $L \times M$  real matrix, degrees of freedom,  $n \geq 1$ ,  $n$  need not be integral.  
                                   $E \times E$  conformable with  $p$ .

**OUTPUT**      $x$               $\max(N, L)$  by  $\max(K, M)$  real matrix, Student's  $t$  deviates, such that  $\text{cdftc}(x, n) = p$ .

**REMARKS**      $\text{cdftc}(\text{cdftci}(p, n)) = p$  to within the errors given below:

0.5e-30      $< p < 0.01$      accurate to  $\pm 1$  in 12th digit  
                  0.01      $< p$              accurate to  $\pm 1e-14$

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**      $x$               $N \times 1$  vector, values of upper limits of integrals.  
                   $v$              scalar, degrees of freedom,  $v > 0$ .

## cdftvn

---

*d* scalar, noncentrality parameter.  
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* N×1 vector, integrals from  $-\infty$  to *x* of noncentral *t*.

REMARKS **cdftc(*x*,*v*) = 1 - cdftnc(*x*,*v*,0).**

The formula used is based on the formula in *SUGI Supplemental Library User's Guide*, SAS Institute, 1983, 232 (which is attributed to Johnson and Kotz, 1970).

The formula used here is a modification of that formula. It has been tested against direct numerical integration, and against simulation experiments in which noncentral *t* random variates were generated and the cdf found directly.

SOURCE cdfnonc.src

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*,*rho21*,*rho31*,*rho32*);**

INPUT *x1* N×1 vector of upper limits of integration for variable 1.  
*x2* N×1 vector of upper limits of integration for variable 2.  
*x3* N×1 vector of upper limits of integration for variable 3.  
*rho21* scalar or N×1 vector of correlation coefficients between the two variables *x1* and *x2*.



	<i>rho31</i>	scalar or N×1 vector of correlation coefficients between the two variables <i>x2</i> and <i>x3</i> .
	<i>rho32</i>	scalar or N×1 vector of correlation coefficients between the two variables <i>x1</i> and <i>x3</i> .
OUTPUT	<i>c</i>	N×1 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:	

$$\begin{array}{lll} -\infty & < x1 & < +\infty \\ -\infty & < x2 & < +\infty \\ -\infty & < x3 & < +\infty \\ -1 & < rho21 & < 1 \\ -1 & < rho31 & < 1 \\ -1 & < rho32 & < 1 \end{array}$$

In addition, *rho12*, *rho23* and *rho31* 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 (*x1*,*x2*,*x3*) must be the same length, N. The second 3 arguments (*rho21*,*rho31*,*rho32*) 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 *x1*,*x2*,*x3*. All inputs must be column vectors.

To find the integral under a general trivariate density, with *x1*, *x2*, and *x3* having nonzero means and any positive standard deviations, transform by subtracting the mean and dividing by the standard deviation. For example:

## **cdir**

---

$x1 = (x1 - \text{mean}(x1)) / \text{std}(x1);$

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.

## **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  $x = \text{cdir}(\emptyset);$   
 $y = \text{cdir}("d:");$

```
print x;
print y;

C:\\gauss{}
D:\
```

ceil

**PURPOSE** Round up toward  $+\infty$ .

**FORMAT**  $y = \text{ceil}(x);$

**INPUT**  $x$   $N \times K$  matrix.

**OUTPUT**  $y$   $N \times K$  matrix.

**REMARKS** This rounds every element in the matrix  $x$  to an integer. The elements are rounded up toward  $+\infty$ .

**EXAMPLE**  $x = 100 * \text{rndn}(2, 2);$   
 $y = \text{ceil}(x);$

$$x = \begin{bmatrix} 77.68 & -14.10 \\ 4.73 & -158.88 \end{bmatrix}$$

$$y = \begin{bmatrix} 78.00 & -14.00 \\ 5.00 & -158.00 \end{bmatrix}$$

**SEE ALSO** **floor**, **trunc**

### ChangeDir

PURPOSE	Changes the working directory.		
FORMAT	$d = \mathbf{ChangeDir}(s);$		
INPUT	$s$	string, directory to change to.	
OUTPUT	$d$	string, new working directory, or null string if change failed.	
SEE ALSO	<b>chdir cdir</b>		

### chdir

PURPOSE	Changes working directory.		
FORMAT	<b>chdir</b> <i>dirstr</i> ;		
INPUT	<i>dirstr</i>	literal or ^string, directory to change to.	
REMARKS	This is for interactive use. Use <b>ChangeDir</b> in a program.  If the directory change fails, <b>chdir</b> prints an error message.  The working directory is listed in the status report on UNIX.		
SEE ALSO	<b>changedir</b> <b>cdir</b>		

**PURPOSE** Compute compute the probability for a chi-bar square statistic from an hypothesis involving parameters under constraints

**FORMAT**  $SLprob = \text{chiBarSquare}(SL, H, a, b, c, d, bounds);$

**INPUT**

$SL$  scalar, chi-bar square statistic

$H$   $K \times K$  matrix, positive covariance matrix

$G$   $K \times K$  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**  $SLprob$  scalar, probability of  $SL$ .

## chol

---

**REMARKS** See Silvapulle and Sen, *Constrained Statistical Inference*, 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

$$\bar{\chi}^2(V, C) = Z'V^{-1}Z - \min_{\theta \in C} (Z - \theta)'V^{-1}(Z - \theta)$$

$C$  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);
```

```
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 `x = moment(rndn(100,4),0);`  
`y = chol(x);`  
`ypy = y'y;`

```

          90.746566  -6.467195  -1.927489  -15.696056
          -6.467195  87.806557   6.319043  -2.435953
    x =      -1.927489   6.319043  101.973276   4.355520
          -15.696056  -2.435953   4.355520   99.042850

```

```

          9.526099  -0.678892  -0.202338  -1.647690
          0.000000   9.345890   0.661433  -0.380334
    y =      0.000000   0.000000  10.074465   0.424211
          0.000000   0.000000   0.000000   9.798130

```

## choldn

```

          90.746566  -6.467195  -1.927489  -15.696056
      ypy =  -6.467195  87.806557   6.319043  -2.435953
          -1.927489   6.319043  101.973276   4.355520
          -15.696056  -2.435953   4.355520   99.042850
```

SEE ALSO **crout**, **solpd**

## choldn

**PURPOSE** Performs a Cholesky downdate of one or more rows on an upper triangular matrix.

**FORMAT**  $r = \text{choldn}(C, x);$

**INPUT**  $C$   $K \times K$  upper triangular matrix.  
 $x$   $N \times K$  matrix, the rows to downdate  $C$  with.

**OUTPUT**  $r$   $K \times K$  upper triangular matrix, the downdated matrix.

**REMARKS**  $C$  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 = choln(C,x);
```

c

```
           18.87055964  15.32294435  8.04947012
r = 0.00000000  9.30682813 -2.12009339
     0.00000000  0.00000000  7.62878355
```

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**  $b$   $N \times K$  matrix.  
 $C$   $N \times N$  matrix.

**OUTPUT**  $x$   $N \times K$  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.,  $A*x[:,i] = b[:,i]$ .

**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**

```
let b[3,1] = 0.03177513 0.41823100 1.70129375;
let C[3,3] = 1.73351215 1.53201723 1.78102499
```

## cholup

---

```
      0      1.09926365 0.63230050
      0      0      0.67015361;
x = cholsol(b,C);
```

```
      -1.94396905
x = -1.52686768
      3.21579513
```

```
      3.00506436 2.65577048 3.08742844
A0 = 2.65577048 3.55545737 3.42362593
      3.08742844 3.42362593 4.02095978
```

SEE ALSO **chol**

## cholup

**PURPOSE** Performs a Cholesky update of one or more rows on an upper triangular matrix.

**FORMAT**  $r = \text{cholup}(C, x);$

**INPUT**  $C$   $K \times K$  upper triangular matrix.  
 $x$   $N \times K$  matrix, the rows to update  $C$  with.

**OUTPUT**  $r$   $K \times K$  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  $C[3,3] = 18.87055964 \quad 15.32294435 \quad 8.04947012$

```

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);

20.16210005  16.50544413  9.86676135
r = 0.00000000  11.16601462  2.97761666
    0.00000000  0.00000000  11.65496052
```

SEE ALSO **choldn**

chrs

- PURPOSE**    Converts a matrix of ASCII values into a string containing the appropriate characters.
- FORMAT**    `y = chrs(x);`
- INPUT**      `x`            `N×K` 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 length strings when formatting printouts, reports, etc.
- EXAMPLE**    `n = 5;`  
              `print chrs(ones(n,1)*42);`  
  
              `*****`

## clear

---

Since the ASCII value of the asterisk character is 42, the program above will print a string of **n** asterisks.

```
y = chrs(67~65~84);  
print y;  
  
CAT
```

SEE ALSO    **vals, ftos, stof**

## clear

**PURPOSE**    Clears space in memory by setting matrices equal to scalar zero.

**FORMAT**    **clear** *x,y*;

**REMARKS**    **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**    clear x;

SEE ALSO    **clearg, new, show, delete**

## clearg

---

PURPOSE	Clears global symbols by setting them equal to scalar zero.
FORMAT	<b>clearg</b> <i>a,b,c</i> ;
OUTPUT	<i>a,b,c</i> scalar global matrices containing 0.
REMARKS	<b>clearg</b> <b>x</b> ; is equivalent to <b>x = 0</b> ;, where <b>x</b> is understood to be a global symbol. <b>clearg</b> 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.
EXAMPLE	<pre>x = 45; clearg x;</pre> <p style="text-align: center;">x = 0.0000000</p>
SEE ALSO	<b>clear, delete, new, show, local</b>

PURPOSE	Closes a <b>GAUSS</b> file.
FORMAT	<i>y</i> = <b>close</b> ( <i>handle</i> );
INPUT	<i>handle</i> scalar, the file handle given to the file when it was opened with the <b>open, create, or fopen</b> command.
OUTPUT	<i>y</i> scalar, 0 if successful, -1 if unsuccessful.
REMARKS	<i>handle</i> 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:

```
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    `open f1 = dat1 for append;  
              y = writer(f1,x);  
              f1 = close(f1);`

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.

EXAMPLE    `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;`

SEE ALSO    **close, open**



PURPOSE Clears the window.

FORMAT **cls;**

PORTABILITY **Windows**

**cls** clears the Command window if you're in Cmnd I/O mode, the Output window if you're in Split I/O mode.

REMARKS This command clears the window and locates the cursor at the upper left hand corner of the window.

SEE ALSO **locate**

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  $y = \text{code}(e, v);$

INPUT  $e$   $N \times K$  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  $e$ . If more than one of these vectors contains a 1 in any given row, the **code** function will terminate with an error message.

$v$   $(K+1) \times 1$  vector containing the values to be assigned to the new variable.

OUTPUT  $y$   $N \times 1$  vector containing the new values.

REMARKS If none of the  $K$  expressions is true, the new variable is assigned the default value, which is given by the last element of  $v$ .

## code

---

```
EXAMPLE  let x1 = 0  /* column vector of original values */
           5
           10
           15
           20;

let v = 1  /* column vector of new values */
       2
       3; /* the last element of v is the "default" */

e1 = (0 .lt x1) .and (x1 .le 5);    /* expression 1 */
e2 = (5 .lt x1) .and (x1 .le 25);  /* expression 2 */

e = e1~e2; /* concatenate e1 & e2 to make a 1,0 mask
           :: with one less column than the number
           :: of new values in v.
           */

y = code(e,v);
```

```

           0
           5
x1[5,1] = 10      (column vector of original values)
           15
           20
```

```
v[3,1] = 1 2 3      (Note: v is a column vector)
```

```

           0 0
           1 0
e[5,2] = 0 1
           0 1
           0 1
```

```

      3
      1
y[5,1] = 2
      2
      2
```

c

For every row in *e*, if a 1 is in the first column, the first element of *v* is used. If a 1 is in the second column, the second element of *v* is used, and so on. If there are only zeros in the row, the last element of *v* is used. This is the default value.

If there is more than one 1 in any row of *e*, the function will terminate with an error message.

SOURCE    `datatran.src`

SEE ALSO    `recode`, `substute`

code (dataloop)

PURPOSE    Creates new variables with different values based on a set of logical expressions.

FORMAT    `code` **[[#]]** **[[ \$]]** *var* **[[default** *defval* **]] with**  
            *val\_1* **for** *expression\_1*,  
            *val\_2* **for** *expression\_2*,  
            .  
            .  
            .  
            *val\_n* **for** *expression\_n*;

INPUT    *var*            literal, the new variable name.

*defval*        scalar, the default value if none of the expressions are TRUE.

*val*            scalar, value to be used if corresponding expression is TRUE.

## cols

---

*expression* 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**    `y = cols(x);`

INPUT     *x*            N×K matrix or sparse matrix.

OUTPUT    *y*            number of columns in *x*.

REMARKS   If *x* is an empty matrix, **rows(*x*)** and **cols(*x*)** both return 0.

EXAMPLE   `x = rndn(100,3);`  
              `y = cols(x);`

`y = 3.000000`

SEE ALSO   **rows, colsf, show**

colsf

PURPOSE   Returns the number of columns in a **GAUSS** data (**.dat**) file or **GAUSS** matrix (**.fmt**) file.

FORMAT    `yf = colsf(fh);`

INPUT     *fh*            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 fp = myfile with x,10,4;`  
              `b = colsf(fp);`

`b = 10.000000`

## combinate

---

SEE ALSO **rowsf, cols, show**

### combinate

**PURPOSE** Computes combinations of  $N$  things taken  $K$  at a time.

**FORMAT**  $y = \text{combinate}(N, K);$

**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**  $n = 4;$   
 $k = 2;$   
 $y = \text{combinate}(n, k);$

$\text{print } y;$

```
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**

**PURPOSE** Writes combinations of  $N$  things taken  $K$  at a time to a **GAUSS** data set.

**FORMAT** `ret = combined(fname, vnames, N, K);`

**INPUT** *fname* string, file name.

*vname*  $1 \times 1$  or  $K \times 1$  string array, names of columns in data set. If  $1 \times 1$  string, names will have column number appended. If null string, names will be X1, X2, ...

*N* scalar.

*K* scalar.

**OUTPUT** *ret* scalar, if data set was successfully written, *ret* = number of rows written to data set. Otherwise, one of the following:

- 0** file already exists.
- 1** data set couldn't be created.
- n** the  $(n-1)^{th}$  write to the data set failed.

**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** `vnames = "Jim"|"Harry"|"Susan"|"Wendy";`

`k = 2;`

`m = combined("couples", vnames, rows(vnames), k);`

`print m;`

`6.0000`

`open f0 = "couples";`

`y = readr(f0, m);`

## comlog

---

```
names = getnamef(f0);  
f0=close(f0);  
  
for i(1,rows(y),1);  
    print names[y[i,.]]';  
endfor;
```

```
Jim    Harry  
Jim    Susan  
Jim    Wendy  
Harry  Susan  
Harry  Wendy  
Susan  Wendy
```

```
print y;
```

```
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 **combinate**, **numCombinations**

## comlog

**PURPOSE** Controls logging of interactive mode commands to a disk file.

**FORMAT** **comlog** [**file=filename**] [**on|off|reset**];

**INPUT** *filename* literal or ^string.



The **file=filename** subcommand selects the file to log interactive mode statements to. This can be any legal file name.  
 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.  
 There is no default file name.

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 Chapter 19.

FORMAT     **compile** *source fname*;

INPUT       *source*       literal or ^string, the name of the file to be compiled.

*fname*      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 *source*. It will have a `.gcg` 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 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 Options menu.

Don't try to include compiled files with **#include**.

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**

C

## complex

**PURPOSE** Converts a pair of real matrices to a complex matrix.

**FORMAT**  $z = \text{complex}(x_r, x_i);$

**INPUT**  $x_r$   $N \times K$  real matrix, the real elements of  $z$ .  
 $x_i$   $N \times K$  real matrix or scalar, the imaginary elements of  $z$ .

**OUTPUT**  $z$   $N \times K$  complex matrix.

**EXAMPLE**  $x = \begin{Bmatrix} 4 & 6, \\ 9 & 8 \end{Bmatrix};$

$y = \begin{Bmatrix} 3 & 5, \\ 1 & 7 \end{Bmatrix};$

$t = \text{complex}(x, y);$

$$t = \begin{array}{cc} 4.0000000 + 3.0000000i & 6.0000000 + 5.0000000i \\ 9.0000000 + 1.0000000i & 8.0000000 + 7.0000000i \end{array}$$

SEE ALSO **imag, real**

**con**

**PURPOSE**     Requests input from the keyboard (console), and returns it in a matrix.

**FORMAT**      $x = \mathbf{con}(r,c);$

**INPUT**      $r$              scalar, row dimension of matrix.

$c$              scalar, column dimension of matrix.

**OUTPUT**      $x$               $r \times c$  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.

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:

<b>u</b>	up one row	<b>U</b>	first row
<b>d</b>	down one row	<b>D</b>	last row
<b>l</b>	left one column	<b>L</b>	first column
<b>r</b>	right one column	<b>R</b>	last column
<b>t</b>	first element		
<b>b</b>	last element		
<b>g #, #</b>	goto element		
<b>g #</b>	goto element of vector		
<b>h</b>	move horizontally, default		
<b>v</b>	move vertically, default		
<b>\</b>	move diagonally, default		
<b>s</b>	show size of matrix		
<b>n</b>	display element as numeric, default		
<b>c</b>	display element as character		
<b>e</b>	exp(1)		
<b>p</b>	pi		
<b>.</b>	missing value		
<b>?</b>	show help screen		
<b>x</b>	exit		

If the desired matrix is  $1 \times N$  or  $N \times 1$ , then **con** will automatically exit after the last element has been entered, allowing you to input the vector quickly.

If the desired matrix is  $N \times K$ , 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);`

## cond

---

? 2

```
-0.148030    0.861562
 1.791516   -0.663392
```

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**  $c = \text{cond}(x);$

**INPUT**  $x$   $N \times K$  matrix.

**OUTPUT**  $c$  scalar, an estimate of the condition number of  $x$ . 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  $10^{300}$ .

**EXAMPLE**  $x = \begin{Bmatrix} 4 & 2 & 6, \\ 8 & 5 & 7, \\ 3 & 8 & 9 \end{Bmatrix};$

$y = \text{cond}(x);$

$y = 9.8436943$

SOURCE    `svd.src`

c

conj

PURPOSE    Returns the complex conjugate of a matrix.

FORMAT    `y = conj(x);`

INPUT      `x`            `N`×`K` matrix.

OUTPUT    `y`            `N`×`K` matrix, the complex conjugate of `x`.

REMARKS    Compare **conj** with the transpose (') operator.

EXAMPLE    `x = { 1+9i 2,`  
              `4+4i 5i,`  
              `7i 8-2i };`  
  
              `y = conj(x);`

$$\mathbf{x} = \begin{matrix} 1.0000000 + 9.0000000i & 2.0000000 \\ 4.0000000 + 4.0000000i & 0.0000000 + 5.0000000i \\ 0.0000000 + 7.0000000i & 8.0000000 - 2.0000000i \end{matrix}$$

$$\mathbf{y} = \begin{matrix} 1.0000000 - 9.0000000i & 2.0000000 \\ 4.0000000 - 4.0000000i & 0.0000000 - 5.0000000i \\ 0.0000000 - 7.0000000i & 8.0000000 + 2.0000000i \end{matrix}$$

## ConScore

---

### cons

PURPOSE	Retrieves a character string from the keyboard.
FORMAT	$x = \text{cons};$
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 is 254 characters. The program will resume execution when the ENTER key is pressed.
EXAMPLE	$x = \text{cons};$  At the cursor enter:  probability  $x = \text{"probability"}$
SEE ALSO	<b>con</b>

### ConScore

PURPOSE	Compute local score statistic and its probability for hypotheses involving parameters under constraints
FORMAT	$\{ SL, Slprob \} = \text{ConScore}(H, G, grad, a, b, c, d, bounds, psi);$



INPUT	$H$	$K \times K$ matrix, Hessian of loglikelihood with respect to 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 $SL$ .
REMARKS		ConScore computes the local score statistic for the hypothesis $H(\theta) = 0$ vs. $H(\theta) \geq 0$ , where $\theta$ 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|-.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;
grad = -out0.gradient; // minus because -logl in log-likelihood

psi = { 7, 8, 9 };

{ sl, slprob } = ConScore(H,G,grad,0,0,0,0,bounds,psi);

```

## continue

---

```
sl = 3.8605086
```

```
slprob = 0.10410000
```

```
SOURCE  hypotest.src
```

## continue

**PURPOSE** Jumps to the top of a **do** or **for** loop.

**FORMAT** **continue;**

**EXAMPLE**

```
x = rndn(4,4);
r = 0;
do while r < rows(x);
    r = r + 1;
    c = 0;
    do while c < cols(x);    /* continue jumps here */
        c = c + 1;
        if c = \,= r;
            continue;
        endif;
        x[r,c] = 0;
    endo;
endo;
```

```

      -1.032195    0.000000    0.000000    0.000000
x =    0.000000   -1.033763    0.000000    0.000000
      0.000000    0.000000    0.061205    0.000000
      0.000000    0.000000    0.000000   -0.225936
```

REMARKS     This command works just as in **C**.

**c**

contour

PURPOSE     Graphs a matrix of contour data.

LIBRARY     pgraph

FORMAT     **contour**(*x,y,z*);

INPUT     *x*            1×K vector, the X axis data. K must be odd.  
            *y*            N×1 vector, the Y axis data. N must be odd.  
            *z*            N×K matrix, the matrix of height data to be plotted.

GLOBAL     **\_plev**            K×1 vector, user-defined contour levels for **contour**.  
INPUT       Default 0.  
            **\_pzclr**            N×1 or N×2 vector. This controls the Z level colors. See  
                                 **surface** for a complete description of how to set this global.

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 **ztics**.

              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 **ztics**.

SOURCE     pcontour.src

SEE ALSO     **surface**

## conv

**PURPOSE** Computes the convolution of two vectors.

**FORMAT**  $c = \text{conv}(b, x, f, l);$

**INPUT**  $b$   $N \times 1$  vector.  
 $x$   $L \times 1$  vector.  
 $f$  scalar, the first convolution to compute.  
 $l$  scalar, the last convolution to compute.

**OUTPUT**  $c$   $Q \times 1$  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.

**EXAMPLE**  $x = \{ 1, 2, 3, 4 \};$   
 $y = \{ 5, 6, 7, 8 \};$   
 $z1 = \text{conv}(x, y, 0, 0);$   
 $z2 = \text{conv}(x, y, 2, 5);$

5  
 16  
 34  
 $z1 =$  60  
 61  
 52  
 32

```

      16
      34
z2 =  60
      61

```

c

SEE ALSO    **polymult**

convertsatostr

PURPOSE    Converts a 1×1 string array to a string.

FORMAT    *str* = **convertsatostr**(*sa*);

INPUT      *sa*            1×1 string array.

OUTPUT    *str*            string, *sa* converted to a string.

SEE ALSO    **convertstrtosa**

convertstrtosa

PURPOSE    Converts a string to a 1×1 string array.

FORMAT    *sa* = **convertstrtosa**(*str*);

INPUT      *str*            string.

OUTPUT    *sa*            1×1 string array, *str* converted to a string array.

EXAMPLE    `str = "This is a string";`

## corrmm, corrvc, corrx

---

```
z = convertstrtos(a(str));
```

SEE ALSO **convertsatostr**

## corrmm, corrvc, corrx

PURPOSE Computes a population correlation matrix.

FORMAT  $cx = \text{corrmm}(m);$   
 $cx = \text{corrvc}(vc);$   
 $cx = \text{corrx}(x);$

INPUT  $m$   $K \times K$  moment ( $x'x$ ) matrix. A constant term **MUST** have been the first variable when the moment matrix was computed.  
 $vc$   $K \times K$  variance-covariance matrix (of data or parameters).  
 $x$   $N \times K$  matrix of data.

OUTPUT  $cx$   $P \times P$  correlation matrix. For **corrmm**,  $P = K-1$ . For **corrvc** and **corrx**,  $P = K$ .

REMARKS Computes population correlation/covariance matrix, that is, it divides by  $N-1$ , rather than  $N$ . For sample correlation/covariance matrix which uses  $N$  rather than  $N-1$ , see **corrms** and **corrxs**.

SOURCE `corr.src`

SEE ALSO **momentd, corrms, corrxs**



## corrms, corrxs

C

PURPOSE	Computes sample correlation matrix.	
FORMAT	$cx = \text{corrms}(m);$ $cx = \text{corrxs}(x);$	
INPUT	$m$	K×K moment ( $x'x$ ) matrix. A constant term <b>MUST</b> have been the first variable when the moment matrix was computed.
	$x$	N×K matrix of data.
OUTPUT	$cx$	P×P correlation matrix. For <b>corrms</b> , P = K-1. For <b>corrxs</b> , P = K.
REMARKS	Computes sample correlation/covariance matrix, that is, it divides the sample size, N, rather than N-1. For population correlation/covariance matrix which uses N-1 rather than N, see <b>corrmm</b> or <b>corrxx</b> .	
SOURCE	corrs.src	
SEE ALSO	momentd, corrmm, corrxx	

## COS

PURPOSE	Returns the cosine of its argument.	
FORMAT	$y = \text{cos}(x);$	
INPUT	$x$	N×K matrix.
OUTPUT	$y$	N×K matrix containing the cosines of the elements of $x$ .

## cosh

---

REMARKS For real matrices,  $x$  should contain angles measured in radians.

To convert degrees to radians, multiply the degrees by  $\frac{\pi}{180}$ .

EXAMPLE  $x = \{ 0, .5, 1, 1.5 \};$   
 $y = \cosh(x);$

$$y = \begin{matrix} 1.00000000 \\ 0.87758256 \\ 0.54030231 \\ 0.07073720 \end{matrix}$$

SEE ALSO **atan, atan2, pi**

## cosh

PURPOSE Computes the hyperbolic cosine.

FORMAT  $y = \cosh(x);$

INPUT  $x$   $N \times K$  matrix.

OUTPUT  $y$   $N \times K$  matrix containing the hyperbolic cosines of the elements of  $x$ .

EXAMPLE  $x = \{ -0.5, -0.25, 0, 0.25, 0.5, 1 \};$   
 $x = x * \pi;$   
 $y = \cosh(x);$

```

      -1.570796
      -0.785398
      0.000000
x =    0.785398
      1.570796
      3.141593

```

c

```

      2.509178
      1.324609
      1.000000
y =    1.324609
      2.509178
      11.591953

```

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×1 vector containing the numbers to be counted.

*v*        P×1 vector containing breakpoints specifying the ranges within  
                 which counts are to be made. The vector *v* MUST be sorted in  
                 ascending order.

OUTPUT    *c*        P×1 vector, the counts of the elements of *x* that fall into the regions:

$x \leq v[1],$   
 $v[1] < x \leq v[2],$   
 $\vdots$

$$v[p - 1] < x \leq v[p]$$

**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

$$x = \begin{matrix} 1 \\ 2 \\ 3 \\ 4 \\ 5 \\ 6 \\ 7 \\ 8 \\ 9 \end{matrix} \quad \text{and} \quad v = \begin{matrix} 4 \\ 5 \\ 8 \end{matrix}$$

then

$$c = \begin{matrix} 4 \\ 1 \\ 3 \end{matrix}$$

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**  $x = \{ 1, 3, 2, 4, 1, 3 \};$   
 $v = \{ 0, 1, 2, 3, 4 \};$

---

```
c = counts(x,v);
```

```

0.0000000
2.0000000
c = 1.0000000
2.0000000
1.0000000

```

c

---

**countwts**

**PURPOSE** Returns a weighted count of the numbers of elements of a vector that fall into specified ranges.

**FORMAT**  $c = \text{countwts}(x, v, w);$

**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**  $c$   $P \times 1$  vector containing the weighted counts of the elements of  $x$  that fall into the regions:

$$\begin{array}{rcl}
 & x & \leq v[1], \\
 v[1] & < x & \leq v[2], \\
 & \dots & \\
 v[p-1] & < x & \leq v[p]
 \end{array}$$

That is, when  $x[i]$  falls into region  $j$ , the weight  $w[i]$  is added to the  $j^{\text{th}}$  counter.

## create

**REMARKS** If any elements of  $x$  are greater than the last element of  $v$ , they will not be counted.

Missing values are not counted unless there is a missing in  $v$ . A missing value in  $v$  **MUST** be the first element in  $v$ .

**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);
```

```
0.000000
0.500000
c = 0.333000
2.000000
0.100000
```

## 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** *vflag* literal, version flag.

- v89** obsoleted, use **-v96**.
- v92** obsoleted, use **-v96**.
- v96** supported on all platforms.

For details on the various versions, see FILE I/O, Chapter 20. The default format can be specified in `gauss.cfg` by setting the

**dat\_fmt\_version** configuration variable. The default, **v96**, should be used.

*filename* literal or ^string  
*filename* 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 **.dat**. If an extension is specified, the **.dat** 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 ^ (caret) operator.

**create... with...**

*vnames* literal or ^string or ^character matrix.  
*vnames* 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 ^ (caret) operator must be placed before the name of the string or character matrix. The number of columns parameter, *col*, 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.

*col* scalar expression.  
*col* is a scalar expression containing the number of columns in the data file. If *col* is 0, the number of columns will be controlled by the contents of *vnames*. If *col* is positive, the file will contain *col* columns and the names to be given each column will be created as necessary depending on the *vnames* parameter. See the examples.

*dtyp* scalar expression.  
*dtyp* 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	$-32768$	$\leq X \leq 32767$
single	6-7	$8.43 \times 10^{-37}$	$\leq  X  \leq 3.37 \times 10^{+38}$
double	15-16	$4.19 \times 10^{-307}$	$\leq  X  \leq 1.67 \times 10^{+308}$

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;  
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 uses 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. 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.

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.

#### EXAMPLE

```
let vnames = age sex educat wage occ;
create f1 = simdat with ^vnames,0,8;
```

## create

---

```
obs = 0;  nr = 1000;
do while obs < 10000;
    data = rndn(nr,colsf(f1));
    if writer(f1,data) /= nr;
        print "Disk Full"; end;
    endif;
    obs = obs+nr;
endo;
closeall f1;
```

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,0,2,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**

**PURPOSE**    Computes the cross-products (vector products) of sets of 3×1 vectors.

**FORMAT**     $z = \text{crossprd}(x,y);$

**INPUT**       $x$             3×K matrix, each column is treated as a 3×1 vector.

$y$             3×K matrix, each column is treated as a 3×1 vector.

**OUTPUT**     $z$             3×K 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 K×1 vectors, all of whose elements are 0 (except for rounding error).

**EXAMPLE**     $x = \{ \begin{array}{l} 10 \ 4, \\ 11 \ 13, \\ 14 \ 13 \end{array} \};$   
               $y = \{ \begin{array}{l} 3 \ 11, \\ 5 \ 12, \\ 7 \ 9 \end{array} \};$   
               $z = \text{crossprd}(x,y);$

                              7.0000000   -39.000000  
 $z = \begin{array}{l} -28.000000 \quad 107.00000 \\ 17.000000 \quad -95.00000 \end{array}$

**SOURCE**    crossprd.src

**PURPOSE** Computes the Crout decomposition of a square matrix without row pivoting, such that:  $X = LU$ .

**FORMAT**  $y = \text{crout}(x);$

**INPUT**  $x$   $N \times N$  square nonsingular matrix.

**OUTPUT**  $y$   $N \times N$  matrix containing the lower ( $L$ ) and upper ( $U$ ) matrices of the Crout decomposition of  $x$ . The main diagonal of  $y$  is the main diagonal of the lower matrix  $L$ . The upper matrix has an implicit main diagonal of 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**  $X = \begin{Bmatrix} 1 & 2 & -1, \\ 2 & 3 & -2, \\ 1 & -2 & 1 \end{Bmatrix};$

```
y = crout(x);
L = lowmat(y);
U = upmat1(y);
```

$$y = \begin{pmatrix} 1 & 2 & -1 \\ 2 & -1 & 0 \\ 1 & -4 & 2 \end{pmatrix}$$

$$L = \begin{pmatrix} 1 & 0 & 0 \\ 2 & -1 & 0 \\ 1 & -4 & 2 \end{pmatrix}$$

## croutp

---

$$U = \begin{bmatrix} 1 & 2 & -1 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

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 = 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 };
```

```
y = croutp(x);
r = rows(y);      /* the number of rows of y */
indx = y[r,.]';   /* get the index vector */
```

---

```

z = y[indx,.];    /* z is indexed RxR matrix y */
L = lowmat(z);    /* obtain L and U of permuted matrix X */
U = upmat1(z);
q = sortc(indx~(L*U),1); /* sort L*U against index */
x2 = q[.,2:cols(q)];    /* remove index column */

```

```

      1   2  -1
X =   2   3  -2
      1  -2   1

```

```

      1   0.5  0.2857
y =   2   1.5    -1
      1  -3.5 -0.5714
      2    3     1

```

```

r = 4

```

```

      2
indx = 3
      1

```

```

      2   1.5    -1
z =   1  -3.5 -0.5714
      1   0.5  0.2857

```

```

      2    0    0
L =   1  -3.5    0
      1   0.5  0.2857

```

```

      1  1.5    -1
U =   0    1 -0.5714
      0    0     1

```

## csrcol, csrlin

---

$$q = \begin{pmatrix} 1 & 1 & 2 & -1 \\ 2 & 2 & 3 & -2 \\ 3 & 1 & -2 & 1 \end{pmatrix}$$

$$x2 = \begin{pmatrix} 1 & 2 & -1 \\ 2 & 3 & -2 \\ 1 & -2 & 1 \end{pmatrix}$$

SEE ALSO **crout, chol, lowmat, lowmat1, lu, upmat, upmat1**

## csrcol, csrlin

**PURPOSE** Returns the position of the cursor.

**FORMAT**  $y = \text{csrcol};$

$y = \text{csrlin};$

**OUTPUT**  $y$  scalar, row or column value.

**PORTABILITY** **Windows** only

**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.

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;  
              cls;  
              locate r,c;`

In this example the screen is cleared without affecting the cursor position.

SEE ALSO    **cls, locate**

## cumprodc

PURPOSE    Computes the cumulative products of the columns of a matrix.

FORMAT    `y = cumprodc(x);`

INPUT      `x`            N×K matrix.

OUTPUT    `y`            N×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);`

## cumsumc

---

$$y = \begin{bmatrix} 1.00 & -3.00 \\ 2.00 & -6.00 \\ 6.00 & 6.00 \end{bmatrix}$$

SOURCE    cumprodc.src

SEE ALSO    **cumsumc**, **recsercp**, **recserar**

## cumsumc

PURPOSE    Computes the cumulative sums of the columns of a matrix.

FORMAT     $y = \text{cumsumc}(x);$

INPUT     $x$              $N \times K$  matrix.

OUTPUT     $y$              $N \times K$  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:

$$\text{recserar}(x, x[1, :], \text{ones}(1, \text{cols}(x)))$$

to accomplish the same thing.

EXAMPLE     $x = \begin{bmatrix} 1 & -3, \\ 2 & 2, \\ 3 & -1 \end{bmatrix};$

$$y = \text{cumsumc}(x);$$

$$y = \begin{pmatrix} 1 & -3 \\ 3 & -1 \\ 6 & -2 \end{pmatrix}$$

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);`

INPUT *x* K×1 vector, x-abscissae (x-axis values).  
*y* K×1 vector, y-ordinates (y-axis values).  
*d* K×1 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.  
*sigma* scalar, tension factor.  
*G* scalar, grid size factor.

OUTPUT *u* (K\*G)×1 vector, x-abscissae, regularly spaced.  
*v* (K\*G)×1 vector, y-ordinates, regularly spaced.

REMARKS *sigma* contains the tension factor. This value indicates the curviness desired. If *sigma* is nearly zero (e.g. .001), the resulting curve is approximately the tensor product of cubic curves. If *sigma* is large, (e.g. 50.0) the resulting curve is approximately bi-linear. If *sigma* equals zero, tensor products of cubic curves result. A standard value for *sigma* is approximately 1.

## cvtos

---

$G$  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       $v$              $N \times 1$  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" };`  
              `s = cvtos(v);`

`s = "Now is the time for all good men"`

SEE ALSO **stocv, vget, vlist, vput, vread**

C

## datacreate

**PURPOSE** Creates a **v96** 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 N×1 string array, names of variables.
<i>col</i>	scalar, number of variables.
<i>dtyp</i>	scalar, data precision, one of the following: <b>2</b> 2-byte, signed integer. <b>4</b> 4-byte, single precision. <b>8</b> 8-byte, double precision.
<i>vtyp</i>	scalar or N×1 vector, types of variables, may contain one or both of the following: <b>0</b> character variable. <b>1</b> numeric variable.

**OUTPUT** *fh* 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 *vnames*1, *vnames*2, ..., *vnames*N (or *vnames*01, *vnames*02, ..., *vnames*N), 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	$-32768 \leq X \leq 32767$
single	6-7	$8.43 \times 10^{-37} \leq  X  \leq 3.37 \times 10^{+38}$
double	15-16	$4.19 \times 10^{-307} \leq  X  \leq 1.67 \times 10^{+308}$

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*×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

d

**PURPOSE** Creates a **v96** complex data set.

**FORMAT** *fh* = **datacreatecomplex**(*filename*, *vnames*, *col*, *dtyp*, *vtyp*);

**INPUT** *filename* string, name of data file.

*vnames* string or N×1 string array, names of variables.

*col* scalar, number of variables.

*dtyp* scalar, data precision, one of the following:

**2** 2-byte, signed integer.

**4** 4-byte, single precision.

**8** 8-byte, double precision.

*vtyp* scalar or N×1 vector, types of variables, may contain one or both of the following:

**0** character variable.

**1** numeric variable.

**OUTPUT** *fh* 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.

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 *vnames*1, *vnames*2, ..., *vnames*N (or *vnames*01, *vnames*02, ..., *vnames*N), 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	$-32768 \leq X \leq 32767$
single	6-7	$8.43 \times 10^{-37} \leq  X  \leq 3.37 \times 10^{+38}$
double	15-16	$4.19 \times 10^{-307} \leq  X  \leq 1.67 \times 10^{+308}$

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  $N \times 1$  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**

d

**datalist**

PURPOSE    List selected variables from a data set.

FORMAT    **datalist** *dataset* `[[var1 [[var2...]]]`;

INPUT    *dataset*    literal, name of the data set.  
           *var#*        literal, the names of the variables to list.

GLOBAL    **\_\_range**        scalar, the range of rows to list. The default is all rows.  
 INPUT    **\_\_miss**            scalar, controls handling of missing values.  
           **0**        display rows with missing values.  
           **1**        do not display rows with missing values.  
           The default is 0.  
           **\_\_prec**        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**

## **dataloop (dataloop)**

---

### **dataload**

**PURPOSE** Loads matrices, N-dimensional arrays, strings and string arrays from a disk file.

**FORMAT** `y = dataload(filename);`

**INPUT** *filename* string, name of data file.

**OUTPUT** *y* 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:

- `.fmt` matrix file
- array file
- `.fst` string file
- string array file

See FILE I/O, Chapter 20, for details on these file types.

**EXAMPLE** `y = dataload("myfile.fmt");`

**SEE ALSO** `load`, `datasave`

### **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	<i>outfile</i>	string variable or literal, the name of the output data set.
REMARKS	The statements between the <b>dataloop... endata</b> commands are assumed to be metacode to be translated at compile time. The data from <i>infile</i> is manipulated by the specified statements, and stored to the data set <i>outfile</i> . Case is not significant within the <b>dataloop... endata</b> section, except for within quoted strings. Comments can be used as in any <b>GAUSS</b> code.	
EXAMPLE	<pre>src = "source"; dataloop ^src dest;     make newvar = x1 + x2 + log(x3);     x6 = sqrt(x4);     keep x6, x5, newvar; endata;</pre> <p>Here, <b>src</b> is a string variable requiring the caret (^) operator, while <b>dest</b> is a string literal.</p>	

d

## dataopen

PURPOSE	Opens a data set.	
FORMAT	<i>fh</i> = <b>dataopen</b> ( <i>filename</i> , <i>mode</i> );	
INPUT	<i>filename</i>	string, name of data file.
	<i>mode</i>	string containing one of the following:
	<b>read</b>	open file for read.
	<b>append</b>	open file for append.
	<b>update</b>	open file for update.
OUTPUT	<i>fh</i>	scalar, file handle.

REMARKS     The file must exist before it can be opened with the **dataopen** command (to create a new file, see **datacreate** or **datasave**).

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.

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.

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 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.

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.

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.

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.

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.

EXAMPLE    `fh = dataopen("myfile.dat","read");`  
              `y = readr(fh,100);`  
              `ret = close(fh);`

This example opens the data file `myfile.dat` in the current directory and reads 100 observations (rows) from the file into the global variable **y**.

SOURCE    `datafile.src`

SEE ALSO    **open, datacreate, writer, readr**

**d**

**datasave**

PURPOSE    Saves matrices, N-dimensional arrays, strings and string arrays to a disk file.

FORMAT    `ret = datasave(filename,x);`

INPUT      *filename*    string, name of data file.  
              *x*                matrix, array, string or string array, data to write to disk.

OUTPUT    *ret*                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

## date

---

See FILE I/O, Chapter 20, for details on these file types.

Use **dataload** to load a data file created with **datsave**.

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;`

```
2005.0000
8.0000000
31.000000
4804392.2
```

SEE ALSO    **time, timestr, ethsec, hsec, etstr**

## datestr

PURPOSE    Returns a date in a string.

FORMAT `str = datestr(d);`

INPUT `d` 4×1 vector, like the **date** function returns. If this is 0, the **date** function will be called for the current system date.

OUTPUT `str` 8 character string containing current date in the form: **mo/dy/yr**

EXAMPLE `d = { 2005, 8, 31, 0 };`  
`y = datestr(d);`  
`print y;`  
  
8/31/05

SOURCE `time.src`

SEE ALSO **date, datestring, datestrynd, time, timestr, ethsec**

## datestring

PURPOSE Returns a date in a string with a 4-digit year.

FORMAT `str = datestring(d);`

INPUT `d` 4×1 vector, like the **date** function returns. If this is 0, the **date** function will be called for the current system date.

OUTPUT `str` 10 character string containing current date in the form: **mm/dd/yyyy**

EXAMPLE `y = datestring(0);`  
`print y;`  
  
8/31/2005

## datestrymd

---

SOURCE    `time.src`

SEE ALSO    `date`, `datestr`, `datestrymd`, `time`, `timestr`, `ethsec`

### datestrymd

PURPOSE    Returns a date in a string.

FORMAT    `str = datestrymd(d);`

INPUT    *d*            4×1 vector, like the **date** function returns. If this is 0, the **date** function will be called for the current system date.

OUTPUT    *str*            8 character string containing current date in the form: **yyyymmdd**

EXAMPLE    `d = { 2005, 8, 31, 0 };`  
             `y = datestrymd(d);`  
             `print y;`

             20050831

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>	3×1 or 4×1 vector, date to check. The date should be in the form returned by <b>date</b> .
OUTPUT	<i>daynum</i>	scalar, the day number of that date in that year.
EXAMPLE	<pre>x = { 2005, 8, 31, 0 }; y = dayinyr(x); print y;  243.000000</pre>	
SOURCE	<code>time.src</code>	
GLOBALS	<code>_isleap</code>	

d

dayofweek

PURPOSE	Returns day of week.	
FORMAT	<i>d</i> = <b>dayofweek</b> ( <i>a</i> );	
INPUT	<i>a</i>	N×1 vector, dates in DT format.
OUTPUT	<i>d</i>	N×1 vector, integers indicating day of week of each date: <div><div>1</div>Sunday</div> <div><div>2</div>Monday</div> <div><div>3</div>Tuesday</div> <div><div>4</div>Wednesday</div> <div><div>5</div>Thursday</div> <div><div>6</div>Friday</div> <div><div>7</div>Saturday</div>

## debug

---

REMARKS    The DT scalar format is a double precision representation of the date and time.  
              In the DT scalar format, the number  
  
              20050901183207  
  
              represents 18:32:07 or 6:32:07 PM on September 1, 2005.

SOURCE    `time.src`

## debug

PURPOSE    Runs a program under the source level debugger.

FORMAT    **debug** *filename*;

INPUT    *filename*    Literal, name of file to debug.

REMARKS    See DEBUGGING, Section 3.3.

## declare

PURPOSE    Initializes global variables at compile time.

FORMAT    **declare** *[[type]] symbol [[aop clist]]*;

INPUT    *type*            optional literal, specifying the type of the symbol.  
              **matrix**  
              **string**  
              **array**  
              **sparse matrix**  
              **struct** *structure\_type*

if *type* is not specified, **matrix** is assumed. Set *type* to **string** to initialize a string or string array variable.

*symbol* the name of the symbol being declared.

*aop* the type of assignment to be made.

= if not initialized, initialize. If already initialized, reinitialize.

!= if not initialized, initialize. If already initialized, reinitialize.

:= if not initialized, initialize. If already initialized, redefinition error.

?= if not initialized, initialize. If already initialized, leave as is.

If *aop* is specified, *clist* must be also.

*clist* a list of constants to assign to *symbol*.

If *aop clist* is not specified, *symbol* is initialized as a scalar 0 or a null string.

REMARKS The **declare** syntax is similar to the **let** statement.

**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:

## declare

---

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:

<b>!=, =</b>	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 <b>declare</b> statement is compiled. The symbol will be reset. This allows mixing <b>declare</b> statements with the procedure definitions that reference the global matrices and strings or placing them in your main file.
<b>:=</b>	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. You need to place <b>declare</b> statements in a separate file from the rest of your program and procedure definitions.
<b>?=</b>	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 <b>declare</b> '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 Windows 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:

d

**declare !=**      Reinitialize and warn.

**declare :=**      Crash 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.";`

`x = "This string."`

`declare matrix x;`

`x = 0`

`declare matrix x != { 1 2 3, 4 5 6, 7 8 9 };`

## declare

---

$$\mathbf{x} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix}$$

```
declare matrix x[3,3] = 1 2 3 4 5 6 7 8 9;
```

$$\mathbf{x} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix}$$

```
declare matrix x[3,3] = 1;
```

$$\mathbf{x} = \begin{bmatrix} 1 & 1 & 1 \\ 1 & 1 & 1 \\ 1 & 1 & 1 \end{bmatrix}$$

```
declare matrix x[3,3];
```

$$\mathbf{x} = \begin{bmatrix} 0 & 0 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix}$$

```
declare matrix x = 1 2 3 4 5 6 7 8 9;
```

---

```

1
2
3
4
x = 5
6
7
8
9

```

**d**

```
declare matrix x = dog cat;
```

```

x = DOG
    CAT

```

```
declare matrix x = "dog" "cat";
```

```

x = dog
    cat

```

```
declare array a;
```

**a** is a 1-dimensional array of 1 containing 0.

```
declare sparse matrix sm;
```

**sm** is an empty sparse matrix.

```

struct mystruct {
    matrix m;
    string s;
}

```

## delete

---

```
        string array sa;  
        array a;  
        sparse matrix sm;  
};  
  
declare struct mystruct ms;
```

**ms** is a **mystruct** structure, with its members set as follows:

```
ms.m    empty matrix  
ms.s    null string  
ms.sa   1×1 string array containing a null string  
ms.a    1-dimensional array of 1 containing 0  
ms.sm   empty sparse matrix
```

SEE ALSO **let, external**

## delete

**PURPOSE** Deletes global symbols from the symbol table.

**FORMAT** **delete** *[-flags]* *[[symbol1]* *[[symbol2]* *[[symbol3]*];

<b>INPUT</b>	<i>flags</i>	specify the type(s) of symbols to be deleted
	<b>p</b>	procedures
	<b>k</b>	keywords
	<b>f</b>	<b>fn</b> functions
	<b>m</b>	matrices
	<b>s</b>	strings
	<b>g</b>	only procedures with global references
	<b>l</b>	only procedures with all local references
	<b>n</b>	no pause for confirmation



*symbol*     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 workspace.

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 deleted, all procedures, keywords and functions with global references to those symbols will be deleted as well.

**EXAMPLE**     `print x;`

```

96.000000
6.0000000
14.000000
3502965.9

```

```
delete -m x;
```

At the Delete?[Yes No Previous Quit] prompt, enter y.

```
show x;
```

**x** no longer exists.

d

## DeleteFile

---

PURPOSE	Removes specific rows in a data loop based on a logical expression.
FORMAT	<b>delete</b> <i>logical expression</i> ;
REMARKS	<p>Deletes only those rows for which <i>logical expression</i> is TRUE. Any variables referenced must already exist, either as elements of the source data set, as <b>extern</b>'s, or as the result of a previous <b>make</b>, <b>vector</b>, or <b>code</b> statement.</p> <p><b>GAUSS</b> expects <i>logical expression</i> to return a row vector of 1's and 0's. The relational and other operators (e.g. &lt;) are already interpreted in terms of their dot equivalents (. &lt;), but it is up to the user to make sure that function calls within <i>logical expression</i> result in a vector.</p>
EXAMPLE	<code>delete age &lt; 40 or sex = \,= 'FEMALE' ;</code>
SEE ALSO	<b>select</b>

## DeleteFile

PURPOSE	Deletes files.
FORMAT	<i>ret</i> = <b>DeleteFile</b> ( <i>name</i> );
INPUT	<i>name</i> string or N×K string array, name of file or files to delete.
OUTPUT	<i>ret</i> scalar or N×K matrix, 0 if successful.
REMARKS	<p>The return value, <i>ret</i>, is scalar if <i>name</i> is a string. If <i>name</i> is an N×K string array, <i>ret</i> will be an N×K matrix reflecting the success or failure of each separate file deletion.</p> <p><b>DeleteFile</b> calls the C library <b>unlink</b> function for each file. If <b>unlink</b> fails it sets the C library errno value. <b>DeleteFile</b> returns the value of errno if <b>unlink</b></p>

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

d

**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** *x*            N×K data matrix.  
*e*                N×1 logical vector (vector of 0's and 1's).

**OUTPUT** *y*            M×K 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.

**REMARKS** The input *e* will usually be generated by a logical expression using dot operators. For instance:

```
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*.

**EXAMPLE** `x = { 0 10 20,  
          30 40 50,  
          60 70 80 };`  
`/* logical vector */`  
`e = (x[.,1] .gt 0) .and (x[.,3] .lt 100);`  
`y = delif(x,e);`

## denseToSp

---

$y = \begin{bmatrix} 0 & 10 & 20 \end{bmatrix}$

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.

SEE ALSO **selif**

## denseToSp

**PURPOSE** Converts a dense matrix to a sparse matrix.

**FORMAT**  $y = \text{denseToSp}(x, \text{eps});$

**INPUT**  $x$   $M \times N$  dense matrix.  
 $\text{eps}$  scalar, elements of  $x$  whose absolute values are less than or equal to  $\text{eps}$  will be treated as zero.

**OUTPUT**  $y$   $M \times N$  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**.

**EXAMPLE** sparse matrix  $y$ ;  
 $x = \begin{bmatrix} 0 & 0 & 0 & 1, \\ 0 & 4 & 0 & 0, \\ 0 & 0 & 0 & 0, \\ 0 & 0 & -2 & 0 \end{bmatrix};$   
  
 $y = \text{denseToSp}(x, 0);$   
 $d = \text{spDenseSubmat}(y, 0, 0);$

$$d = \begin{pmatrix} 0 & 0 & 0 & 1 \\ 0 & 4 & 0 & 0 \\ 0 & 0 & 0 & 0 \\ 0 & 0 & -2 & 0 \end{pmatrix}$$

d

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 *x* M×N dense matrix.  
*reps* scalar, relative epsilon. Elements of *x* will be treated as zero if their absolute values are less than or equal to *reps* multiplied by the mean of the absolute values of the non-zero values in *x*.

OUTPUT *y* M×N 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 0 };`

## denToZero

---

```
y = denseToSpRE(x, .5);  
d = spToDense(y);
```

```
      -9  0  0  0  
d =    0  4  0  0  
      5  0  0  7  
      0  0  0  0
```

SEE ALSO    **denseToSp**, **spCreate**, **spToDense**

## denToZero

**PURPOSE**    Converts every denormal to a 0 in a matrix or array.

**FORMAT**     $y = \text{denToZero}(x);$

**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. column.

**EXAMPLE**     $x = \{ 1, \exp(-724.5), 3 \};$   
               $y = \text{isden}(x);$

$y = 1, 0, 3;$

SEE ALSO    **isden**

design

d

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×1 vector.

OUTPUT      *y*             N×K matrix, where **K = maxc(x)**; each row of *y* will contain a single 1, and the rest 0's. The one in the *i<sup>th</sup>* row will be in the **round(x[i,1])** column.

REMARKS     Note that *x* does not have to contain integers: it will be rounded to nearest if necessary.

EXAMPLE     `x = { 1, 1.2, 2, 3, 4.4 };`  
              `y = design(x);`

                  1 0 0 0  
                  1 0 0 0  
y = 0 1 0 0  
      0 0 1 0  
      0 0 0 1

SOURCE      design.src

SEE ALSO     cumprodc, cumsumc, recserrc

**det**

**PURPOSE** Returns the determinant of a square matrix.

**FORMAT**  $y = \text{det}(x);$

**INPUT**  $x$   $N \times N$  square matrix or K-dimensional array where the last two dimensions are  $N \times N$ .

**OUTPUT**  $y$  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  $10 \times 4 \times 4$  array, the result will be a 1-dimensional array of 10 elements containing the determinants of each of the 10  $4 \times 4$  arrays contained in  $x$ .

**det** computes a LU decomposition.

**det1** can be much faster in many applications.

**EXAMPLE**  $x = \begin{Bmatrix} 3 & 2 & 1, \\ 0 & 1 & -2, \\ 1 & 3 & 4 \end{Bmatrix};$   
 $y = \text{det}(x);$

$$x = \begin{bmatrix} 3 & 2 & 1 \\ 0 & 1 & -2 \\ 1 & 3 & 4 \end{bmatrix}$$



$y = 25$

SEE ALSO **detl**

d

**detl**

**PURPOSE** Returns the determinant of the last matrix that was passed to one of the intrinsic matrix decomposition routines.

**FORMAT**  $y = \mathbf{detl};$

**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**( $y, x$ )

$y/x$

determinant of  $x$

determinant of  $x$  when neither argument is a scalar

or

determinant of  $x'x$  if  $x$  is not square

**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:

$\mathbf{xi} = \mathbf{inv}(x);$

```
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 10×10:

```
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**     `x`            N×1 vector.

**OUTPUT**    `y`            N×1 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**    `dfft.src`

SEE ALSO **dffti**, **fft**, **ffti**

## dfft

d

**PURPOSE** Computes inverse discrete Fourier transform.

**FORMAT** `y = dfft(x);`

**INPUT** `x`  $N \times 1$  vector.

**OUTPUT** `y`  $N \times 1$  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** `dfft.src`

SEE ALSO **fft**, **dffti**, **ffti**

## diag

**PURPOSE** Creates a column vector from the diagonal of a matrix.

**FORMAT** `y = diag(x);`

**INPUT** `x`  $N \times K$  matrix or  $L$ -dimensional array where the last two dimensions are  $N \times K$ .

## diagrv

---

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  $4 \times 4$  arrays contained in  $x$ .

**diagrv** reverses the procedure and puts a vector into the diagonal of a matrix.

EXAMPLE     $x = \text{rndu}(3,3);$   
               $y = \text{diag}(x);$

```
          0.660818  0.367424  0.302208
x = 0.204800  0.077357  0.145755
      0.712284  0.353760  0.642567
```

```
          0.660818
y = 0.077357
      0.642567
```

SEE ALSO    **diagrv**

## diagrv

PURPOSE    Inserts a vector into the diagonal of a matrix.

FORMAT      $y = \text{diagrv}(x,v);$

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 = rndu(3,3);
v = ones(3,1);
y = diagrv(x,v);
```

```
      0.660818  0.367424  0.302208
x =  0.204800  0.077357  0.145755
      0.712284  0.353760  0.642567
```

```
      1.000000
v =  1.000000
      1.000000
```

```
      1.000000  0.367424  0.302208
y =  0.204800  1.000000  0.145755
      0.712284  0.353760  1.000000
```

SEE ALSO **diag**

digamma

PURPOSE Computes the digamma function.

FORMAT  $y = \text{digamma}(x);$

## dlibrary

---

INPUT	$x$	$M \times N$ matrix or $N$ -dimensional array.
OUTPUT	$y$	$M \times N$ 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 *lib1 lib2...* literal, the base name of the library or the pathed name of the library.  
**dlibrary** 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 `.so` or `.dll`, 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.

**-a** 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 **-a** flag, any previously linked libraries are dumped.

**-d** dump flag, ALL shared libraries are unlinked and the functions they contain are no longer available to your programs. If you use **dllcall** to call one of your functions after executing a **dlibrary** **-d**, 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 always active and are not unlinked when you execute **dlibrary -d**.

For more information, see FOREIGN LANGUAGE INTERFACE, Chapter 21.

**SEE ALSO** **dllcall**, **sysstate**—case 24

## **dllcall**

**PURPOSE** Calls functions located in dynamic libraries.

**FORMAT** **dllcall** **[-r] [-v] func[(arg1,arg2...)]**;

d

**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.

INPUT	<i>func</i>	the name of a function contained in a shared library (linked into <b>GAUSS</b> with <b>dlibrary</b> ). If <i>func</i> is not specified or cannot be located in a shared library, <b>dllcall</b> will fail.
	<i>arg#</i>	arguments to be passed to <i>func</i> , optional. These must be simple variable references; they cannot be expressions.
	<b>-r</b>	optional flag. If <b>-r</b> is specified, <b>dllcall</b> examines the value returned by <i>func</i> , and fails if it is nonzero.
	<b>-v</b>	optional flag. Normally, <b>dllcall</b> passes parameters to <i>func</i> in a list. If <b>-v</b> is specified, <b>dllcall</b> 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 [,double \*arg2,... ]);**
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 21.

SEE ALSO **dlibrary**, **sysstate**—case 24

d

## do while, do until

**PURPOSE** Executes a series of statements in a loop as long as a given expression is true (or false).

**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

## do while, do until

---

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;
```

```
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.

d

First loop:

```
format /rd 1,0;  
i = 1;  
do while i <= 10;  
    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
```

## dos

---

SEE ALSO **continue, break**

## dos

**PURPOSE** Provides access to the operating system from within **GAUSS**.

**FORMAT** **dos** `[[s]]`;

**INPUT** `s` literal or ^string, the OS command to be executed.

**PORTABILITY** **UNIX/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.

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 26.

```
> 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 UNIX/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**

## DOSWinCloseall

---

### doswin

PURPOSE    Opens the DOS compatibility window with default settings.

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.

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;

```

d

## DOSWinOpen

**PURPOSE**    Opens the DOS compatibility window and gives it the specified title and attributes.

**FORMAT**    *ret* = **DOSWinOpen**(*title*,*attr*);

**INPUT**    *title*        string, window title.

*attr*        5×1 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 bit)    issue dialog

                  bit 1 (2's bit)    close window

                  bit 2 (4's bit)    stop program

**OUTPUT**    *ret*        scalar, success flag, 1 if successful, 0 if not.

**PORTABILITY**    **Windows** only

**REMARKS**    If *title* is a null string (""), the window will be titled "GAUSS-DOS".

## dotfeq, dotfge, dotfgt, dotfle, dotflt, dotfne

---

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.



dotfeq, dotfge, dotfgt, dotfle, dotflt, dotfne

d

PURPOSE	Fuzzy comparison functions. These functions use <b>_fcmptol</b> to fuzz the comparison operations to allow for roundoff error.		
FORMAT	<pre>y = dotfeq(a,b);  y = dotfge(a,b);  y = dotfgt(a,b);  y = dotfle(a,b);  y = dotflt(a,b);  y = dotfne(a,b);</pre>		
INPUT	<i>a</i>	N×K matrix, first matrix.	
	<i>b</i>	L×M matrix, second matrix, E×E compatible with <i>a</i> .	
GLOBAL INPUT	<b>_fcmptol</b>	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:		
	<pre>y = dotfeq(a,b);</pre>		
	is equivalent to:		
	<pre>y = a .eq b;</pre>		

## dotfeqmt, dotfgemt, dotfgtmt, dotflemt, dotflmt, dotfnemt

---

The calling program can reset **\_fcmptol** before calling these procedures:

```
_fcmptol = 1e-12;
```

EXAMPLE    `x = rndu(2,2);`  
            `y = rndu(2,2);`  
            `t = dotfge(x,y);`

```
      x = 0.85115559  0.98914218  
          0.12703276  0.43365175
```

```
      y = 0.41907226  0.49648058  
          0.58039125  0.98200340
```

```
      t = 1.0000000  1.0000000  
          0.0000000  0.0000000
```

SOURCE    `fcompare.src`

GLOBALS   **\_fcmptol**

SEE ALSO   **feq--fne**

## dotfeqmt, dotfgemt, dotfgtmt, dotflemt, dotflmt, dotfnemt

PURPOSE    Fuzzy comparison functions. These functions use the *fcmptol* argument to fuzz the comparison operations to allow for roundoff error.

FORMAT    `y = dotfeqmt(a,b,fcmptol);`

```
y = dotfgemt(a,b,fcmtol);
```

```
y = dotfgtmt(a,b,fcmtol);
```

```
y = dotflemt(a,b,fcmtol);
```

```
y = dotflmt(a,b,fcmtol);
```

```
y = dotfnemt(a,b,fcmtol);
```

INPUT    *a*            N×K matrix, first matrix.  
           *b*            L×M matrix, second matrix, E×E compatible with *a*.  
           *fcmtol*      scalar, comparison tolerance.

OUTPUT   *y*            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* = rndu(2,2);  
              *t* = dotfge(*x*,*y*,1e-15);

```

x =  0.85115559  0.98914218
     0.12703276  0.43365175

```

## draw

---

```
y = 0.41907226 0.49648058
     0.58039125 0.98200340
```

```
t = 1.0000000 1.0000000
     0.0000000 0.0000000
```

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.

**LIBRARY** `pgraph`

**FORMAT** **`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.

## dsCreate

---

EXAMPLE     drop age, pay, sex;

SEE ALSO     **keep** (**dataloop**)

### dsCreate

PURPOSE     Creates an instance of a structure of type **DS** set to default values.

INCLUDE     ds.sdf

FORMAT     *s* = **dsCreate**;

OUTPUT     *s*             instance of structure of type **DS**.

SOURCE     ds.src

### dstat

PURPOSE     Computes descriptive statistics.

FORMAT     { *vnam,mean,var,std,min,max,valid,mis* } = **dstat**(*dataset,vars*);

INPUT     *dataset*     string, name of data set.  
                              If *dataset* is null or 0, *vars* will be assumed to be a matrix containing the data.

*vars*             the variables.  
                              If *dataset* contains the name of a **GAUSS** data set, *vars* will be interpreted as:

K×1 character vector, names of variables.

- or -

K×1 numeric vector, indices of 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.

If *dataset* is null or 0, *vars* will be interpreted as:

N×K matrix, the data on which to compute the descriptive statistics.

GLOBAL INPUT	<b>__altnam</b>	matrix, default 0. This can be a K×1 character vector of alternate variable names for the output.
	<b>__maxbytes</b>	scalar, the maximum number of bytes to be read per iteration of the read loop. Default = 1e9.
	<b>__maxvec</b>	scalar, the largest number of elements allowed in any one matrix. Default = 20000.
	<b>__miss</b>	scalar, default 0. <b>0</b> there are no missing values (fastest). <b>1</b> listwise deletion, drop a row if any missings occur in it. <b>2</b> pairwise deletion.
	<b>__row</b>	scalar, the number of rows to read per iteration of the read loop. if 0, (default) the number of rows will be calculated using <b>__maxbytes</b> and <b>__maxvec</b> .
	<b>__output</b>	scalar, controls output, default 1. <b>1</b> print output table. <b>0</b> do not print output.
OUTPUT	<i>vnam</i>	K×1 character vector, the names of the variables used in the statistics.
	<i>mean</i>	K×1 vector, means.
	<i>var</i>	K×1 vector, variance.

## dstatmt

---

<i>std</i>	K×1 vector, standard deviation.
<i>min</i>	K×1 vector, minima.
<i>max</i>	K×1 vector, maxima.
<i>valid</i>	K×1 vector, the number of valid cases.
<i>mis</i>	K×1 vector, the number of missing cases.

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.

SOURCE    `dstat.src`

## dstatmt

PURPOSE    Compute descriptive statistics.

FORMAT    `dout = dstatmt(dc0,dataset,vars);`

INPUT    *dc0*       instance of a **dstatmtControl** structure containing the following members:

<i>dc0.altnames</i>	K×1 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>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, default 0. <b>0</b> there are no missing values (fastest).



		<div><div>1</div><div>listwise deletion, drop a row if any missings occur in it.</div></div> <div><div>2</div><div>pairwise deletion.</div></div>
	<i>dc0.row</i>	<div>scalar, the number of rows to read per iteration of the read loop.</div> <div>If 0, (default) the number of rows will be calculated using <i>dc0.maxbytes</i> and <i>dc0.maxvec</i>.</div>
	<i>dc0.output</i>	<div>scalar, controls output, default 1.</div> <div><div>1</div><div>print output table.</div></div> <div><div>0</div><div>do not print output.</div></div>
	<i>dataset</i>	<div>string, name of data set.</div> <div>If <i>dataset</i> is null or 0, <i>vars</i> will be assumed to be a matrix containing the data.</div>
	<i>vars</i>	<div>the variables.</div> <div>If <i>dataset</i> contains the name of a <b>GAUSS</b> data set, <i>vars</i> will be interpreted as:</div> <div><div>K×1 string array, names of variables.</div><div>- or -</div><div>K×1 numeric vector, indices of variables.</div></div> <div>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.</div> <div>If <i>dataset</i> is null or 0, <i>vars</i> will be interpreted as:</div> <div><div>N×K matrix, the data on which to compute the descriptive statistics.</div></div>
OUTPUT	<i>dout</i>	<div>instance of a <b>dstatmtOut</b> structure containing the following members:</div> <div><div><i>dout.vnames</i></div><div>K×1 string array, the names of the variables used in the statistics.</div></div> <div><div><i>dout.mean</i></div><div>K×1 vector, means.</div></div> <div><div><i>dout.var</i></div><div>K×1 vector, variance.</div></div> <div><div><i>dout.std</i></div><div>K×1 vector, standard deviation.</div></div>

## dstatmtControlCreate

---

<i>dout.min</i>	K×1 vector, minima.
<i>dout.max</i>	K×1 vector, maxima.
<i>dout.valid</i>	K×1 vector, the number of valid cases.
<i>dout.missing</i>	K×1 vector, the number of missing cases.
<i>dout.errcode</i>	scalar, error code, 0 if successful; otherwise, one of the following: <b>2</b> Can't open file. <b>7</b> Too many missings - no data left after packing. <b>9</b> <b>altnames</b> member of <b>dstatmtControl</b> structure wrong size. <b>10</b> <b>vartype</b> member of <b>dstatmtControl</b> structure wrong size.

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.

SOURCE `dstatmt.src`

SEE ALSO **dstatmtControlCreate**

## dstatmtControlCreate

PURPOSE Creates default **dstatmtControl** structure.

INCLUDE `dstatmt.sdf`

FORMAT `c = dstatmtControlCreate;`

OUTPUT `c` instance of **dstatmtControl** structure with members set to default values.

SOURCE    `dstatmt.src`

SEE ALSO    **dstatmt**

**d**

**dtdate**

PURPOSE    Creates a matrix in DT scalar format.

FORMAT    `dt = dtdate(year,month,day,hour,minute,second);`

INPUT    *year*        N×K matrix of years.  
           *month*      N×K matrix of months, 1-12.  
           *day*        N×K matrix of days, 1-31.  
           *hour*       N×K matrix of hours, 0-23.  
           *minute*    N×K matrix of minutes, 0-59.  
           *second*    N×K matrix of seconds, 0-59.

OUTPUT    *dt*            N×K matrix of DT scalar format dates.

REMARKS    The arguments must be E×E conformable.

SOURCE    `time.src`

SEE ALSO    **dtday, dttime, utctodt, dttostr**

## datetime

### dtday

PURPOSE	Creates a matrix in DT scalar format containing only the year, month and day. Time of day information is zeroed out.		
FORMAT	<i>dt</i> = <b>dtday</b> ( <i>year</i> , <i>month</i> , <i>day</i> );		
INPUT	<i>year</i>	N×K matrix of years.	
	<i>month</i>	N×K matrix of months, 1-12.	
	<i>day</i>	N×K matrix of days, 1-31.	
OUTPUT	<i>dt</i>	N×K matrix of DT scalar format dates.	
REMARKS	This amounts to 00:00:00 or midnight on the given day. The arguments must be E×E conformable.		
SOURCE	time.src		
SEE ALSO	<b>dttime</b> , <b>dtdate</b> , <b>utctodt</b> , <b>dttostr</b>		

### dttime

PURPOSE	Creates a matrix in DT scalar format containing only the hour, minute and second. The date information is zeroed out.		
FORMAT	<i>dt</i> = <b>dttime</b> ( <i>hour</i> , <i>minute</i> , <i>second</i> );		
INPUT	<i>hour</i>	N×K matrix of hours, 0-23.	
	<i>minute</i>	N×K matrix of minutes, 0-59.	

---

*second*      N×K matrix of seconds, 0-59.

OUTPUT    *dt*            N×K matrix of DT scalar format times.

REMARKS    The arguments must be E×E conformable.

SOURCE    `time.src`

SEE ALSO    **dtday, dtdate, utctodt, dttostr**

## dttodtv

PURPOSE    Converts DT scalar format to DTV vector format.

FORMAT    *dtv* = **dttodtv**(*dt*);

INPUT      *dt*            N×1 vector, DT scalar format.

OUTPUT    *dtv*            N×8 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

## dttostr

---

[N,8] Days since Jan 1 of current year, 0-365

EXAMPLE    `dt = 20010326110722;`  
            `print "dt = " dt;`

            20010326110722

`dtv = dttodtv(dt);`  
            `print "dtv = " dtv;`

            2001      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      *x*            N×K matrix containing dates in DT scalar format.  
            *fmt*        string containing date/time format characters.

OUTPUT     *sa*           N×K string array.

REMARKS    The DT scalar format is a double precision representation of the date and time.  
            In the DT scalar format, the number

            20050703105031

represents 10:50:31 or 10:50:31 AM on July 3, 2005. **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
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    `s0 = dttostr(utctodt(timeutc), "YYYY-MO-DD HH:MI:SS");`  
              `print ("Date and Time are: " $+ s0);`

Date and time are: 2005-09-14 11:49:10

`print dttostr(utctodt(timeutc), "Today is DD-MO-YR");`

Today is 14-09-05

`s = dttostr(x, "YYYY-MO-DD");`

                 20000317060424  
                  20010427031213  
                  20010517020437  
                  20011117161422  
      If x = 20010717120448  
                  20010817043451  
                  20010919052320  
                  20011017032203  
                  20011107071418

## dttoutc

---

```
                2000 - 03 - 17
                2001 - 04 - 27
                2001 - 05 - 17
                2001 - 11 - 17
then s = 2001 - 07 - 17
                2001 - 08 - 17
                2001 - 09 - 19
                2001 - 10 - 17
                2001 - 11 - 07
```

SEE ALSO **strtodt**, **dttoutc**, **utctodt**

## dttoutc

**PURPOSE**     Converts DT scalar format to UTC scalar format.

**FORMAT**     *utc* = **dttoutc**(*dt*);

**INPUT**       *dt*            N×1 vector, DT scalar format.

**OUTPUT**      *utc*            N×1 vector, UTC scalar format.

**REMARKS**     In DT scalar format, 10:50:31 on July 15, 2005 is 20050703105031. 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*;  
  
                 *tc* = 985633642;



SOURCE `time.src`

SEE ALSO `dtvnormal`, `timeutc`, `utctodtv`, `dtctodtv`, `dtvtodt`, `dtvtoutc`, `dtvtodt`, `strtodt`, `dttostr`

d

**dtvnormal**

PURPOSE Normalizes a date and time (DTV) vector.

FORMAT  $d = \text{dtvnormal}(t);$

INPUT  $t$   $1 \times 8$  date and time vector that has one or more elements outside the normal range.

OUTPUT  $d$  Normalized  $1 \times 8$  date and time vector.

REMARKS The date and time vector is a  $1 \times 8$  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.

The last two elements are ignored on input.

EXAMPLE `format /rd 10,2;`  
`x = { 2005 14 21 6 21 37 0 0 };`  
`d = dtvnormal(x);`

$d = 2006 \ 2 \ 21 \ 6 \ 21 \ 37 \ 2 \ 51$

## dtvtodt

---

SEE ALSO **date, ethsec, etstr, time, timestr, timeutc, utctodtv**

### dtvtodt

**PURPOSE** Converts DT vector format to DT scalar format.

**FORMAT** *dt* = **dtvtodt**(*dtv*);

**INPUT** *dtv* N×8 matrix, DTV vector format.

**OUTPUT** *dt* N×1 vector, DT scalar format.

**REMARKS** In DT scalar format, 11:06:47 on March 15, 2001 is 20010315110647.

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 = { 2005 3 26 11 7 22 1 84 };  
dt = dtvtodt(dtv);
```

```
dtv = 2005 3 26 11 7 22 1 84;
```

```
dt = 20050326110722
```

SOURCE

time.src

SEE ALSO

dtvnormal, timeutc, utctodtv, dttodtv, dtvtodt, dttoutc, dtvtodt, strtodt, dttostr

d

dtvtoutc

PURPOSE

Converts DTV vector format to UTC scalar format.

FORMAT

*utc* = **dtvtoutc**(*dtv*);

INPUT

*dtv*            N×8 matrix, DTV vector format.

OUTPUT

*utc*            N×1 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
- [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 = 2005   9   14   14   40   18   3   256

utc = 1126734018

SEE ALSO **dtvnormal**, **timeutc**, **utctodt**, **dttdtv**, **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**  $y = \text{dummy}(x, v);$

**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**  $x = \{ 0, 2, 4, 6 \};$   
 $v = \{ 1, 5, 7 \};$   
 $y = \text{dummy}(x, v);$

The result **y** looks like this:

1	0	0	0
0	1	0	0
0	1	0	0
0	0	1	0

d

The vector **v** will produce 4 dummies satisfying the following conditions:

	<b>x</b>	≤	1
1	< <b>x</b>	≤	5
5	< <b>x</b>	≤	7
7	< <b>x</b>		

SOURCE    `datatran.src`

SEE ALSO    `dummybr`, `dummydn`

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*        N×1 vector of data that is to be broken up into dummy variables.  
          *v*        K×1 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*        N×K 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**  $x = \{ 0, \\ 2, \\ 4, \\ 6 \};$

$$v = \{ 1, \\ 5, \\ 7 \};$$
$$y = \text{dummybr}(x, v);$$

The resulting matrix **y** looks like this:

$$\begin{array}{ccc} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{array}$$

The vector  $v = 1 \ 5 \ 7$  will produce 3 dummies satisfying the following conditions:

$$\begin{array}{rcl} & x & \leq 1 \\ 1 & < x & \leq 5 \\ 5 & < x & \leq 7 \end{array}$$

SOURCE    `datatran.src`

SEE ALSO    `dummydn`, `dummy`

## dummydn

d

**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 = dummydn(x,v,p);`

**INPUT**

$x$	$N \times 1$ vector of data to be broken up into dummy variables.
$v$	$K \times 1$ 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.

**EXAMPLE**    `x = { 0, 2, 4, 6 };`

```
v = { 1, 5, 7 };  
p = 2;  
y = dummydn(x,v,p);
```

The resulting matrix **y** looks like this:

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

The vector **v** = 1 5 7 will produce 4 dummies satisfying the following conditions:

```
          x ≤ 1  
1 < x ≤ 5  
5 < x ≤ 7  
7 < x
```

SOURCE    `datatran.src`

SEE ALSO    **dummy**, **dummybr**

## ed

PURPOSE    Accesses an alternate editor.

FORMAT    **ed** *filename*;

INPUT    *filename*    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 type:

```
ed = editor_name flags;
```

or

```
ed = ‘‘editor_name flags’’;
```

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.

This command can be placed in the startup file, so it will be set for you automatically when you start **GAUSS**.

edit

PURPOSE Edits a disk file.

FORMAT **edit** *filename*;

INPUT *filename* literal, the name of the file to be edited.

PORTABILITY **Windows** and **Mac** only

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.

EXAMPLE `edit test1.e;`

## eig

---

SEE ALSO **run**

## eig

**PURPOSE** Computes the eigenvalues of a general matrix.

**FORMAT**  $va = \mathbf{eig}(x);$

**INPUT**  $x$   $N \times N$  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  $4 \times 4$  arrays contained in  $x$ .

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 **scalerr**( $va[1]$ )+1 to N 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

The eigenvalues are unordered except that complex conjugate pairs of eigenvalues will appear consecutively with the eigenvalue having the positive imaginary part first.

**EXAMPLE**  $x = \begin{Bmatrix} 4 & 8 & 1, \\ 9 & 4 & 2, \end{Bmatrix}$

```

      5 5 7 };

  va = eig(x);

      -4.4979246
  va =  14.475702
      5.0222223

```

e

SEE ALSO **eigh**, **eighv**, **eigv**

## eigh

**PURPOSE** Computes the eigenvalues of a complex hermitian or real symmetric matrix.

**FORMAT**  $va = \mathbf{eigh}(x);$

**INPUT**  $x$   $N \times N$  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  $4 \times 4$  arrays contained in  $x$ .

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.

## eighv

---

**trap 0** set *va*[1] and terminate with message  
**trap 1** set *va*[1] and continue execution

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

**PURPOSE** Computes eigenvalues and eigenvectors of a complex hermitian or real symmetric matrix.

**FORMAT** { *va*, *ve* } = **eighv**(*x*);

**INPUT** *x* N×N matrix or K-dimensional array where the last two dimensions are N×N.

**OUTPUT** *va* N×1 vector or K-dimensional array where the last two dimensions are N×1, the eigenvalues of *x*.

*ve* N×N matrix or K-dimensional array where the last two dimensions are N×N, 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 10×4×4 array, *va* will be a 10×4×1 array containing the eigenvalues and *ve* a 10×4×4 array containing the eigenvectors of each of the 10 4×4 arrays contained in *x*.

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 va[1] and terminate with message
trap 1  set va[1] and continue execution
```

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**

e

eigv

PURPOSE	Computes eigenvalues and eigenvectors of a general matrix.		
FORMAT	{ <i>va</i> , <i>ve</i> } = <b>eigv</b> ( <i>x</i> );		
INPUT	<i>x</i>	N×N matrix or K-dimensional array where the last two dimensions are N×N.	
OUTPUT	<i>va</i>	N×1 vector or K-dimensional array where the last two dimensions are N×1, the eigenvalues of <i>x</i> .	
	<i>ve</i>	N×N matrix or K-dimensional array where the last two dimensions are N×N, the eigenvectors of <i>x</i> .	
REMARKS	If <i>x</i> is an array, <i>va</i> will be an array containing the eigenvalues of each 2-dimensional array described by the two trailing dimensions of <i>x</i> , and <i>ve</i> will be an array containing the corresponding eigenvectors. In other words, for a		

10×4×4 array, *va* will be a 10×4×1 array containing the eigenvalues and *ve* a 10×4×4 array containing the eigenvectors of each of the 10 4×4 arrays contained in *x*.

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 **scalerr**(*va*[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 *va*[1] and terminate with message  
**trap 1** set *va*[1] and continue execution

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 *ve* contain the eigenvectors of *x* in the same order as the eigenvalues. The eigenvectors are not normalized.

EXAMPLE    *x* = { 4 8 1,  
                  9 4 2,  
                  5 5 7 };

              {*y*,*n*} = eigv(*x*);

                  -4.4979246  
*y* =     14.475702  
                  5.0222223

                  -0.66930459    -0.64076622    -0.40145623  
*n* =     0.71335708    -0.72488533    -0.26047487  
                  -0.019156716    -0.91339349     1.6734214

SEE ALSO    **eig, eigh, eighv**

end

## elapsedTradingDays

**PURPOSE** Computes number of trading days between two dates inclusively.

**FORMAT**  $n = \text{elapsedTradingDays}(a, b);$

**INPUT**  $a$  scalar, date in DT scalar format.  
 $b$  scalar, date in DT scalar format.

**OUTPUT**  $n$  number of trading days between dates inclusively, that is, elapsed time includes the dates  $a$  and  $b$ .

**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** `elapsedTradingDays`, `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**

## endp

---

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;

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 12.)

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);
```

```
          1.00000000  3.00000000  2.00000000  
x =       7.00000000  4.00000000  9.00000000  
          1.00000000  1.00000000  6.00000000  
          3.00000000  3.00000000  2.00000000
```

```
          3.00000000  
y =       5.00000000  
          2.00000000  
          7.00000000
```

```
          0.15456890  
b =       1.50276345  
          -0.12840825
```

SEE ALSO **proc, keyword, retp**

**endwind**

**PURPOSE** Ends graphic panel manipulation; displays graphs with **rerun**.

**LIBRARY** pgraph

**FORMAT** **endwind;**

**REMARKS** This function uses **rerun** to display the most recently created .tkf file.

## envget

---

SOURCE    `pwindow.src`

SEE ALSO    **begwind, window, makewind, setwind, nextwind, getwind**

## envget

PURPOSE    Searches the environment table for a defined name.

FORMAT    `y = envget(s);`

INPUT    `s`            string, the name to be searched for.

OUTPUT    `y`            string, the string that corresponds to that name in the environment table or a null string if it is not found.

EXAMPLE    

```
proc dopen(file);
    local fname,fp;
    fname = envget("DPATH");
    if fname $=\,= "";
        fname = file;
    else;
        if strsect(fname,strlen(fname),1) $=\,= "\\\";
            fname = fname $+ file;
        else;
            fname = fname $+ "\\\" $+ file;
        endif;
    endif;
    open fp = ^fname;
    retp(fp);
endp;
```

This is an example of a procedure that will open a data file using a path stored in an environment string called DPATH. The procedure returns the file handle and is called as follows:

---

```
fp = dopen('myfile');
```

SEE ALSO **cdir**

eof

e

**PURPOSE** Tests if the end of a file has been reached.

**FORMAT**  $y = \text{eof}(fh);$

**INPUT**  $fh$  scalar, file handle.

**OUTPUT**  $y$  scalar, 1 if end of file has been reached, else 0.

**REMARKS** This function is used with the **readr** and **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 ( $\mathbf{x}'\mathbf{x}$ ) 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.

## eqSolve

---

**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 *start* K×1 vector, starting values.  
&*F* scalar, a pointer to a procedure which computes the value at *x* of the equations to be solved.

GLOBAL INPUT The following are set by **eqSolveSet**:

**\_eqs\_JacobianProc** pointer to a procedure which computes the analytical Jacobian. By default, **eqSolve** will compute the Jacobian numerically.

**\_eqs\_MaxIters** scalar, the maximum number of iterations. Default = 100.

**\_eqs\_StepTol** scalar, the step tolerance. Default = **\_\_macheps**<sup>2/3</sup>.

**\_eqs\_TypicalF** K×1 vector of the typical *F*(*x*) values at a point not near a root, used for scaling. This becomes important when the magnitudes of the components of *F*(*x*) are expected to be very different. By default, function values are not scaled.

**\_eqs\_TypicalX** K×1 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.

**\_\_eqs\_IterInfo** scalar, if nonzero, iteration information is printed. Default = 0.

The following are set by **gausset**:

**\_\_Tol** scalar, the tolerance of the scalar function  $f = 0.5 * \|F(x)\|^2$  required to terminate the algorithm. Default = 1e-5.

**\_\_altnam** K×1 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 **\_\_vpad** is set) will be used.

**\_\_output** scalar. If non-zero, final results are printed.

**\_\_title** string, a custom title to be printed at the top of the iterations report. By default, only a generic title will be printed.

**\_\_vpad** scalar. If **\_\_altnam** is not set, variable names are automatically created. Two types of names can be created:

- 0** Variable names are not padded to give them equal length. For example, X1, X2,...,X10,...
- 1** Variable names are padded with zeros to give them an equal number of characters. For example, X01,X02,...,X10,... This is useful if you want the variable names to sort properly.

OUTPUT  $x$  K×1 vector, solution.

*retcode* scalar, the return code:

- 1** Norm of the scaled function value is less than **\_\_Tol**.  $x$  given is an approximate root of  $F(x)$  (unless **\_\_Tol** is too large).
- 2** The scaled distance between the last two steps is less than the step-tolerance (**\_\_eqs\_StepTol**).  $x$  may be an approximate root of  $F(x)$ , 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  $\text{norm2}(F(x))$  sufficiently; either  $x$  is close to a root of  $F(x)$  and no more accuracy is

e

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  $\text{norm2}(F(x))$  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  $\text{norm2}(F(x))$  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 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 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 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 Version 6.5.0                                8/18/2005   3:33 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-06
X2	12.00000	1.4085912	-6.6263102e-06
X3	-1.00000	1.1111111	4.4175402e-06

e

## eqSolvemt

---

SOURCE    eqsolve.src

### eqSolvemt

PURPOSE    Solves a system of nonlinear equations.

INCLUDE    eqsolvemt.sdf

FORMAT    *out* = **eqSolvemt**(&*fct*,*par*,*data*,*c*);

INPUT    *&fct*      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.

*par*        an instance of a **PV** structure. The *par* instance is passed to the user-provided procedure pointed to by *&fct*. *par* is constructed using the **pvPack** functions.

*data*      an array of instances of a **DS** structure. This array is passed to the user-provided procedure pointed to by *&fct* to be used in the objective function. **eqSolvemt** does not look at this structure. Each instance contains the the following members which can be set in whatever way that is convenient for computing the objective function:

*data1*[*i*].**dataMatrix**     $N \times K$  matrix, data matrix.  
                    *data1*[*i*].**dataArray**     $N \times K \times L \dots$  array, data array.  
                    *data1*[*i*].**vnames**        string array, variable names (optional).  
                    *data1*[*i*].**dsname**        string, data name (optional).  
                    *data1*[*i*].**type**        scalar, type of data (optional).

*c*         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 *c*, the members are:



	<i>c.jacobianProc</i>	pointer to a procedure which computes the analytical Jacobian. By default, <b>eqSolvemt</b> 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>	$K \times 1$ vector of the typical $fct(X)$ values at a point not near a root, used for scaling. This becomes important when the magnitudes of the components of $fct(X)$ are expected to be very different. By default, function values are not scaled.
	<i>c.typicalX</i>	$K \times 1$ 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.
	<i>c.printIters</i>	scalar, if nonzero, iteration information is printed. Default = 0.
	<i>c.tolerance</i>	scalar, the tolerance of the scalar function $f = 0.5 * \ fct(X)\ ^2$ required to terminate the algorithm. That is, the condition that $ f(x)  \leq c.\text{tolerance}$ must be met before that algorithm can terminate successfully. Default = $1e-5$ .
	<i>c.altnam</i>	$K \times 1$ character vector of alternate names to be used by the printed output. By default, the names "X1,X2,X3..." will be used.
	<i>c.title</i>	string, printed as a title in output.
	<i>c.output</i>	scalar. If non-zero, final results are printed.
OUTPUT	<i>out</i>	an instance of an <b>eqsolvemtOut</b> structure. For an instance named <i>out</i> , the members are:
	<i>out.par</i>	an instance of a <b>PV</b> structure containing the parameter estimates.

---

<code>out.fct</code>	scalar, function evaluated at $X$ .
<code>out.retcode</code>	scalar, return code: <ul style="list-style-type: none"><li>-1 Jacobian is singular.</li><li>1 Norm of the scaled function value is less than <code>c.tolerance</code>. <math>X</math> given is an approximate root of <math>fct(X)</math> (unless <code>c.tolerance</code> is too large).</li><li>2 The scaled distance between the last two steps is less than the step-tolerance (<code>c.stepTolerance</code>). <math>X</math> may be an approximate root of <math>fct(X)</math>, 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.</li><li>3 The last global step failed to decrease <math>\text{norm2}(fct(X))</math> sufficiently; either <math>X</math> is close to a root of <math>fct(X)</math> 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.</li><li>4 Iteration limit exceeded.</li><li>5 Five consecutive steps of maximum step length have been taken; either <math>\text{norm2}(fct(X))</math> asymptotes from above to a finite value in some direction or the maximum step length is too small.</li><li>6 <math>X</math> seems to be an approximate local minimizer of <math>\text{norm2}(fct(X))</math> that is not a root of <math>fct(X)</math>. To find a root of <math>fct(X)</math>, restart <b>eqSolvemt</b> from a different region.</li></ul>

REMARKS     The equation procedure should return a column vector containing the result for each equation.

If there is no data, you can pass an empty **DS** structure in the second argument:

```
call eqSolvemt(&fct,par,dsCreate,c);
```

**EXAMPLE**

Equation 1:  $x_1^2 + x_2^2 - 2 = 0$

Equation 2:  $\exp(x_1 - 1) + x_2^3 - 2 = 0$

e

```
#include eqSolvemt.sdf;
struct eqSolvemtControl c;
c = eqSolvemtControlCreate;

c.printIters = 1;

struct PV par;
par = pvPack(pvCreate,1,"x1");
par = pvPack(par,1,"x2");

struct eqSolvemtOut out1;
out1 = eqSolvemt(&fct,par,dsCreate,c);

proc fct(struct PV p, struct DS d);
  local x1, x2, z;
  x1 = pvUnpack (p, "x1");
  x2 = pvUnpack (p, "x2");
  z = x1^2+x2^2-2 | exp(x1-1)+x2^3-2;
  retp(z);
endp;
```

**SOURCE** eqsolvemt.src

**SEE ALSO** eqsolvemtControlCreate, eqsolvemtOutCreate

## eqSolvemtOutCreate

---

### eqSolvemtControlCreate

**PURPOSE**    Creates default **eqSolvemtControl** structure.

**INCLUDE**    eqsolvemt.sdf

**FORMAT**    *c* = **eqSolvemtControlCreate**;

**OUTPUT**    *c*            instance of **eqSolvemtControl** structure with members set to default values.

**SOURCE**    eqsolvemt.src

**SEE ALSO**    **eqsolvemt**

### eqSolvemtOutCreate

**PURPOSE**    Creates default **eqSolvemtOut** structure.

**FORMAT**    *c* = **eqSolvemtOutCreate**;

**OUTPUT**    *c*            instance of **eqSolvemtOut** structure with members set to default values.

**SOURCE**    eqsolvemt.src

**SEE ALSO**    **eqsolvemt**

eqSolveSet

PURPOSE     Sets global input used by **eqSolve** to default values.

FORMAT     **eqSolveset;**

GLOBAL     **\_\_eqs\_TypicalX**     Set to 0.  
OUTPUT     **\_\_eqs\_TypicalF**     Set to 0.  
             **\_\_eqs\_IterInfo**     Set to 0.  
             **\_\_eqs\_JacobianProc**     Set to 0.  
             **\_\_eqs\_MaxIters**     Set to 100.  
             **\_\_eqs\_StepTol**     Set to **\_\_macheps**<sup>2/3</sup>

e

erf, erfc

PURPOSE     Computes the Gaussian error function (**erf**) and its complement (**erfc**).

FORMAT     **y = erf(x);**  
             **y = erfc(x);**

INPUT     *x*             N×K matrix.

OUTPUT     *y*             N×K matrix.

REMARKS     The allowable range for *x* is:

$$x \geq 0$$

## error

---

The **erf** and **erfc** functions are closely related to the Normal distribution:

$$\text{cdfn}(x) = \begin{cases} \frac{1}{2}(1 + \text{erf}(\frac{x}{\sqrt{2}})) & x \geq 0 \\ \frac{1}{2}\text{erfc}(\frac{-x}{\sqrt{2}}) & x < 0 \end{cases}$$

EXAMPLE    `x = { .5 .4 .3,  
                  .6 .8 .3 };`  
              `y = erf(x);`

```
      0.52049988  0.42839236  0.32862676  
y =  0.60385609  0.74210096  0.32862676
```

`x = { .5 .4 .3,  
                  .6 .8 .3 };`  
              `y = erfc(x);`

```
      0.47950012  0.57160764  0.67137324  
y =  0.39614391  0.25789904  0.67137324
```

SEE ALSO    **cdfn**, **cdfnc**

TECHNICAL    **erf** and **erfc** are computed by summing the appropriate series and continued  
NOTES        fractions. They are accurate to about 10 digits.

## 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 <b>scalerr</b> function.
REMARKS	<p>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 <b>scalerr</b>.</p> <p>The <b>scalerr</b> function will return the value of the error code and so is the reverse of <b>error</b>. These user-generated error codes work in the same way as the intrinsic <b>GAUSS</b> error codes which are generated automatically when <b>trap 1</b> is on and certain <b>GAUSS</b> functions detect a numerical error such as a singular matrix.</p> <p><b>error(0)</b> is equal to the missing value code.</p>	
EXAMPLE	<pre>proc syminv(x);     local oldtrap,y;     if not x =\,= x';         retp(error(99));     endif;     oldtrap = trapchk(0xffff);     trap 1;     y = invpd(x);     if scalerr(y);         y = inv(x);     endif;     trap oldtrap,0xffff;     retp(y); endp;</pre> <p>The procedure <b>syminv</b> returns error code 99 if the matrix is not symmetric. If <b>invpd</b> fails, it returns error code 20. If <b>inv</b> fails, it returns error code 50. The original trap state is restored before the procedure returns.</p>	
SEE ALSO	<b>scalerr</b> , <b>trap</b> , <b>trapchk</b>	

## errorlogat

---

### errorlog

PURPOSE	Prints an error message to the window and error log file.
FORMAT	<b>errorlog</b> <i>str</i> ;
INPUT	<i>str</i> string, the error message to print.
REMARKS	This command enables you to do your own error handling in your <b>GAUSS</b> programs. To print an error message to the window and error log file along with file name and line number information, use <b>errorlogat</b> .
SEE ALSO	<b>errorlogat</b>

### 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	<b>errorlogat</b> <i>str</i> ;
INPUT	<i>str</i> string, the error message to print.
REMARKS	This command enables you to do your own error handling in your <b>GAUSS</b> programs. To print an error message to the window and error log file without file name and line number information, use <b>errorlog</b> .
SEE ALSO	<b>errorlog</b>



## etdays

e

**PURPOSE**     Computes the difference between two times, as generated by the **date** command, in days.

**FORMAT**     *days* = **etdays**(*tstart*,*tend*);

**INPUT**     *tstart*     3×1 or 4×1 vector, starting date, in the order: yr, mo, day. (Only the first 3 elements are used.)

*tend*     3×1 or 4×1 vector, ending date, in the order: yr, mo, day. (Only the first 3 elements are used.) **MUST** be later than *tstart*.

**OUTPUT**     *days*     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 = 2004 1 2;`  
                  `let date2 = 2005 9 14;`  
                  `d = etdays(date1,date2);`

`d = 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.

**FORMAT**     *hs* = **ethsec**(*tstart*,*tend*);

**INPUT**       *tstart*       4×1 vector, starting date, in the order: yr, mo, day, hundredths of a second.

*tend*       4×1 vector, ending date, in the order: yr, mo, day, hundredths of a second. **MUST** be later date than *tstart*.

**OUTPUT**     *hs*           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 = 2004 1 2 0;
let date2 = 2005 9 14 0;
t = ethsec(date1,date2);
```

$$t = 5365440000$$

**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**      *tothsecs*    scalar, an elapsed time measured in hundredths of a second, as given, for instance, by the **ethsec** function.
- OUTPUT**    *str*            string containing the elapsed time in the form:  
                              # days    # hours    # minutes    #,## seconds
- EXAMPLE**    `d1 = { 2004, 1, 2, 0 };`  
                 `d2 = { 2004, 9, 14, 815642 };`  
                 `t = ethsec(d1,d2);`  
                 `str = etstr(t);`
- `t = 5366255642`
- `str = 34 days 2 hours 15 minutes 56.42 seconds`
- SOURCE**    `time.src`
- SEE ALSO**    **ethsec**

**EuropeanBinomCall**

- PURPOSE**    Prices European call options using binomial method.

## EuropeanBinomCall\_Greeks

---

FORMAT     $c = \text{EuropeanBinomCall}(S0, K, r, div, tau, sigma, N);$

INPUT     $S0$        scalar, current price.  
           $K$          $M \times 1$  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    $c$          $M \times 1$  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    $S0 = 718.46;$   
             $K = \{ 720, 725, 730 \};$   
             $r = .0498;$   
             $sigma = .2493;$   
             $t0 = \text{dtday}(2001, 1, 30);$   
             $t1 = \text{dtday}(2001, 2, 16);$   
             $tau = \text{elapsedTradingDays}(t0, t1) / \text{annualTradingDays}(2001);$   
             $c = \text{EuropeanBinomCall}(S0, K, r, 0, tau, sigma, 60);$   
             $\text{print } c;$   
  
            17.344044  
            15.058486  
            12.817427

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**

$S0$	scalar, current price.
$K$	M×1 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.

**GLOBAL INPUT**

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

**OUTPUT**

$d$	M×1 vector, delta.
$g$	M×1 vector, gamma.
$t$	M×1 vector, theta.
$v$	M×1 vector, vega.
$rh$	M×1 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.

e

## EuropeanBinomCall\_ImpVol

---

```
EXAMPLE  S0 = 305;
          K = 300;
          r = .08;
          sigma = .25;
          tau = .33;
          div = 0;
          print EuropeanBinomcall_Greeks(S0,K,r,0,tau,sigma,30);

          0.70631204
          0.00076381912
          -44.616125
          68.703851
          76.691829
```

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  $\sigma = \text{EuropeanBinomCall\_ImpVol}(c, S0, K, r, \text{div}, \text{tau}, N);$

INPUT	$c$	M×1 vector, call premiums.
	$S0$	scalar, current price.
	$K$	M×1 vector, strike prices.
	$r$	scalar, risk free rate.
	$\text{div}$	continuous dividend yield.
	$\text{tau}$	scalar, elapsed time to exercise in annualized days of trading.
	$N$	number of time segments.

OUTPUT *sigma* M×1 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 = EuropeanBinomCall_ImpVol(c,S0,K,r,0,tau,30);
print sigma;

0.19632223
0.16988741
0.12879447

```

SOURCE finprocs.src

## EuropeanBinomPut

PURPOSE Prices European put options using binomial method.

FORMAT *c* = **EuropeanBinomPut**(*S0*,*K*,*r*,*div*,*tau*,*sigma*,*N*);

INPUT

<i>S0</i>	scalar, current price.
<i>K</i>	M×1 vector, strike prices.
<i>r</i>	scalar, risk free rate.

## EuropeanBinomPut\_Greeks

---

	<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>	M×1 vector, put premiums.
REMARKS	The binomial method of Cox, Ross, and Rubinstein (“Option pricing: a simplified approach”, <i>Journal of Financial Economics</i> , 7:229:264) as described in <i>Options, Futures, and other Derivatives</i> by John C. Hull is the basis of this procedure.	
EXAMPLE	<pre>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 = EuropeanBinomPut(S0,K,r,0,tau,sigma,60); print c;</pre> <pre>16.851815 19.580390 22.353464</pre>	
SOURCE	finprocs.src	

## EuropeanBinomPut\_Greeks

PURPOSE	Computes Delta, Gamma, Theta, Vega, and Rho for European put options using binomial method.
---------	---



FORMAT {  $d, g, t, v, rh$  } =  
**EuropeanBinomPut\_Greeks**( $S0, K, r, div, tau, sigma, N$ );

INPUT  $S0$  scalar, current price.  
 $K$  M×1 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.

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  $d$  M×1 vector, delta.  
 $g$  M×1 vector, gamma.  
 $t$  M×1 vector, theta.  
 $v$  M×1 vector, vega.  
 $rh$  M×1 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**( $S0, K, r, 0, tau, sigma, 60$ );

## EuropeanBinomPut\_ImpVol

---

-0.36885112  
0.0011457287  
6.6396424  
68.979259  
-33.796807

SOURCE    `finprocs.src`

SEE ALSO    `EuropeanBinomPut_Impvol`, `EuropeanBinomPut`,  
              `EuropeanBinomCall_Greeks`, `EuropeanBSPut_Greeks`

## EuropeanBinomPut\_ImpVol

PURPOSE    Computes implied volatilities for European put options using binomial method.

FORMAT     $\sigma = \text{EuropeanBinomPut\_ImpVol}(c, S_0, K, r, \text{div}, \text{tau}, N);$

INPUT	$c$	M×1 vector, put premiums.
	$S_0$	scalar, current price.
	$K$	M×1 vector, strike prices.
	$r$	scalar, risk free rate.
	$\text{div}$	continuous dividend yield.
	$\text{tau}$	scalar, elapsed time to exercise in annualized days of trading.
	$N$	number of time segments.

OUTPUT     $\sigma$         M×1 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 = EuropeanBinomPut_ImpVol(p,S0,K,r,0,tau,30);
          print sigma;

          0.13006393
          0.17043648
          0.21499803

```

SOURCE finprocs.src

## EuropeanBSCall

PURPOSE Prices European call options using Black, Scholes and Merton method.

FORMAT  $c = \text{EuropeanBSCall}(S0, K, r, div, tau, sigma);$

INPUT	$S0$	scalar, current price.
	$K$	$M \times 1$ 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.

OUTPUT  $c$   $M \times 1$  vector, call premiums.

## EuropeanBSCall\_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 = EuropeanBSCall(S0,K,r,0,tau,sigma);
          print c;
```

17.249367

14.908466

12.796356

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>	M×1 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** **\_fin\_thetaType** scalar, if 1, one day look ahead, else, infinitesimal. Default = 0.

**\_fin\_epsilon** scalar, finite difference stepsize. Default = 1e-8.

OUTPUT *d* M×1 vector, delta.  
*g* M×1 vector, gamma.  
*t* M×1 vector, theta.  
*v* M×1 vector, vega.  
*rh* M×1 vector, rho.

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

```
0.64458005
0.0085029307
-38.505439
65.256273
56.872007
```

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);`

## EuropeanBSPut

---

INPUT    *c*            M×1 vector, call premiums.  
          *S0*           scalar, current price.  
          *K*            M×1 vector, strike prices.  
          *r*            scalar, risk free rate.  
          *div*          continuous dividend yield.  
          *tau*          scalar, elapsed time to exercise in annualized days of trading.

OUTPUT   *sigma*       M×1 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* = EuropeanBSCall\_ImpVol(*c*,*S0*,*K*,*r*,0,*tau*);  
            print *sigma*;

            0.19724517  
            0.17084848  
            0.12978762

SOURCE   finprocs.src

## EuropeanBSPut

PURPOSE   Prices European put options using Black, Scholes, and Merton method.

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

INPUT    *S0*           scalar, current price.

e

*K*            M×1 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.

OUTPUT    *c*            M×1 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 = EuropeanBSPut(S0,K,r,0,tau,sigma);  
            print c;  
  
            16.759909  
            19.404914  
            22.278710
```

SOURCE    finprocs.src

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      *S0*            scalar, current price.

## EuropeanBSPut\_ImpVol

---

*K*            M×1 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.

GLOBAL    **\_fin\_thetaType**    scalar, if 1, one day look ahead, else, infinitesimal. Default  
INPUT                    = 0.  
             **\_fin\_epsilon**    scalar, finite difference stepsize. Default = 1e-8.

OUTPUT    *d*            M×1 vector, delta.  
            *g*            M×1 vector, gamma.  
            *t*            M×1 vector, theta.  
            *v*            M×1 vector, vega.  
            *rh*          M×1 vector, rho.

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

```
-0.35541995  
0.0085029307  
-15.130748  
65.256273  
-39.548591
```

SOURCE    `finprocs.src`

SEE ALSO    **EuropeanBSPut\_Impvol, EuropeanBSPut, EuropeanBSCall\_Greeks,  
EuropeanBinomPut\_Greeks**



## EuropeanBSPut\_ImpVol

**PURPOSE** Computes implied volatilities for European put options using Black, Scholes, and Merton method.

**FORMAT**  $\sigma = \text{EuropeanBSPut\_ImpVol}(c, S_0, K, r, \text{div}, \tau);$

**INPUT**

$c$	M×1 vector, put premiums
$S_0$	scalar, current price.
$K$	M×1 vector, strike prices.
$r$	scalar, risk free rate.
$\text{div}$	continuous dividend yield.
$\tau$	scalar, elapsed time to exercise in annualized days of trading.

**OUTPUT**  $\sigma$  M×1 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 = EuropeanBSPut_ImpVol(p,S0,K,r,0,tau);
print sigma;
```

```
0.13222999
0.17115393
0.21666496
```

**SOURCE** finprocs.src

e

## exctsmpl

**PURPOSE**     Computes a random subsample of a data set.

**FORMAT**      $n = \text{exctsmpl}(\text{infile}, \text{outfile}, \text{percent});$

**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.

**OUTPUT**      $n$      scalar, number of rows in output data set.  
 Error returns are controlled by the low bit of the trap flag:

<b>trap 0</b>	terminate with error message
<b>trap 1</b>	return scalar negative integer
<b>-1</b>	can't open input file
<b>-2</b>	can't open output file
<b>-3</b>	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**      $n = \text{exctsmpl}(\text{"freqdata.dat"}, \text{"rout"}, 30);$

$n = 120$

*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**

**e**

PURPOSE    Executes an executable program and returns the exit code to **GAUSS**.

FORMAT    `y = exec(program,comline);`

INPUT    *program*    string, the name of the program, including the extension, to be executed.

*comline*    string, the arguments to be placed on the command line of the program being executed.

OUTPUT    *y*            scalar, the exit code returned by *program*.

          If **exec** can't 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 = exec("atog","comd1.cmd");`  
              `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, Chapter 26.

---

**execbg**

**PURPOSE**     Executes an executable program in the background and returns the process id to **GAUSS**.

**FORMAT**     *pid* = **execbg**(*program*,*comline*);

**INPUT**        *program*     string, the name of the program, including the extension, to be executed.

*comline*     string, the arguments to be placed on the command line of the program being executed.

**OUTPUT**      *pid*            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 26.

exp

PURPOSE     Calculates the exponential function.

FORMAT      $y = \text{exp}(x);$

INPUT        $x$               $N \times K$  matrix or  $N$ -dimensional array.

OUTPUT      $y$               $N \times K$  matrix or  $N$ -dimensional array containing  $e$ , the base of natural logs, raised to the powers given by the elements of  $x$ .

EXAMPLE     $x = \text{eye}(3);$   
               $y = \text{exp}(x);$

$x =$

1.000000	0.000000	0.000000
0.000000	1.000000	0.000000
0.000000	0.000000	1.000000

  

$y =$

2.718282	1.000000	1.000000
1.000000	2.718282	1.000000
1.000000	1.000000	2.718282

This example creates a 3×3 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)

## external

---

**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 olddata newdata;
    extern base, exvec;
    make ndvar = exvec[sega(base,1,rows(x_x))];
    # base = base + rows(x_x); /* execute command */
                                /* literally */
enddata;
```

## 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,spb*;

REMARKS    See PROCEDURES AND KEYWORDS, Chapter 12.

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**  $y = \text{eye}(n);$

**INPUT**  $n$  scalar, size of identity matrix to be created.

**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**  $x = \text{eye}(3);$

```
      1.000000  0.000000  0.000000  
x =  0.000000  1.000000  0.000000  
      0.000000  0.000000  1.000000
```



SEE ALSO **zeros, ones**

## fcheckerr

e

**PURPOSE** Gets the error status of a file.

**FORMAT** *err* = **fcheckerr**(*f*);

**INPUT** *f* scalar, file handle of a file opened with **fopen**.

**OUTPUT** *err* 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

**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

## feq, fge, fgt, fle, flt, fne

---

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**  $y = \text{feq}(a,b);$   
 $y = \text{fge}(a,b);$   
 $y = \text{fgt}(a,b);$   
 $y = \text{fle}(a,b);$   
 $y = \text{flt}(a,b);$   
 $y = \text{fne}(a,b);$

**INPUT**  $a$   $N \times K$  matrix, first matrix.  
 $b$   $L \times M$  matrix, second matrix,  $E \times E$  compatible with  $a$ .

**GLOBAL INPUT** **\_fcmptol** scalar, comparison tolerance. The default value is 1.0e-15.

OUTPUT     *y*            scalar, 1 (TRUE) or 0 (FALSE).

REMARKS    The return value is TRUE if every comparison is TRUE.

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 **\_fcmtol** before calling these procedures:

```
_fcmtol = 1e-12;
```

EXAMPLE    *x* = rndu(2,2);  
              *y* = rndu(2,2);  
              *t* = fge(*x*,*y*);

```

x =      0.038289504  0.072535275
      0.014713947   0.96863611

```

```

y =      0.25622293  0.70636474
      0.0036191244  0.35913385

```

```
t = 0.0000000
```

## feqmt, fgemt, fgtmt, flemt, fltmt, fnemt

---

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**    *a*            N×K matrix, first matrix.

*b*            L×M matrix, second matrix, E×E compatible with *a*.

*fcmtol*      scalar, comparison tolerance.

**OUTPUT**    *y*            scalar, 1 (TRUE) or 0 (FALSE).

**REMARKS**    The return value is TRUE if every comparison is TRUE.

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    `x = rndu(2,2);`  
              `y = rndu(2,2);`  
              `t = fgemt(x,y,1e-14);`

```
x =    0.038289504  0.072535275
      0.014713947  0.96863611
```

```
y =    0.25622293  0.70636474
      0.0036191244 0.35913385
```

```
t = 0.0000000
```

SOURCE    `fcomparemt.src`

SEE ALSO    `dotfeqmt--dotfnemt`

fflush

PURPOSE    Flushes a file's output buffer.

FORMAT    `ret = fflush(f);`

## fft

---

INPUT	<i>f</i>	scalar, file handle of a file opened with <b>fopen</b> .
OUTPUT	<i>ret</i>	scalar, 0 if successful, -1 if not.
REMARKS	If <b>fflush</b> fails, you can call <b>fstderr</b> to find out why.  If you pass <b>fflush</b> the handle of a file opened with <b>open</b> (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	<i>y</i> = <b>fft</b> ( <i>x</i> );	
INPUT	<i>x</i>	N×K matrix.
OUTPUT	<i>y</i>	L×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 <i>x</i> , scaled by 1/N.  This uses a Temperton Fast Fourier algorithm.  If N or K is not a power of 2, <i>x</i> will be padded out with zeros before computing the transform.	
EXAMPLE	<i>x</i> = { 22 24, 23 25 }; <i>y</i> = <b>fft</b> ( <i>x</i> );  <i>y</i> =     23.500000   -1.0000000 -0.500000   0.0000000	

SEE ALSO **ffti, rfft, rffti**

## ffti

**PURPOSE** Computes an inverse 1- or 2-D Fast Fourier transform.

**FORMAT**  $y = \text{ffti}(x);$

**INPUT**  $x$   $N \times K$  matrix.

**OUTPUT**  $y$   $L \times M$  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.

**EXAMPLE**  $x = \begin{Bmatrix} 22 & 24, \\ 23 & 25 \end{Bmatrix};$   
 $y = \text{fft}(x);$

$$y = \begin{bmatrix} 23.500000 & -1.0000000 \\ -0.5000000 & 0.0000000 \end{bmatrix}$$

$$fi = \text{ffti}(y);$$

$$fi = \begin{bmatrix} 22.000000 & 24.000000 \\ 23.000000 & 25.000000 \end{bmatrix}$$

SEE ALSO **fft, rfft, rffti**

## fftm

**PURPOSE** Computes a multi-dimensional FFT.

**FORMAT**  $y = \text{fftm}(x, dim);$

**INPUT**  $x$   $M \times 1$  vector, data.  
 $dim$   $K \times 1$  vector, size of each dimension.

**OUTPUT**  $y$   $L \times 1$  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  $\mathbf{x}$  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,  $\mathbf{x}$  would look something like this:

$$\begin{aligned} X_{hyper} &= X_{cube1} | X_{cube2} \\ X_{cube1} &= X_{mat1} | X_{mat2} \\ X_{mat1} &= X_{row1} | X_{row2} \\ X_{row1} &= X_{col1} | X_{col2} \end{aligned}$$

Or, in an extended **GAUSS** notation,  $\mathbf{x}$  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,  $\mathbf{x}$  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  $\mathbf{x}$ , 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  $2 \times 3 \times 2 \times 3$  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  $2^{nd}$  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  $2 \times 3 \times 3$  three-dimensional array, i.e., two  $3 \times 3$  matrices of data. Suppose that **x1** is the first  $3 \times 3$  matrix and **x2** the second  $3 \times 3$  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  $2 \times 3 \times 2 \times 3$  hypercube example, **x** would be padded from  $2 \times 3 \times 2 \times 3$  out to  $2 \times 4 \times 2 \times 4$ . 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 = fftmi(x,dim);`

INPUT    *x*             $M \times 1$  vector, data.

	<i>dim</i>	K×1 vector, size of each dimension.
OUTPUT	<i>y</i>	L×1 vector, inverse FFT of <i>x</i> .
REMARKS	<p>The multi-dimensional data are laid out in a recursive or heirarchical fashion in the vector <i>x</i>. 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 2×2×2×2 hypercubic <b>x</b> 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, <b>x</b> would look something like this:</p>	

$$\begin{aligned}
 X_{hyper} &= X_{cube1} \mid X_{cube2} \\
 X_{cube1} &= X_{mat1} \mid X_{mat2} \\
 X_{mat1} &= X_{row1} \mid X_{row2} \\
 X_{row1} &= X_{col1} \mid X_{col2}
 \end{aligned}$$

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  $2 \times 3 \times 2 \times 3$  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  $2^{nd}$  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  $2 \times 3 \times 3$  three-dimensional array, i.e., two  $3 \times 3$  matrices of data. Suppose that **x1** is the first  $3 \times 3$  matrix and **x2** the second  $3 \times 3$

matrix, then  $\mathbf{x} = \mathbf{vecr}(\mathbf{x1})|\mathbf{vecr}(\mathbf{x2})$ .

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 2×3×2×3 hypercube example,  $\mathbf{x}$  would be padded from 2×3×2×3 out to 2×4×2×4. The input vector would contain 36 elements, while the output vector would contain 64 elements.

SOURCE     `fftm.src`

SEE ALSO    `fftm`, `fft`, `ffti`, `fftn`

f

fftn

PURPOSE    Computes a complex 1- or 2-D FFT.

FORMAT     `y = fftn(x);`

INPUT       *x*            N×K matrix.

OUTPUT      *y*            L×M 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

$$2^p \times 3^q \times 5^r \times 7^s, \quad p,q,r \text{ nonnegative integers} \\ s=0 \text{ or } 1$$

If a dimension of  $x$  does not meet this requirement, it will be padded with zeros to the next allowable size before the 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  $33600 \times 1$  vector can compute as much as 20% faster than a  $32768 \times 1$  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**, **fftn**, **rfft**, **rffti**, **rfftm**, **fftn**, **rfftn**, **rfftnp**, **rfftp**

## fgets

PURPOSE Reads a line of text from a file.

FORMAT *str* = **fgets**(*f*, *maxsize*);

INPUT *f* scalar, file handle of a file opened with **fopen**.  
*maxsize* scalar, maximum size of string to read in, including the terminating null byte.

OUTPUT *str* 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.

SEE ALSO **fgetst**, **fgetsa**, **fopen**

f

## fgetsa

**PURPOSE** Reads lines of text from a file into a string array.

**FORMAT** *sa* = **fgetsa**(*f*,*numl*);

**INPUT** *f* scalar, file handle of a file opened with **fopen**.  
*numl* scalar, number of lines to read.

**OUTPUT** *sa* N×1 string array, N ≤ *numl*.

**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 **fgetsa**, your program will

## fgetsat

---

terminate with an error. Use **eof** in conjunction with **fgetsa** 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 **fgetsa**, **fgets**, **fopen**

## fgetsat

PURPOSE Reads lines of text from a file into a string array.

FORMAT *sa* = **fgetsat**(*f*,*numl*);

INPUT *f* scalar, file handle of a file opened with **fopen**.  
*numl* scalar, number of lines to read.

OUTPUT *sa* N×1 string array, N ≤ *numl*.

REMARKS **fgetsat** operates identically to **fgetsa**, except that newlines are not retained as 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** *f* scalar, file handle of a file opened with **fopen**.  
*maxsize* scalar, maximum size of string to read in, including the null terminating byte.

**OUTPUT** *str* 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** *fspec* string, file specification. Can include path. Wildcards are allowed in *fspec*.

OUTPUT    *fnames*    N×1 string array of all file names that match, null string if none are found.

*finfo*    N×13 matrix, information about matching files.

### UNIX/Linux

- [N, 1] filesystem ID
- [N, 2] inode number
- [N, 3] mode bit mask
- [N, 4] number of links
- [N, 5] user ID
- [N, 6] group ID
- [N, 7] device ID (char/block special files only)
- [N, 8] size in bytes
- [N, 9] last access time
- [N,10] last data modification time
- [N,11] last file status change time
- [N,12] preferred I/O block size
- [N,13] number of 512-byte blocks allocated

### Windows

- [N, 1] drive number (A = 0, B = 1, etc.)
- [N, 2] n/a, 0
- [N, 3] mode bit mask
- [N, 4] number of links, always 1
- [N, 5] n/a, 0
- [N, 6] n/a, 0
- [N, 7] n/a, 0
- [N, 8] size in bytes
- [N, 9] last access time
- [N,10] last data modification time
- [N,11] creation time
- [N,12] n/a, 0
- [N,13] n/a, 0

*finfo* 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

f

PURPOSE Returns a string array of file names.

FORMAT  $y = \text{filesa}(n);$

INPUT  $n$  string, file specification to search for. Can include path. Wildcards are allowed in  $n$ .

OUTPUT  $y$   $N \times 1$  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.

EXAMPLE  $y = \text{filesa}(\text{"ch*"});$

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.

## floor

---

SEE ALSO **fileinfo, shell**

### floor

PURPOSE Round down toward  $-\infty$ .

FORMAT  $y = \mathbf{floor}(x);$

INPUT  $x$   $N \times K$  matrix or N-dimensional array.

OUTPUT  $y$   $N \times K$  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  $x = 100 * \mathbf{rndn}(2,2);$

$$x = \begin{array}{cc} 77.68 & -14.10 \\ 4.73 & -158.88 \end{array}$$
$$f = \mathbf{floor}(x);$$
$$f = \begin{array}{cc} 77.00 & -15.00 \\ 4.00 & -159.00 \end{array}$$

SEE ALSO **ceil, round, trunc**

fmod

f

PURPOSE    Computes the floating-point remainder of  $x/y$ .

FORMAT     $r = \text{fmod}(x,y);$

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 11.)

EXAMPLE     $x = \text{seqa}(1.7,2.3,5)';$   
               $y = 2;$   
               $r = \text{fmod}(x,y);$

$x = \begin{bmatrix} 1.7 & 4 & 6.3 & 8.6 & 10.9 \end{bmatrix}$

$r = \begin{bmatrix} 1.7 & 0 & 0.3 & 0.6 & 0.9 \end{bmatrix}$

PURPOSE    Allows user to create one-line functions.

## fonts

---

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);`

`a = 50.265482`

## fonts

PURPOSE   Loads fonts to be used in the graph.

LIBRARY   `pgraph`

FORMAT    **fonts**(*str*);

INPUT    *str*            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.

For more information on how to select fonts within a text string, see PUBLICATION QUALITY GRAPHICS, Chapter [24](#).

SOURCE `pgraph.src`

SEE ALSO `title, xlabel, ylabel, zlabel`

## fopen

f

PURPOSE Opens a file.

FORMAT `f = fopen(filename,omode);`

INPUT *filename* string, name of file to open.  
*omode* string, file I/O mode. (See Remarks, below.)

OUTPUT *f* scalar, file handle.

PORTABILITY **UNIX**

Carriage return-linefeed conversion for files opened in text mode is unnecessary, because in UNIX 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 opened for reading and writing, or update, as follows:

## for

---

- 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-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.  
           *start*       scalar expression, the initial value of the counter.  
           *stop*        scalar expression, the final value of the counter.  
           *step*        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. They are converted to integers and 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

```

x = zeros(10, 5);
for i (1, rows(x), 1);
    for j (1, cols(x), 1);
        x[i,j] = i*j;
    endfor;
endfor;

```

Example 2

## format

---

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

```
li = 0;
x = rndn(100,1);
y = rndn(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    */typ*       literal, symbol type flag(s). Indicate which symbol types you are setting the output format for.

*/mat, /sa, /str*    Formatting parameters are maintained separately for matrices and arrays (*/mat*), string arrays (*/sa*), and strings (*/str*). You can specify more than one */typ* flag; the format will be set for all types indicated. If no */typ* flag is listed, **format** assumes */mat*.

*/fmted*    literal, enable formatting flag.

*/on, /off*        Enable/disable formatting. 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 1×1 string array.

*/mf*            literal, matrix row format flag.

*/m0*            no delimiters before or after rows when printing out matrices.

*/m1* or */mb1*    print 1 carriage return/line feed pair before each row of a matrix with more than 1 row.

*/m2* or */mb2*    print 2 carriage return/line feed pairs before each row of a matrix with more than 1 row.

*/m3* or */mb3*    print “Row 1”, “Row 2”... before each row of a matrix with more than one row.

*/ma1*            print 1 carriage return/line feed pair after each row of a matrix with more than 1 row.

*/ma2*            print 2 carriage return/line feed pairs after each row of a matrix with more than 1 row.

---

<b>/a1</b>	print 1 carriage return/line feed pair after each row of a matrix.
<b>/a2</b>	print 2 carriage return/line feed pairs after each row of a matrix.
<b>/b1</b>	print 1 carriage return/line feed pair before each row of a matrix.
<b>/b2</b>	print 2 carriage return/line feed pairs before each row of a matrix.
<b>/b3</b>	print “Row 1”, “Row 2”... before each row of a matrix.
<b>/jnt</b>	literal, matrix element format flag – controls justification, notation and trailing character.
<b>Right-Justified</b>	
<b>/rd</b>	Signed decimal number in the form <code>[[ - ]]#####.####</code> , where <code>#####</code> 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.
<b>/re</b>	Signed number in the form <code>[[ - ]]#.##E±###</code> , where <code>#</code> is one decimal digit, <code>##</code> is one or more decimal digits depending on the precision, and <code>###</code> is three decimal digits. If precision is 0, the form will be <code>[[ - ]]#E±###</code> with no decimal point printed.
<b>/ro</b>	This will give a format like <b>/rd</b> or <b>/re</b> depending on which is most compact for the number being printed. A format like <b>/re</b> will be used only if the exponent value is less than -4 or greater than the precision. If a <b>/re</b> format is used, a decimal point will always appear. The precision signifies the number of significant digits displayed.
<b>/rz</b>	This will give a format like <b>/rd</b> or <b>/re</b> depending on which is most compact for the number being printed. A format like <b>/re</b> 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  $[[ - ] #. ##E \pm ###$ , 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. 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

f

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:

	<code>format /rdn 1,3;</code>
<b>s</b>	The number will be followed immediately by a space character. This is the default.
<b>c</b>	The number will be followed immediately by a comma.
<b>t</b>	The number will be followed immediately by a tab character.
<b>n</b>	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:

-1.63533465	1.61350700	-1.06295179
0.26171282	0.27972294	-1.38937242
0.58891114	0.46812202	1.08805960

## format

---

This code:

```
format /m1 /rzs 1,10;  
print x;
```

produces:

```
-1.635334648 1.613507002 -1.062951787  
0.2617128159 0.2797229414 -1.389372421  
0.5889111366 0.4681220206 1.088059602
```

This code:

```
format /m3 /rdn 16,4;  
print x;
```

produces:

Row 1	-1.6353	1.6135	-1.0630
Row 2	0.2617	0.2797	-1.3894
Row 3	0.5889	0.4681	1.0881

This code:

```
format /m1 /ldn 16,4;  
print x;
```



produces:

-1.6353	1.6135	-1.0630
0.2617	0.2797	-1.3894
0.5889	0.4681	1.0881

This code:

```
format /ml /res 12,4;
print x;
```

produces:

-1.6353E+000	1.6135E+000	-1.0630E+000
2.6171E-001	2.7972E-001	-1.3894E+000
5.8891E-001	4.6812E-001	1.0881E+000

SEE ALSO **formatcv**, **formatnv**, **print**, **output**

## formatcv

**PURPOSE** Sets the character data format used by **printfmt**.

**FORMAT** *oldfmt* = **formatcv**(*newfmt*);

**INPUT** *newfmt* 1×3 vector, the new format specification.

**OUTPUT** *oldfmt* 1×3 vector, the old format specification.

**REMARKS** See **printfm** for details on the format vector.

## formatnv

---

**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:

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** `newfmt` 1×3 vector, the new format specification.

**OUTPUT** `oldfmt` 1×3 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("*. *lf" ~ 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** `formatcv`, `printfm`, `printfmt`

## fputs

**PURPOSE** Writes strings to a file.

**FORMAT** `numl = fputs(f,sa);`

**INPUT** *f* scalar, file handle of a file opened with **fopen**.  
*sa* string or string array.

**OUTPUT** *numl* scalar, the number of lines written to the file.

## fputst

---

PORTABILITY    **UNIX**

Carriage return-linefeed conversion for files opened in text mode is unnecessary, because in **UNIX** a newline is simply a linefeed.

**REMARKS**    **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 *numl* 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.

**SEE ALSO**    **fputst**, **fopen**

## fputst

**PURPOSE**    Writes strings to a file.

**FORMAT**    *numl* = **fputst** (*f*,*sa*);

**INPUT**    *f*            scalar, file handle of a file opened with **fopen**.  
          *sa*            string or string array.

**OUTPUT**    *numl*          scalar, the number of lines written to the file.

PORTABILITY    **UNIX**

Carriage return-linefeed conversion for files opened in text mode is unnecessary, because in **UNIX** a newline is simply a linefeed.

**REMARKS** **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

f

**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 <b>fopen</b> .
<i>offs</i>	scalar, offset (in bytes).
<i>base</i>	scalar, base position.
<b>0</b>	beginning of file.
<b>1</b>	current position of file pointer.
<b>2</b>	end of file.

**OUTPUT** *ret* scalar, 0 if successful, 1 if not.

**PORTABILITY** **UNIX**

Carriage return-linefeed conversion for files opened in text mode is unnecessary, because in UNIX 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 *s* 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.

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 *f* scalar, file handle of a file opened with **fopen**.

OUTPUT *pos* 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.

## ftocv

---

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**  $y = \text{ftocv}(x, \text{field}, \text{prec});$

**INPUT**  $x$   $N \times K$  matrix containing numeric data to be converted.  
 $\text{field}$  scalar, minimum field width.  
 $\text{prec}$  scalar, the numbers created will have  $\text{prec}$  places after the decimal point.

**OUTPUT**  $y$   $N \times K$  matrix containing the decimal character equivalent of the corresponding elements in  $x$  in the format defined by  $\text{field}$  and  $\text{prec}$ .

**REMARKS** If a number is narrower than  $\text{field}$ , it will be padded on the left with zeros.  
If  $\text{prec} = 0$ , the decimal point will be suppressed.

**EXAMPLE**  $y = \text{seqa}(6, 1, 5);$   
 $x = 0 \ \$+ \text{"cat"} \ \$+ \text{ftocv}(y, 2, 0);$

```
      cat06
      cat07
x =  cat08
      cat09
      cat10
```



Notice that the ( 0 \$+ ) above was necessary to force the type of the result to matrix because the string constant “**cat**” would be of type string. The left operand in an expression containing a \$+ operator controls the type of the result.

SEE ALSO **ftos**

ftos

f

**PURPOSE** Converts a scalar into a string containing the decimal character representation of that number.

**FORMAT**  $y = \mathbf{ftos}(x, fmat, field, prec);$

**INPUT**  $x$  scalar, the number to be converted.

$fmat$  string, the format string to control the conversion.

$field$  scalar or 2×1 vector, the minimum field width. If  $field$  is 2×1, it specifies separate field widths for the real and imaginary parts of  $x$ .

$prec$  scalar or 2×1 vector, the number of places following the decimal point. If  $prec$  is 2×1, it specifies separate precisions for the real and imaginary parts of  $x$ .

**OUTPUT**  $y$  string containing the decimal character equivalent of  $x$  in the format specified.

**REMARKS** The format string corresponds to the **format** /*jnt* (justification, notation, trailing character) slash parameter as follows:

```
/rdn    ‘%*.1f’  
/ren    ‘%*.1E’  
/ron    ‘%#*.1G’  
/rzn    ‘%*.1G’  
  
/ldn    ‘%- *.1f’  
/len    ‘%- *.1E’  
/lon    ‘%-# *.1G’  
/lzn    ‘%- *.1G’
```

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:

```
‘%*.1f,’    right-justified decimal followed by a comma.  
‘%-*.1s ’   left-justified string followed by a space.  
‘%*.1f’     right-justified decimal followed by nothing.
```

You can embed the format specification in the middle of other text:

```
"Time: %*.1f seconds."
```

If you want the beginning of the field padded with zeros, then put a “0” before the first “\*” in the format string:

```
‘%0*.1f’    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 = "%*.*lE";
xx = ftos(x,fmat,field,prec);

field = 7;
prec = 2;
```

## ftostrC

---

```
fmat = "%*.*lf seconds";
s1 = ftos(x,fmat,field,prec);
s2 = ftos(y,fmat,field,prec);

field = 1;
prec = 2;
fmat = "The maximum resistance is %*.*lf ohms.";
om = ftos(x,fmat,field,prec);
```

The results:

```
zz = "5"
xx = "9.2985743532E+02"
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**  $sa = \mathbf{ftostrC}(x,fmt);$

**INPUT**  $x$   $N \times K$  matrix, real or complex.  
 $fmt$   $K \times 1$ ,  $1 \times K$  or  $1 \times 1$  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:

```

          -0.166      1.05565441
          -1.590      -0.79283296
           0.130      -1.84886957
           0.789      0.86089687

( 0.000, -0.407) ( 1.02745044, 0.000000000)
( 0.000, -1.261) ( 0.000000000, -0.89041168)
( 0.361, 0.000) ( 0.000000000, -1.35973143)
( 0.888, 0.000) ( 0.92784529, 0.000000000)

```

SEE ALSO **strtof, strtocplx**

### gamma

**PURPOSE** Returns the value of the gamma function.

**FORMAT** `y = gamma(x);`

**INPUT** `x` N×K matrix or N-dimensional array.

**OUTPUT** `y` N×K matrix or N-dimensional array.

**REMARKS** For each element of `x` this function returns the integral

$$\int_0^{\infty} t^{(x-1)} e^{-t} dt$$

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);`

`y = 1.32934`

**SEE ALSO** `cdfchic, cdfbeta, cdfcfc, cdfn, cdfnc, cdftc, erf, erfc, lnfact`

### gamma

PURPOSE Computes the inverse incomplete gamma function.

FORMAT  $x = \text{gammaii}(a, p);$

INPUT  $a$  M×N matrix, exponents.  
 $p$  K×L matrix, E×E conformable with  $a$ , incomplete gamma values.

OUTPUT  $x$  max(M,K) by max(N,L) matrix, abscissae.

SOURCE cdfchii.src

GLOBALS `_ginvinc, __macheps`

g

## 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);`

## gdaAppend

---

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.
	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	<p>This command appends the data contained in <i>x</i> to the variable <i>varname</i> in <i>filename</i>. Both <i>x</i> and the variable referenced by <i>varname</i> must be the same data type, and they must both contain the same number of columns.</p> <p>Because <b>gdaAppend</b> 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 <b>gdaPack</b> to pack the data in a GDA, so it contains no empty bytes.</p>	
EXAMPLE	<pre>x = rndn(100,50); ret = gdaCreate("myfile.gda",1); ret = gdaWrite("myfile.gda",x,"x1");</pre>	



```
y = rndn(25,50);
ret = gdaAppend("myfile.gda",y,"x1");
```

This example adds 25\*50=1250 elements to **x1**, making it a 125×50 matrix.

SEE ALSO **gdaWriteSome**, **gdaUpdate**, **gdaWrite**

## gdaCreate

g

**PURPOSE** Creates a **GAUSS** Data Archive.

**FORMAT** *ret* = **gdaCreate**(*filename*,*overwrite*);

**INPUT** *filename* string, name of data file to create.  
*overwrite* scalar, one of the following:  
**0** error out if file already exists.  
**1** overwrite file if it already exists.

**OUTPUT** *ret* 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.

**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);`

## gdaDStat

---

SEE ALSO **gdaWrite**

### gdaDStat

**PURPOSE** Computes descriptive statistics on multiple N×1 variables in a **GAUSS** Data Archive.

**FORMAT** *dout* = **gdaDStat**(*dc0*,*filename*,*vars*);

**INPUT** *dc0* an instance of a **dstatmtControl** structure with the following members:

<i>dc0.altnames</i>	K×1 string array of alternate variable names for the output. Default = “”.
<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: <b>0</b> There are no missing values (fastest). <b>1</b> Listwise deletion, drop a row if any missings occur in it. <b>2</b> Pairwise deletion. Default = 0.
<i>dc0.output</i>	scalar, one of the following: <b>0</b> Do not print output table. <b>1</b> 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> .

*filename* string, name of data file.

*vars* K×1 string array, names of variables  
 - or -  
 K×1 vector, indices of variables.

OUTPUT *dout* an instance of a **dstatmtOut** structure with the following members:

<i>dout.vnames</i>	K×1 string array, the names of the variables used in the statistics.
<i>dout.mean</i>	K×1 vector, means.
<i>dout.var</i>	K×1 vector, variance.
<i>dout.std</i>	K×1 vector, standard deviation.
<i>dout.min</i>	K×1 vector, minima.
<i>dout.max</i>	K×1 vector, maxima.
<i>dout.valid</i>	K×1 vector, the number of valid cases.
<i>dout.missing</i>	K×1 vector, the number of missing cases.
<i>dout.errcode</i>	scalar, error code, 0 if successful, or one of the following: <ol style="list-style-type: none"> <li>1 No GDA indicated.</li> <li>4 Not implemented for complex data.</li> <li>5 Variable must be type matrix.</li> <li>6 Too many variables specified.</li> <li>7 Too many missings - no data left after packing.</li> <li>8 Name variable wrong size.</li> <li>9 <b>altnames</b> member of <b>dstatmtControl</b> structure wrong size.</li> <li>11 Data read error.</li> </ol>

REMARKS The variables referenced by *vars* must all be N×1.

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.

## gdaDStatMat

---

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;`  
  
              `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`

SEE ALSO     **gdaDStatMat**, **dstatmtControlCreate**

## gdaDStatMat

PURPOSE     Computes descriptive statistics on a selection of columns in a variable in a **GAUSS** Data Archive.

FORMAT     `dout = gdaDStatMat(dc0,filename,var,colind,vnamevar);`

INPUT     *dc0*            an instance of a **dstatmtControl** structure with the following members:

<i>dc0.altnames</i>	K×1 string array of alternate variable names for the output. Default = "".
<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: <b>0</b> There are no missing values (fastest). <b>1</b> Listwise deletion, drop a row if any missings occur in it. <b>2</b> Pairwise deletion. Default = 0.
	<i>dc0.output</i>	scalar, one of the following: <b>0</b> Do not print output table. <b>1</b> 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>var</i>	string, name of variable - or - scalar, index of variable.
	<i>colind</i>	K×1 vector, indices of columns in variable to use.
	<i>vnamevar</i>	string, name of variable containing names for output - or - scalar, index of variable containing names for output.
OUTPUT	<i>dc0</i>	an instance of a <b>dstatmtOut</b> structure with the following members: <i>dout.vnames</i> K×1 string array, the names of the variables used in the statistics. <i>dout.mean</i> K×1 vector, means. <i>dout.var</i> K×1 vector, variance. <i>dout.std</i> K×1 vector, standard deviation. <i>dout.min</i> K×1 vector, minima.

<i>dout.max</i>	K×1 vector, maxima.
<i>dout.valid</i>	K×1 vector, the number of valid cases.
<i>dout.missing</i>	K×1 vector, the number of missing cases.
<i>dout.errcode</i>	scalar, error code, 0 if successful, otherwise one of the following: <b>1</b> No GDA indicated. <b>3</b> Variable must be N×1. <b>4</b> Not implemented for complex data. <b>5</b> Variable must be type matrix. <b>7</b> Too many missings - no data left after packing. <b>9</b> <b>altnames</b> member of <b>dstatmtControl</b> structure wrong size. <b>11</b> Data read error.

REMARKS     Set *colind* to a scalar 0 to use all of the columns in *var*.

*vnamevar* must either reference an M×1 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 M×1 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 **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;

dc0 = dstatmtControlCreate;
var = 3;
index = { 1,3,4,7 };
dout = gdaDStatMat(dc0,"myfile.gda",var,index,"");
```

This example computes descriptive statistics on the first, third, fourth and seventh columns of the third variable in `myfile.gda`, generating names for the output automatically.

SOURCE `gdadstat.src`

SEE ALSO `gdaDStat`, `dstatmtControlCreate`

## gdaGetIndex

g

PURPOSE Gets the index of a variable in a **GAUSS** Data Archive.

FORMAT `ind = gdaGetIndex(filename, varname);`

INPUT *filename* string, name of data file.  
*varname* string, name of variable in the GDA.

OUTPUT *ind* 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.
- 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`

## gdaGetNames

---

### gdaGetName

- PURPOSE** Gets the name of a variable in a **GAUSS** Data Archive.
- FORMAT** `varname = gdaGetName(filename, varind);`
- INPUT** *filename* string, name of data file.  
*varind* scalar, index of variable in the GDA.
- OUTPUT** *varname* 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** *filename* string, name of data file.



OUTPUT     *varnames*   N×1 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**

g

## gdaGetOrders

PURPOSE     Gets the orders of a variable in a **GAUSS** Data Archive.

FORMAT     `ord = gdaGetOrders(filename, varname);`

INPUT     *filename*   string, name of data file.  
             *varname*   string, name of variable in the GDA.

OUTPUT     *ord*           M×1 vector, orders of the variable in the GDA.

REMARKS     If the specified variable is a matrix or string array, then *ord* will be a 2×1 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 N×1 vector containing the sizes of each dimension.

## gdaGetType

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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.

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    *filename*    string, name of data file.  
          *varname*    string, name of variable in the GDA.

OUTPUT    *vartype*    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**

g

## gdaGetTypes

PURPOSE     Gets the types of all the variables in a **GAUSS** Data Archive.

FORMAT     `vartypes = gdaGetTypes(filename);`

INPUT     *filename*     string, name of data file.

OUTPUT     *vartypes*     N×1 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:

## gdaGetVarInfo

---

- 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     *filename*     string, name of data file.

OUTPUT     *vtab*     N×1 array of **gdavartable** structures, where N is the number of variables in *filename*, containing the following members:

*vtab*[*i*].**name**     string, name of variable.

*vtab*[*i*].**type**     scalar, type of variable.

*vtab*[*i*].**orders**     M×1 vector or scalar, orders of the variable.

REMARKS     The size of *vtab.orders* is dependent on the type of the variable as follows:

Variable Type	<i>vtab.orders</i>
array	M×1 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	2×1 vector containing the rows and columns of the matrix, respectively.
string	scalar containing the length of string, excluding the null terminating byte.
string array	2×1 vector containing the rows and columns of the string array, respectively.

*vtab.type* may contain any of the following:

- 6**    matrix
- 13**   string
- 15**   string array
- 21**   array

EXAMPLE    `#include gdafns.sdf`

`struct gdavartable vtab;`

`vtab = gdaGetVarInfo("myfile.gda");`

SOURCE    `gdafns.src`

SEE ALSO    **gdaReportVarInfo**, **gdaGetNames**, **gdaGetTypes**, **gdaGetOrders**

PURPOSE    Checks to see if a variable in a **GAUSS** Data Archive is complex.

FORMAT     `y = gdaIsCplx(filename, varname);`

INPUT       *filename*    string, name of data file.

## gdaLoad

---

*varname*    string, name of variable in the GDA.

OUTPUT    *y*            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    *filename*    string, name of data file.

*create*    scalar, create flag:

- 0**    do not create any new variables in the workspace.
- 1**    create new variables in the workspace.

*modify*    scalar, modify flag:

- 0**    do not modify any variables in the workspace.
- 1**    if the name of a variable in the data file matches the name of a variable already in the workspace, modify that variable.

*rename*    scalar, rename flag:

		<ul style="list-style-type: none"> <li><b>0</b> do not rename a variable retrieved from the data file when copying it into the workspace.</li> <li><b>1</b> 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.</li> </ul>
	<i>ftypes</i>	<p>scalar, type force flag:</p> <ul style="list-style-type: none"> <li><b>0</b> do not force a type change on any variables in the workspace when modifying.</li> <li><b>1</b> 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, <b>gdaLoad</b> will follow regular type change rules. The types of sparse matrix and structure variables will NOT be changed.</li> </ul>
	<i>errh</i>	<p>scalar, controls the error handling of <b>gdaLoad</b>:</p> <ul style="list-style-type: none"> <li><b>0</b> skip operations that cannot be performed, without setting an error return.</li> <li><b>1</b> return an error code if operations are skipped.</li> <li><b>2</b> terminate program if operations are skipped.</li> </ul>
	<i>report</i>	<p>scalar, controls reporting:</p> <ul style="list-style-type: none"> <li><b>0</b> no reporting.</li> <li><b>1</b> report only name changes and operations that could not be performed.</li> <li><b>2</b> report type changes, name changes, and operations that could not be performed.</li> <li><b>3</b> report everything.</li> </ul>
OUTPUT	<i>ret</i>	<p>scalar, return code, 0 if successful, otherwise one of the following error codes:</p> <ul style="list-style-type: none"> <li><b>4</b> File read error.</li> <li><b>5</b> Invalid file type.</li> <li><b>10</b> File contains no variables.</li> <li><b>14</b> File too large to be read on current platform.</li> <li><b>24</b> Variables skipped.</li> <li><b>26</b> Cannot add structure definition.</li> </ul>

**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 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**

g

gdaPack

PURPOSE     Packs the data in a **GAUSS** Data Archive, removing all empty bytes and truncating the file.

FORMAT     `ret = gdaPack(filename);`

INPUT       *filename*     string, name of data file.

OUTPUT      *ret*           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.

## gdaRead

---

EXAMPLE     `ret = gdaPack("myfile.gda");`

SEE ALSO     **gdaUpdate, gdaWrite**

## gdaRead

PURPOSE     Gets a variable from a **GAUSS** Data Archive.

FORMAT     `y = gdaRead(filename, varname);`

INPUT     *filename*     string, name of data file.  
            *varname*     string, name of variable in the GDA.

OUTPUT     *y*             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** *filename* string, name of data file.  
*varind* scalar, index of variable in the GDA.

**OUTPUT** *y* matrix, array, string or string array, variable data.

**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);`

## gdaReadSome

---

INPUT    *filename*    string, name of data file.

*varname*    string, name of variable in the GDA.

*index*      scalar or  $N \times 1$  vector, index into variable where read is to begin.

*orders*     scalar or  $K \times 1$  vector, orders of object to output.

OUTPUT   *y*            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  $10 \times 5$  matrix, an *index* of 42 would indicate the 42<sup>nd</sup> 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  $N \times 1$  vector, then *N* must equal the number of dimensions in the variable referenced by *varname*.

If *orders* is a  $K \times 1$  vector, then *y* will be a *K*-dimensional object. If *orders* is a scalar *r*, then *y* will be an  $r \times 1$  column vector. To specify a  $1 \times r$  row vector, set *output* = { 1, *r* }.

If the variable referenced by *varname* is numeric (a matrix or array) and *orders* is a scalar or  $2 \times 1$  vector, then *y* will of type matrix. If the variable is numeric and *orders* is an  $N \times 1$  vector where  $N > 2$ , then *y* will be of type array.

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\*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       *filename*    string, name of data file.  
             *varname*    string, name of sparse matrix variable in the GDA.

OUTPUT      *sm*            sparse matrix.

## gdaReadStruct

---

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.

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 *filename* string, name of data file.  
*varname* string, name of structure instance in the GDA.  
*structure\_type* string, structure type.

OUTPUT *instance* instance of the structure.  
*retcode* 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 = rndn(500,25);
msw.a = areshape(rndn(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

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 *filename* string, name of data file.

OUTPUT *vinfo* N×1 string array containing variable information.

## gdaSave

---

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	Orders
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 N×K string array, names of variables in the workspace to include or exclude.
	<i>exclude</i>	scalar, include/exclude flag: <ul style="list-style-type: none"> <li><b>0</b> include all variables contained in <i>varnames</i>.</li> <li><b>1</b> exclude all variables contained in <i>varnames</i>.</li> </ul>
	<i>overwrite</i>	scalar, controls the overwriting of the file and variables in the file: <ul style="list-style-type: none"> <li><b>0</b> if file exists, return with an error code.</li> <li><b>1</b> if file exists, overwrite completely.</li> <li><b>2</b> if file exists, append to file, appending to variable names if necessary to avoid name conflicts.</li> <li><b>3</b> if file exists, update file. When a name conflict occurs, update the existing variable in the file with the new variable.</li> </ul>
	<i>report</i>	scalar, controls reporting: <ul style="list-style-type: none"> <li><b>0</b> no reporting.</li> <li><b>1</b> report only name changes (note that name changes occur only when <i>overwrite</i> is set to 2).</li> <li><b>3</b> report everything.</li> </ul>
OUTPUT	<i>ret</i>	scalar, return code, 0 if successful, otherwise one of the following error codes: <ul style="list-style-type: none"> <li><b>1</b> Null file name.</li> <li><b>3</b> File write error.</li> <li><b>4</b> File read error.</li> <li><b>5</b> Invalid file type.</li> <li><b>6</b> File exists and <i>overwrite</i> set to 0.</li> <li><b>7</b> Cannot create file.</li> <li><b>14</b> File too large to be read on current platform.</li> <li><b>16</b> Cannot write to GDA – version outdated.</li> <li><b>17</b> Type mismatch.</li> </ul>

REMARKS Only initialized variables are written to the GDA with **gdaSave**.

## gdaUpdate

---

If *varnames* is a null string, 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.

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    *filename*    string, name of data file.  
           *x*            matrix, array, string or string array, data.  
           *varname*    string, variable name.

OUTPUT    *ret*            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.

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,

## gdaUpdateAndPack

---

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      *filename*    string, name of data file.  
              *x*                matrix, array, string or string array, data.  
              *varname*    string, variable name.

OUTPUT    *ret*                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.

**12** File truncate error.

**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*. **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.

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.

g

**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** *filename* string, name of data file.

**OUTPUT** *nvars* scalar, the number of variables in *filename*.

**EXAMPLE** *nvars* = **gdaVars**("myfile.gda");

## gdaWrite

---

SOURCE    `gdafns.src`

SEE ALSO    `gdaReportVarInfo`, `gdaGetNames`,

## gdaWrite

PURPOSE    Writes a variable to a **GAUSS** Data Archive.

FORMAT    `ret = gdaWrite(filename,x,varname);`

INPUT    *filename*    string, name of data file.  
          *x*            matrix, array, string or string array, data to write to the GDA.  
          *varname*    string, variable name.

OUTPUT    *ret*            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.

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**    *filename*    string, name of data file.  
             *x*            matrix, array, string or string array, data to write to the GDA.  
             *varname*    string, variable name.

**OUTPUT**    *ret*            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 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

g

## gdaWriteSome

---

Windows XP than the 32-bit file write commands that were used for binary writes in **GAUSS** 6.0. **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     *filename*     string, name of data file.  
              *x*                 matrix, array, string or string array, data.  
              *varname*     string, variable name.  
              *index*         scalar or N×1 vector, index into variable where new data is to be written.

OUTPUT     *ret*               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 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*<sup>th</sup> element of the variable. Thus if *varname* references a 10×5 matrix, an *index* of 42 would indicate the 42<sup>nd</sup> 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 N×1 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 N×K matrix, then *x* may be any L×M 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).

## getarray

---

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\*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 *a* N-dimensional array.

	<i>loc</i>	M×1 vector of indices into the array to locate the subarray of interest, where $1 \leq M \leq N$ .
OUTPUT	<i>y</i>	[N-M]-dimensional subarray or scalar.
REMARKS		If N-M>0, <b>getarray</b> will return an array of [N-M] dimensions, otherwise, if N-M=0, it will return a scalar.
EXAMPLE		<pre>a = seqa(1,1,720); a = areshape(a,2 3 4 5 6); loc = { 2,1 }; y = getarray(a,loc);</pre> <p><b>y</b> will be a 4×5×6 array of sequential values, beginning at [1,1,1] with 361, and ending at [4,5,6] with 480.</p>
SEE ALSO		<b>getmatrix</b>

g

## getdims

PURPOSE		Gets the number of dimensions in an array.
FORMAT	<i>y</i>	<b>getdims(a);</b>
INPUT	<i>a</i>	N-dimensional array.
OUTPUT	<i>y</i>	scalar, the number of dimensions in the array.
EXAMPLE		<pre>a = arrayinit(3 4 5 6 7 2,0); dims = getdims(a);</pre> <p style="text-align: center;">dims = 6</p>

## getf

---

SEE ALSO **getorders**

## getf

**PURPOSE** Loads an ASCII or binary file into a string.

**FORMAT** `y = getf(filename,mode);`

**INPUT** *filename* string, any valid file name.  
*mode* scalar 1 or 0 which determines if the file is to be loaded in ASCII mode (0) or binary mode (1).

**OUTPUT** *y* 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** Create a file `examp.e` containing the following program:

```
library pgraph;  
graphset;  
x = seqa(0,0.1,100);  
y = sin(x);  
xy(x,y);
```

Then execute the following:

```
y = getf("examp.e",0);  
  
print y;
```

This produces:

```
library pgraph;  
graphset;  
x = seqa(0,0.1,100);  
y = sin(x);  
xy(x,y);
```



SEE ALSO    **load, save, let, con**

getmatrix

PURPOSE	Gets a contiguous matrix from an N-dimensional array.	
FORMAT	$y = \text{getmatrix}(a, loc);$	
INPUT	$a$	N-dimensional array.
	$loc$	M×1 vector of indices into the array to locate the matrix of interest, where M equals N, N-1 or N-2.
OUTPUT	$y$	K×L or 1×L 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	Inputting an N×1 locator vector will return a scalar, an (N-1)×1 locator vector will return a 1×L matrix, and an (N-2)×1 locator vector will return a K×L matrix.	

## getmatrix4D

---

EXAMPLE    `a = seqa(1,1,120);`  
             `a = areshape(a,2|3|4|5);`  
             `loc = { 1,2 };`  
             `y = getmatrix(a,loc);`

$y =$

21	22	23	24	25
26	27	28	29	30
31	32	33	34	35
36	37	38	39	40

SEE ALSO    `getarray`, `getmatrix4D`

## getmatrix4D

PURPOSE    Gets a contiguous matrix from a 4-dimensional array.

FORMAT    `y = getmatrix4D(a,i1,i2);`

INPUT    *a*            4-dimensional array.  
          *i1*           scalar, index into the slowest moving dimension of the array.  
          *i2*           scalar, index into the second slowest moving dimension of the array.

OUTPUT    *y*            K×L 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     `a = seqa(1,1,120);`  
              `a = areshape(a,2|3|4|5);`  
              `y = getmatrix4D(a,2,3);`

                 101 102 103 104 105  
                 106 107 108 109 110  
      `y =`       111 112 113 114 115  
                 116 117 118 119 120

SEE ALSO     `getmatrix`, `getscalar4D`, `getarray`

g

getname

PURPOSE     Returns a column vector containing the names of the variables in a **GAUSS** data set.

FORMAT     `y = getname(dset);`

INPUT       *dset*       string specifying the name of the data set from which the function will obtain the variable names.

OUTPUT      *y*           N×1 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("olsdat");`  
              `format 8,8;`  
              `print $y;`  
  
              produces:

## getnamef

---

```
TIME
DIST
TEMP
FRICT
```

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** `f` scalar, file handle of an open data set

**OUTPUT** `y` N×1 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**

```
open f = olsdat for read;
y = getnamef(f);
t = vartypef(f);
print y;
```

produces:



time  
dist  
temp  
frict

The above example assumes that the data set `olsdat` contains the variables: **time**, **dist**, **temp**, **frict**.

Note the use of **vartypef** to determine the types of these variables.

SEE ALSO **getname**, **indcv**, **vartypef**

g

getNextTradingDay

- PURPOSE Returns the next trading day.
- FORMAT `n = getNextTradingDay(a);`
- INPUT `a` scalar, date in DT scalar format.
- OUTPUT `n` 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**

## getnr

### getNextWeekDay

**PURPOSE** Returns the next day that is not on a weekend.

**FORMAT**  $n = \text{getNextWeekDay}(a);$

**INPUT**  $a$  scalar, date in DT scalar format.

**OUTPUT**  $n$  scalar, next week day in DT scalar format.

**SOURCE** finutils.src

**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**  $nr = \text{getnr}(nsets, ncols);$

**INPUT**  $nsets$  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.

$ncols$  scalar, columns in the data file.

**OUTPUT**  $nr$  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

## g

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 *nsets* 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.

*ncols* scalar, columns in the data file.

*row* scalar, if row is greater than 0, *nr* will be set to *row*.

*rowfac* scalar, *nr* 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).

*maxv* scalar, the largest number of elements allowed in any one matrix.

OUTPUT *nr* scalar, number of rows **readr** should read per iteration of the read loop.

SOURCE `gaussmt.src`

## getpath

---

### getorders

**PURPOSE** Gets the vector of orders corresponding to an array.

**FORMAT**  $y = \text{getorders}(a);$

**INPUT**  $a$  N-dimensional array.

**OUTPUT**  $y$  N×1 vector of orders, the sizes of the dimensions of the array.

**EXAMPLE**  $a = \text{arrayalloc}(7|6|5|4|3,0);$   
 $\text{orders} = \text{getorders}(a);$

```

              7
              6
orders = 5
              4
              3
```

**SEE ALSO** [\*\*getdims\*\*](#)

### getpath

**PURPOSE** Returns an expanded filename including the drive and path.

**FORMAT**  $fname = \text{getpath}(pname);$

**INPUT**  $pname$  string, partial filename with only partial or missing path information.

**OUTPUT**  $fname$  string, filename with full drive and path.

REMARKS This function handles relative path references.

EXAMPLE `y = getpath("temp.e");  
print y;`

produces:

`/gauss/temp.e`

assuming that `/gauss` is the current directory.

SOURCE `getpath.src`

g

## getPreviousTradingDay

PURPOSE Returns the previous trading day.

FORMAT `n = getPreviousTradingDay(a);`

INPUT *a* scalar, date in DT scalar format.

OUTPUT *n* 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**  $n = \text{getPreviousWeekDay}(a);$

**INPUT**  $a$  scalar, date in DT scalar format.

**OUTPUT**  $n$  scalar, previous week day in DT scalar format.

**SOURCE** finutils.src

**SEE ALSO** getNextWeekDay

### getRow

**PURPOSE** Returns a specified row from a matrix.

**FORMAT**  $y = \text{getRow}(a, row);$

**INPUT**  $a$   $N \times K$  matrix  
 $row$  The row of the matrix to extract.

**OUTPUT**  $y$  A  $1 \times K$  row vector.

**REMARKS** **getRow** is designed to give an alternative access to rows in a matrix than indexing the matrix by brackets.

**EXAMPLE**  $a = \text{rndn}(10, 10);$   
 $y = \text{getRow}(a, 3);$

SEE ALSO **geTrRow**

## getscalar3D

**PURPOSE** Gets a scalar from a 3-dimensional array.

**FORMAT** `y = getscalar3D(a,i1,i2,i3);`

<b>INPUT</b>	<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** *y* 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**

```

a = seqa(1,1,24);
a = areshape(a,2|3|4);
y = getscalar3D(a,1,3,2);

```

`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);`

**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** *y* 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);
```

`y = 50`

**SEE ALSO** **getmatrix**, **getscalar3D**, **getarray**



## getTrRow

PURPOSE	Transposes a matrix and then returns a single row from it.	
FORMAT	$y = \text{getTrRow}(a, row);$	
INPUT	$a$	$N \times K$ matrix
	$row$	The row of the matrix to extract.
OUTPUT	$y$	A $1 \times K$ row vector.
REMARKS	<b>getRow</b> is designed to give an alternative access to rows in a matrix than indexing the matrix by brackets.	
EXAMPLE	$a = \text{rndn}(10, 10);$ $y = \text{getTrRow}(a, 3);$	
SEE ALSO	<b>getRow</b>	

g

## getwind

PURPOSE	Retrieve the current graphic panel number.	
LIBRARY	pgraph	
FORMAT	$n = \text{getwind};$	
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.	

## gosub

---

SOURCE    `pwindow.src`

SEE ALSO    **endwind**, **begwind**, **window**, **setwind**, **nextwind**

## gosub

PURPOSE    Causes a branch to a subroutine.

FORMAT    **gosub** *label*;  
          .  
          .  
          .  
          *label*:  
          .  
          .  
          .  
          **return**;

REMARKS    For multi-line recursive user-defined functions, see PROCEDURES AND KEYWORDS, Chapter [12](#).

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 1-32 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 12.)

**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**.
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**

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 = seqa(.1,.1,5);  
n = { 1 2 3 };  
goto fip;  
print x;  
end;`

`fip:  
    print n;`

produces:

## gradMT

---

1.00000000

2.00000000

3.00000000

SEE ALSO **gosub, if**

## gradMT

**PURPOSE** Computes numerical gradient.

**INCLUDE** `optim.sdf`

**FORMAT** `g = gradMT(&fct,par1,data1);`

**INPUT**

<b>&amp;fct</b>	scalar, pointer to procedure returning either N×1 vector or 1×1 scalar.
<b>par1</b>	an instance of structure of type <b>PV</b> containing parameter vector at which gradient is to be evaluated.
<b>data1</b>	structure of type <b>DS</b> containing any data needed by <i>fct</i> .

**OUTPUT** `g` N×K Jacobian or 1×K gradient.

**REMARKS** *par1* must be created using the **pvpPack** 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

g

gradMTm

PURPOSE Computes numerical gradient with mask.

INCLUDE optim.sdf

FORMAT  $g = \text{gradMTm}(\&fct, par1, data1, mask);$

INPUT	$\&fct$	scalar, pointer to procedure returning either $N \times 1$ vector or $1 \times 1$ scalar.
	$par1$	an instance of structure of type <b>PV</b> containing parameter vector at which gradient is to be evaluated.
	$data1$	structure of type <b>DS</b> containing any data needed by $fct$ .
	$mask$	$K \times 1$ matrix, elements in $g$ corresponding to elements of $mask$ set to zero are not computed, otherwise they are computed.

OUTPUT  $g$   $N \times K$  Jacobian or  $1 \times K$  gradient.

REMARKS  $par1$  must be created using the **pvPack** procedures.

EXAMPLE `#include optim.sdf`

## gradp

---

```
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

## gradp

**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.

**FORMAT**  $g = \text{gradp}(\&f, x0);$

**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);  
    retp( exp(x.*b) );  
endp;
```

*x0*            K×1 vector of points at which to compute gradient.

OUTPUT      *g*            N×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×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.



EXAMPLE      

```
proc myfunc(x);  
    retp( x.*2 .* exp( x.*x./3 ) );  
endp;
```

```
x0 = 2.5|3.0|3.5;  
y = gradp(&myfunc,x0);
```

```
      82.98901842      0.00000000      0.00000000  
y =   0.00000000  281.19752975      0.00000000  
      0.00000000      0.00000000  1087.95414117
```

It is a 3×3 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**

### graphprt

PURPOSE Controls automatic printer hardcopy and conversion file output.

LIBRARY pgraph

FORMAT **graphprt**(*str*);

INPUT *str* 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.

**-P** print graph.  
**-PO=*c*** set print orientation:  
    **L** landscape.  
    **P** portrait.  
**-C=*n*** convert to another file format:  
    **1** Encapsulated PostScript file.  
    **3** HPGL Plotter file.  
    **5** BMP (Windows Bitmap).  
    **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  $n$  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 24.

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 */
```

## graphset

---

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

PURPOSE    Reset graphics global variables to default values.

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`

hasimag

g

PURPOSE Tests whether the imaginary part of a complex matrix is negligible.

FORMAT `y = hasimag(x);`

INPUT `x`  $N \times K$  matrix.

OUTPUT `y` scalar, 1 if the imaginary part of `x` has any nonzero elements, 0 if it consists entirely of 0's.

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.

**iscplx** is useful as a preliminary check because for large matrices it is much faster than **hasimag**.

EXAMPLE `x = { 1 2 3i,`

## header

---

```
      4-i 5 6i,  
      7 8i 9 };  
  
y = hasimag(x);  
  
y = 1.0000000
```

SEE ALSO **iscplx**

## header

**PURPOSE** Prints a header for a report.

**FORMAT** **header**(*prcnm*,*dataset*,*ver*);

<b>INPUT</b>	<i>prcnm</i>	string, name of procedure that calls <b>header</b> .
	<i>dataset</i>	string, name of data set.
	<i>ver</i>	2×1 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 ( <b>__??_ver</b> ) associated with the module within which header is called. This argument will be ignored if set to 0.

<b>GLOBAL INPUT</b>	<b>__header</b>	string, containing one or more of the following letters:
	<b>t</b>	title is to be printed
	<b>l</b>	lines are to bracket the title
	<b>d</b>	a date and time is to be printed
	<b>v</b>	version number of program is to be printed

---

**f** file name being analyzed is to be printed

**\_\_title** string, title for header.

SOURCE gauss.src

## headermt

PURPOSE Prints a header for a report.

FORMAT **headermt**(*prcnm*,*dataset*,*ver*,*header*,*title*);

INPUT *prcnm* string, name of procedure that calls **header**.

*dataset* string, name of data set.

*ver* 2×1 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 (**\_\_??\_ver**) associated with the module within which header is called. This argument will be ignored if set to 0.

*header* string, containing one or more of the following letters:

- t** title is to be printed
- l** lines are to bracket the title
- d** a date and time is to be printed
- v** version number of program is to be printed
- f** file name being analyzed is to be printed

*title* string, title for header.

SOURCE gaussmt.src

h

## hess

**PURPOSE** Computes the Hessenberg form of a square matrix.

**FORMAT** {  $h, z$  } = **hess**( $x$ );

**INPUT**  $x$   $K \times K$  matrix.

**OUTPUT**  $h$   $K \times K$  matrix, Hessenberg form.  
 $z$   $K \times K$  matrix, transformation matrix.

**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  $x$  into  $h$  and vice versa. Thus:

$$h = z' x z$$

and since  $z$  is orthogonal,

$$x = z h z'$$

$x$  is reduced to upper Hessenberg form using orthogonal similarity transformations. This preserves the Frobenious norm of the matrix and the condition numbers of the eigenvalues.

**hess** uses the ORTRAN and ORTHES functions from EISPACK.



```
EXAMPLE  let x[3,3] = 1 2 3
              4 5 6
              7 8 9;

          { h,z } = hess(x);
```

```
          1.00000000  -3.59700730  -0.24806947
h =  -8.06225775  14.04615385   2.83076923
          0.00000000   0.83076923  -0.04615385

          1.00000000   0.00000000   0.00000000
z =  0.00000000  -0.49613894  -0.86824314
          0.00000000  -0.86824314   0.49613894
```

h

SEE ALSO **schur**

hessMT

PURPOSE	Computes numerical Hessian.	
INCLUDE	optim.sdf	
FORMAT	<i>h</i> = <b>hessMT</b> (& <i>fct</i> , <i>par1</i> , <i>data1</i> );	
INPUT	& <i>fct</i>	scalar, pointer to procedure returning either N×1 vector or 1×1 scalar.
	<i>par1</i>	an instance of structure of type <b>PV</b> containing parameter vector at which Hessian is to be evaluated.
	<i>data1</i>	structure of type <b>DS</b> containing any data needed by <i>fct</i> .
OUTPUT	<i>h</i>	K×K matrix, Hessian.

## hessMTg

---

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 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    &*gfct*      scalar, pointer to procedure computing either 1×K gradient or N×K Jacobian.

*par1* an instance of structure of type **PV** containing parameter vector at which Hessian is to be evaluated.

*data1* structure of type **DS** containing any data needed by *gfct*.

OUTPUT *h*  $K \times K$  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 = hessMTg(&gfct,p1,d0);

```

SOURCE `hessmt.src`

h

hessMTgw

PURPOSE Computes numerical Hessian using gradient procedure with weights.

## hessMTgw

---

INCLUDE    `optim.sdf`

FORMAT    `h = hessMTgw(&gfct,par1,data1,wgts);`

INPUT    **&gfct**      scalar, pointer to procedure computing either N×K Jacobian.  
         *par1*        an instance of structure of type **PV** containing parameter vector at  
                        which Hessian is to be evaluated.  
         *data1*        structure of type **DS** containing any data needed by *gfct*.  
         *wgts*         N×1 vector.

OUTPUT    *h*            K×K 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);
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.

INCLUDE optim.sdf

FORMAT  $h = \text{hessMTm}(\&fct, par1, data1, mask);$

INPUT	$\&fct$ $par1$  $data1$ $mask$	scalar, pointer to procedure returning either $N \times 1$ vector or scalar. an instance of structure of type <b>PV</b> containing parameter vector at which Hessian is to be evaluated.  structure of type <b>DS</b> containing any data needed by $fct$ . $K \times K$ matrix, elements in $h$ corresponding to elements of mask set to zero are not computed, otherwise are computed.
-------	--	--

h

OUTPUT  $h$   $K \times K$  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 p1;
p1 = pvCreate;
p1 = pvPack(p1, 0.1 | 0.2, "P");
struct DS d0;
d0 = dsCreate;
d0.dataMatrix = seqa(1, 1, 15);

mask = { 1 1,
```

## hessMTmw

---

```
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  $h = \mathbf{hessMTmw}(\&fct,par1,data1,mask,wgts);$

INPUT	$\&fct$	scalar, pointer to procedure returning $N \times 1$ vector.
	$par1$	an instance of structure of type <b>PV</b> containing parameter vector at which Hessian is to be evaluated.
	$data1$	structure of type <b>DS</b> containing any data needed by $fct$ .
	$mask$	$K \times K$ matrix, elements in $h$ corresponding to elements of mask set to zero are not computed, otherwise are computed.
	$wgts$	$N \times 1$ vector, weights.

OUTPUT  $h$   $K \times K$  matrix, Hessian.

REMARKS *fct* must evaluate to an  $N \times 1$  vector conformable to the weight vector. *par1* must be created using the **pvpPack** 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 = hessMTw(&fct,p1,d0,mask,wgt);

```

h

SOURCE `hessmt.src`

hessMTw

PURPOSE Computes numerical Hessian with weights.

INCLUDE `optim.sdf`

## hessp

---

FORMAT    *h* = **hessMTw**(&*fct*,*par1*,*data1*,*wgts*);

INPUT    **&*fct***       scalar, pointer to procedure returning N×1 vector.  
         *par1*       an instance of structure of type **PV** containing parameter vector at  
                       which Hessian is to be evaluated.  
         *data1*       structure of type **DS** containing any data needed by *fct*.  
         *wgts*       N×1 vector, weights.

OUTPUT    *h*           K×K matrix, Hessian.

REMARKS   *fct* must evaluate to an N×1 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;

    p = pvUnpack(p0,"P");
    y = p[1] * exp( -p[2] * d0.dataMatrix );
    retp(y);
endp;

h = hessMTw(&fct,p1,d0,wgt);
```

SOURCE    `hessmt.src`



**PURPOSE** Computes the matrix of second partial derivatives (Hessian matrix) of a function defined as a procedure.

**FORMAT**  $h = \text{hessp}(\&f, x0);$

**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  $x0$ ; 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);
    retp( exp(x'b) );
endp;
```

$b0 = \{ 3, 2, 1 \};$

h

## hist

---

```
h = hessp(&g,b0);
```

The resulting matrix of second partial derivatives of **g(b)** evaluated at **b=b0** is:

```
22027.12898372  44054.87238165  66083.36762901
44054.87238165  88111.11102645  132168.66742899
66083.36762901  132168.66742899  198256.04087836
```

SOURCE `hessp.src`

SEE ALSO **gradp**

## hist

**PURPOSE** Computes and graphs a frequency histogram for a vector. The actual frequencies are plotted for each category.

**LIBRARY** `pgraph`

**FORMAT** `{ b,m,freq } = hist(x,v);`

**INPUT** *x*            M×1 vector of data.  
*v*                N×1 vector, the breakpoints to be used to compute the frequencies  
                 - or -  
                 scalar, the number of categories.

**OUTPUT** *b*            P×1 vector, the breakpoints used for each category.  
*m*                P×1 vector, the midpoints of each category.  
*freq*            P×1 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:

$$\begin{aligned} freq[1] &= & x &\leq b[1] \\ freq[2] &= b[1] & < x &\leq b[2] \\ freq[3] &= b[2] & < x &\leq b[3] \\ &\vdots \\ &\vdots \\ freq[P] &= b[P-1] & < x &\leq b[P] \end{aligned}$$

h

EXAMPLE    `library pgraph;`  
              `x = rndn(5000,1);`  
              `{ b,m,f } = hist(x,20);`

SOURCE    `phist.src`

SEE ALSO    **histp, histf, bar**

histf

PURPOSE    Graphs a histogram given a vector of frequency counts.

LIBRARY    `pgraph`

FORMAT    **histf(f,c);**

INPUT       $f$              $N \times 1$  vector, frequencies to be graphed.

## histp

---

	<i>c</i>	N×1 vector, numeric labels for categories. If this is a scalar 0, a sequence from 1 to <b>rows(<i>f</i>)</b> will be created.
REMARKS		The axes are not automatically labeled. Use <b>xlabel</b> for the category axis and <b>ylabel</b> for the frequency axis.
SOURCE		phist.src
SEE ALSO		<b>hist</b> , <b>bar</b> , <b>xlabel</b> , <b>ylabel</b>

## histp

PURPOSE		Computes and graphs a percent frequency histogram of a vector. The percentages in each category are plotted.
LIBRARY		pgraph
FORMAT		{ <i>b,m,freq</i> } = <b>histp</b> ( <i>x,v</i> );
INPUT	<i>x</i>	M×1 vector of data.
	<i>v</i>	N×1 vector, the breakpoints to be used to compute the frequencies - or - scalar, the number of categories.
OUTPUT	<i>b</i>	P×1 vector, the breakpoints used for each category.
	<i>m</i>	P×1 vector, the midpoints of each category.
	<i>freq</i>	P×1 vector of computed frequency counts. This is the vector of counts, not percentages.
REMARKS		If a vector of breakpoints is specified, a final breakpoint equal to the maximum value of <i>x</i> 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:

$$\begin{aligned}
 freq[1] &= & x &\leq b[1] \\
 freq[2] &= b[1] & < x &\leq b[2] \\
 freq[3] &= b[2] & < x &\leq b[3] \\
 &\vdots \\
 &\vdots \\
 freq[P] &= b[P-1] & < x &\leq b[P]
 \end{aligned}$$

h

SOURCE `phist.src`

SEE ALSO **hist**, **histf**, **bar**

hsec

PURPOSE Returns the number of hundredths of a second since midnight.

FORMAT  $y = \mathbf{hsec};$

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 = rndu(1000,1000);`  
`ts = hsec;`

## if

---

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

In this example, **hsec** is used to time a 1000×1000 multiplication in **GAUSS**. A 1000×1000 matrix, **x**, is created, and the current time, in hundredths of a second since midnight, is stored in the variable **ts**. Then the multiplication is carried out. Finally, **ts** is subtracted from **hsec** to give the time difference which is assigned to **et**.

SEE ALSO **date, time, timestr, ethsec, etstr**

## if

**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**

i

## imag

PURPOSE    Returns the imaginary part of  $x$ .

FORMAT     $zi = \text{imag}(x);$

INPUT       $x$              $N \times K$  matrix or  $N$ -dimensional array.

OUTPUT     $zi$              $N \times K$  matrix or  $N$ -dimensional array, the imaginary part of  $x$ .

REMARKS    If  $x$  is real,  $zi$  will be an  $N \times K$  matrix or  $N$ -dimensional array of zeros.

## #include

---

EXAMPLE     $x = \begin{Bmatrix} 4i & 9 & 3, \\ 2 & 5-6i & 7i \end{Bmatrix};$   
              $y = \text{imag}(x);$

$$y = \begin{bmatrix} 4.0000000 & 0.0000000 & 0.0000000 \\ 0.0000000 & -6.0000000 & 7.0000000 \end{bmatrix}$$

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;` No additional search will be made if the file is not found.

`#include myprog.prc;` 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.

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**

i

## 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>	N×1 character vector which contains the elements to be found in vector <i>where</i> .
<i>where</i>	M×1 character vector to be searched for matches to the elements of <i>what</i> .

**OUTPUT**

<i>z</i>	N×1 vector of integers containing the indices of the corresponding element of <i>what</i> in <i>where</i> .
----------	---

## indexcat

---

**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 what = AGE PAY SEX;`  
              `let where = AGE SEX JOB date PAY;`  
              `z = indcv(what,where);`

```
          AGE
what =    PAY
          SEX
```

```
          AGE
          SEX
where =   JOB
          date
          PAY
```

```
          1
z =       5
          2
```

**SEE ALSO**    **indnv, indsav**

## indexcat

**PURPOSE**    Returns the indices of the elements of a vector which fall into a specified

category

FORMAT `y = indexcat(x,v);`

INPUT `x`        `N`×1 vector.  
      `v`        scalar or 2×1 vector.  
              If scalar, the function returns the indices of all elements of `x` equal to `v`.  
              If 2×1, then the function returns the indices of all elements of `x` that fall into the range:  
                       $v[1] < x \leq v[2]$   
              If `v` is scalar, it can contain a single missing to specify the missing value as the category.

OUTPUT `y`        `L`×1 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;`  
              `y = indexcat(x,v);`

              1.0  
              4.0  
              3.3  
              4.2  
`x =`        6.0  
              5.7  
              8.1  
              5.5

              4  
`v =`        6



## indices

---

$y = \begin{matrix} 4 \\ 5 \\ 6 \\ 8 \end{matrix}$

### 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** *dataset* string, the name of the data set.  
*vars* N×1 vector, a character vector of names or a numeric vector of column indices.  
If scalar 0, all variables in the data set will be selected.

**OUTPUT** *name* N×1 character vector, the names associated with *vars*.  
*indx* N×1 numeric vector, the column indices associated with *vars*.

**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.<br>This can be either the name of the variable, or the column index of the variable.<br>If null or 0, the last variable in the data set will be used. |
| <i>var2</i>    | N×1 vector, a character vector of names or a numeric vector of column indices.<br>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> . |
| <i>indx1</i> | scalar, the column index of <i>var1</i> .   |
| <i>name2</i> | N×1 character vector, the names associated with <i>var2</i> .                             |
| <i>indx2</i> | N×1 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:

## indicesf

---

- 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    *fp*            scalar, file handle of an open data set.  
          *namein*    N×1 string array, names of selected columns in the data set. If set to a null string, columns are selected using *indxin*  
          *indxin*    N×1 vector, indices of selected columns in the data set. If set to 0, columns are selected using *namein*.

OUTPUT    *name*           N×1 string array, the names of the selected columns.  
          *indx*           N×1 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    **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>	N×1 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>	N×1 vector, indices of selected columns in the data set. If set to 0, columns are selected using <i>namein</i> .

**OUTPUT**

<i>name</i>	N×1 string array, the names of the selected columns.
<i>indx</i>	N×1 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.

**FORMAT**  $z = \text{indnv}(\text{what}, \text{where});$

**INPUT** *what*  $N \times 1$  numeric vector which contains the values to be found in vector *where*.

*where*  $M \times 1$  numeric vector to be searched for matches to the values in *what*.

**OUTPUT**  $z$   $N \times 1$  vector of integers, the indices of the corresponding elements of *what* in *where*.

**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**

```
let what = 8 7 3;
let where = 2 7 8 4 3;
z = indnv(what,where);
```

```
      8
what = 7
      3
```



```

      2
      7
where = 8
      4
      3

      3
z =    2
      5
```

SEE ALSO    **indcv**

indsav

i

- 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

*what*        N×1 string array which contains the values to be found in vector *where*.

*where*        M×1 string array to be searched for the corresponding elements of *what*.
- OUTPUT

*indx*        N×1 vector of indices, the values of *what* in *where*.
- 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.

## intgrat2

**PURPOSE** Integrates the following double integral, using user-defined functions  $f$ ,  $g_1$  and  $g_2$  and scalars  $a$  and  $b$ :

$$\int_a^b \int_{g_2(x)}^{g_1(x)} f(x, y) dy dx$$

**FORMAT**  $y = \text{intgrat2}(\&f, xl, gl);$

**INPUT**  $\&f$  scalar, pointer to the procedure containing the function to be integrated.

$xl$   $2 \times 1$  or  $2 \times N$  matrix, the limits of  $x$ . These must be scalar limits.

$gl$   $2 \times 1$  or  $2 \times N$  matrix of function pointers, the limits of  $y$ .  
For  $xl$  and  $gl$ , 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.

$\_intrec$  scalar. This variable is used to keep track of the level of recursion of **intgrat2** and may start out with a different value if your program terminated inside of the integration function on a previous run. Always set  $\_intrec$  explicitly to 0 before any call to **intgrat2**.

**OUTPUT**  $y$   $N \times 1$  vector of the estimated integral(s) of  $f(x, y)$ , evaluated between the limits given by  $xl$  and  $gl$ .

**REMARKS** The user-defined functions specified by  $f$  and  $gl$  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 f(x,y);
    retp(cos(x) + 1).*(sin(y) + 1));
endp;

proc g1(x);
    retp(sqrt(1-x^2));
endp;

proc g2(x);
    retp(0);
endp;

x1 = 1|-1;
g0 = &g1|&g2;
_intord = 40;
_intrec = 0;
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, \_intrec

SEE ALSO intgrat3, intquad1, intquad2, intquad3, intsimp

## intgrat3

**PURPOSE** Integrates the following triple integral, using user-defined functions and scalars for bounds:

$$\int_a^b \int_{g_2(x)}^{g_1(x)} \int_{h_2(x,y)}^{h_1(x,y)} f(x,y,z) dz dy dx$$

**FORMAT** `y = intgrat3(&f,xl,gl,hl);`

**INPUT**

- &f** scalar, pointer to the procedure containing the function to be integrated. *f* is a function of (*x,y,z*).
- xl** 2×1 or 2×N matrix, the limits of *x*. These must be scalar limits.
- gl** 2×1 or 2×N matrix of function pointers. These procedures are functions of *x*.
- hl** 2×1 or 2×N matrix of function pointers. These procedures are functions of *x* and *y*.

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**

- \_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.
- \_intrec** scalar. This variable is used to keep track of the level of recursion of **intgrat3** and may start out with a different value if your program terminated inside of the integration function on a previous run. Always set **\_intrec** explicitly to 0 before any call to **intgrat3**.

**OUTPUT** `y` N×1 vector of the estimated integral(s) of *f(x,y,z)* evaluated between the limits given by *xl*, *gl* and *hl*.

REMARKS    User-defined functions  $f$ , and those used in  $gl$  and  $hl$  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;`

`proc h2(x,y);`  
                  `retp(-h1(x,y));`  
              `endp;`

`x1 = 5|-5;`  
              `g0 = &g1|&g2;`  
              `h0 = &h1|&h2;`  
              `_intrec = 0;`  
              `_intord = 40;`  
              `y = intgrat3(&f,x1,g0,h0);`

This will integrate the function  $f(x,y,z) = 2$  over the sphere of radius 5. The

## inthp1

---

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, _intrec`

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    *&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:

*pds->dataMatrix*     $N \times K$  matrix.

*pds->dataArray*     $N \times K \times L \dots$  array.

*pds->vnames*        string array.

*pds->dsname*        string.

*pds->type*          scalar.

The contents, if any, are set by the user and are passed by **inthp1** to the user-provided function without modification.

*ctl*       instance of an **inthpControl** structure with members

*ctl.maxEvaluations*    scalar, maximum number of function evaluations, default = 1e5;

A default *ctl* can be generated by calling **inthpControlCreate**.

REMARKS The user-provided function must have the following format

where

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.

```
EXAMPLE  #include inthp.sdf

proc fct(struct DS *pds, x);
```

## inthp2

---

```
        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);
```

```
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  $y = \text{inthp2}(\&f, pds, ctl, a);$

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:

**pds->dataMatrix** N×K matrix.

**pds->dataArray** N×K×L... array.

**pds->vnames** string array.

**pds->dsname** string.

**pds->type** scalar.

The contents, if any, are set by the user and are passed by **inthp1** to the user-provided function without modification.

**ctl** instance if an **inthpControl** structure with members

**ctl.maxEvaluations** scalar, maximum number of function evaluations, default = 1e5;

**ctl.p** scalar, termination parameter

**0** heuristic termination, default.

**1** deterministic termination with infinity norm.

**2,...** deterministic termination with p-th norm.

**ctl.d** scalar termination parameter

**1** if heuristic termination

**0 < ctl.d ≤ π/2** if deterministic termination

**ctl.eps** scalar, relative error bound. Default = 1e-6.

A default **ctl** can be generated by calling **inthpControlCreate**.

**a** 1×N vector, lower limits of integration

OUTPUT      $y$               $N \times 1$  vector, the estimated integrals of  $f(x)$  evaluated over the interval  $[a, +\infty)$ .

REMARKS     The user-provided function must have the following format

$f(\text{struct DS } *pds, x)$

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 normal(struct DS *pd0, x);
    local var;
    var = pd0->dataMatrix;
    retp( (1/sqrt(2*pi*var))*exp( -(x*x) / (2*var) ) );
endp;

struct DS d0;
struct DS *pd0;

pd0 = &d0;
d0.dataMatrix = var;

struct inthpControl c0;
c0 = inthpControlCreate;
```

```

lim = 2;

c0.d = pi/4;
c0.p = 2;

var = 1;

r = inthp2(&normal,pd0,c0,lim);

format /ld 16,10;
print r;
print cdfnc(2);

0.0227501281
0.0227501319

```

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  $y = \text{inthp3}(\&f, pds, ctl, a);$

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:

*pds->dataMatrix* N×K matrix.  
*pds->dataArray* N×K×L... array.  
*pds->vnames* string array.  
*pds->dsname* string.  
*pds->type* scalar.

The contents, if any, are set by the user and are passed by **inthp1** to the user-provided function without modification.

*ctl* instance if an **inthpControl** structure with members

*ctl.maxEvaluations* scalar, maximum number of function evaluations, default = 1e5;  
*ctl.p* scalar, termination parameter  
**0** heuristic termination, default.  
**1** deterministic termination with infinity norm.  
**2,...** deterministic termination with p-th norm.  
*ctl.d* scalar termination parameter  
**1** if heuristic termination  
 $0 < \text{ctl.d} \leq \pi/2$  if deterministic termination  
*ctl.eps* scalar, relative error bound. Default = 1e-6.

A default *ctl* can be generated by calling **inthpControlCreate**.

*a* 1×N vector, lower limits of integration

OUTPUT *y* N×1 vector, the estimated integrals of  $f(x)$  evaluated over the interval  $[a, +\infty)$ .

REMARKS This procedure is designed especially for oscillatory functions.

The user-provided function must have the following format

$f(\text{struct DS } *pds, x)$

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);
    
```

## inthp4

---

```
format /ld 16,10;  
print r;  
print a/(a*a + m*m);
```

```
0.20000000000  
0.20000000000
```

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:

$pds \rightarrow dataMatrix$   $N \times K$  matrix.

$pds \rightarrow dataArray$   $N \times K \times L \dots$  array.

$pds \rightarrow vnames$  string array.

$pds \rightarrow dsname$  string.

$pds \rightarrow type$  scalar.

The contents, if any, are set by the user and are passed by **inthp1** to the user-provided function without modification.

*ctl* instance if an **inthpControl** structure with members

*ctl.maxEvaluations* scalar, maximum number of function evaluations, default = 1e5;

*ctl.p* scalar, termination parameter

**0** heuristic termination, default.

**1** deterministic termination with infinity norm.

**2,...** deterministic termination with p-th norm.

*ctl.d* scalar termination parameter

**1** if heuristic termination

**0 < ctl.d ≤ π/2** if deterministic termination

*ctl.eps* scalar, relative error bound. Default = 1e-6.

A default *ctl* can be generated by calling **inthpControlCreate**.

*c* 2×N vector, upper and lower limits of integration, the first row contains upper limits and the second row the lower.

OUTPUT    *y*    N×1 vector, the estimated integrals of  $f(x)$  evaluated over the interval  $[a,b]$ .

REMARKS    The user-provided function must have the following format

$f(\text{struct DS } *pds, x)$

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 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 inthpControl c0;
c0 = inthpControlCreate;

c0.p = 2;
c0.d = pi/2;

a = -1;
b = -2;
c = 3;
pd0 = &d0;
d0.dataMatrix = a|b|c;

lim = 1 | -1;

r = inthp4(&fct,pd0,c0,lim);

format /ld 16,10;
print r;
print pi/2;
```



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 inthp.src

SEE ALSO inthpControlCreate, inthp1, inthp2, inthp3

## inthpControlCreate

PURPOSE Creates default **inthpControl** structure.

INCLUDE inthp.sdf

FORMAT  $c = \text{inthpControlCreate};$

OUTPUT  $c$  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.

## intquad2

---

FORMAT	$y = \text{intquad1}(\&f, xl);$	
INPUT	$\&f$	scalar, pointer to the procedure containing the function to be integrated. This must be a function of $x$ .
	$xl$	2×N matrix, the limits of $x$ . 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	$y$	N×1 vector of the estimated integral(s) of $f(x)$ evaluated between the limits given by $xl$ .
REMARKS	The user-defined function $f$ must return a vector of function values. <b>intquad1</b> 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	<pre>proc f(x);     retp(x.*sin(x)); endp;  xl = 1 0; y = intquad1(&amp;f,xl);</pre> <p>This will integrate the function <math>f(x) = x\sin(x)</math> between 0 and 1. Note the use of the <math>\_*</math> instead of <math>*</math>.</p>	
SOURCE	integral.src	
GLOBALS	$\_intord, \_intq12, \_intq16, \_intq2, \_intq20, \_intq24, \_intq3, \_intq32, \_intq4, \_intq40, \_intq6, \_intq8$	
SEE ALSO	<b>intsimp, intquad2, intquad3, intgrat2, intgrat3</b>	

## 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**  $y = \text{intquad2}(\&f, xl, yl);$

**INPUT**  $\&f$  scalar, pointer to the procedure containing the function to be integrated.

$xl$   $2 \times 1$  or  $2 \times N$  matrix, the limits of  $x$ .

$yl$   $2 \times 1$  or  $2 \times N$  matrix, the limits of  $y$ .

For  $xl$  and  $yl$ , 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.

$\_intrec$  scalar. This variable is used to keep track of the level of recursion of **intquad2** and may start out with a different value if your program terminated inside of the integration function on a previous run. Always set  $\_intrec$  explicitly to 0 before any calls to **intquad2**.

**OUTPUT**  $y$   $N \times 1$  vector of the estimated integral(s) of  $f(x, y)$  evaluated between the limits given by  $xl$  and  $yl$ .

**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

## intquad3

---

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**.

```
EXAMPLE  proc f(x,y);
          retp(x.*sin(x+y));
        endp;

        x1 = 1|0;
        y1 = 1|0;

        _intrec = 0;
        y = intquad2(&f,x1,y1);
```

This will integrate the function **`x.*sin(x+y)`** between **`x = 0`** and **`1`**, and between **`y = 0`** and **`1`**.

SOURCE `integral.src`

GLOBALS `_intord, _intq12, _intq16, _intq2, _intq20, _intq24, _intq3, _intq32, _intq4, _intq40, _intq6, _intq8, _intrec`

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);`

INPUT `&f` scalar, pointer to the procedure containing the function to be

For  $xl$ ,  $yl$ , and  $zl$ , the first row is the upper limit and the second row is the lower limit.  $N$  integrations are computed.

OUTPUT	$y$	N×1 vector of the estimated integral(s) of $f(x, y, z)$ evaluated between the limits given by $xl$ , $yl$ , and $zl$ .
--------	-----	--

**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.

```
EXAMPLE  proc f(x,y,z);
           ret(x.*y.*z);
        endp;
```

## intrleav

---

```
x1 = 1|0;  
y1 = 1|0;  
z1 = { 1 2 3, 0 0 0 };  
  
_intrec = 0;  
y = intquad3(&f,x1,y1,z1);
```

This will integrate the function  $f(x) = x * y * z$  over 3 sets of limits, since  $z1$  is defined to be a 2×3 matrix.

SOURCE    `integral.src`

GLOBALS    `_intord, _intq12, _intq16, _intq2, _intq20, _intq24, _intq3, _intq32,`  
             `_intq4, _intq40, _intq6, _intq8, _intrec`

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    *infile1*    string, name of input file 1.  
          *infile2*    string, name of input file 2.  
          *outfile*    string, name of output file.  
          *keyvar*    string, name of key variable; this is the column the files are sorted on.  
          *keytyp*    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

i

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

FORMAT     `y = intrleavsa(sa1,sa2,ike y);`

INPUT       *sa1*        N×K string array 1.  
              *sa2*        M×K string array 2.  
              *ike y*       scalar integer, index of the key column the string arrays are sorted on.

OUTPUT     *y*         L×K 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.

## intrsect

---

SOURCE    `sortd.src`

SEE ALSO    **intrleav**

## intrsect

PURPOSE    Returns the intersection of two vectors, with duplicates removed.

FORMAT    `y = intrsect(v1,v2,flag);`

INPUT	<i>v1</i>	N×1 vector.
	<i>v2</i>	M×1 vector.
	<i>flag</i>	scalar, if 1, <i>v1</i> and <i>v2</i> are numeric; if 0, character.

OUTPUT    *y*        L×1 vector containing all unique values that are in both *v1* and *v2*, sorted in ascending order.

REMARKS    Place smaller vector first for fastest operation.

If there are a lot of duplicates within a vector, it is faster to remove them with **unique** before calling **intrsect**.

SOURCE    `intrsect.src`

EXAMPLE    `v1 = { 3, 9, 5, 2, 10, 15 };`  
              `v2 = { 4, 9, 8, 5, 12, 3, 1 };`  
              `y = intrsect(v1,v2,1);`

$$y = \begin{matrix} 3 \\ 5 \\ 9 \end{matrix}$$



---

SEE ALSO **intrsectsa**

## intrsectsa

**PURPOSE** Returns the intersection of two string vectors, with duplicates removed.

**FORMAT**  $y = \text{intrsectsa}(sv1, sv2);$

**INPUT**  $sv1$   $N \times 1$  or  $1 \times N$  string vector.  
 $sv2$   $M \times 1$  or  $1 \times M$  string vector.

**OUTPUT**  $sy$   $L \times 1$  vector containing all unique strings that are in both  $sv1$  and  $sv2$ , sorted in ascending order.

**REMARKS** Place smaller vector first for fastest operation.

If there are a lot of duplicates it is faster to remove them with `unique` before calling `intrsectsa`.

**EXAMPLE**

```
string sv1 = { "mary", "jane", "linda", "dawn" };  
string sv2 = { "mary", "sally", "jane", "lisa", "ruth" };  
sy = intrsectsa(sv1,sv2);
```

**SOURCE** `intrsect.src`

SEE ALSO **intrsect**

i

## intsimp

**PURPOSE** Integrates a specified function using Simpson's method with end correction. A single integral is computed in one function call.

## inv, invpd

---

FORMAT     $y = \text{intsimp}(\&f, xl, tol);$

INPUT     $\&f$             pointer to the procedure containing the function to be integrated.  
           $xl$              $2 \times 1$  vector, the limits of  $x$ .  
                         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  $xl[1]$  and  $xl[2]$ .

EXAMPLE   `proc f(x);`  
            `retp(sin(x));`  
            `endp;`  
  
            `let xl = { 1,`  
                    `0 };`  
  
            `y = intsimp(&f,xl,1E-8);`

$y = 0.45969769$

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     $y = \text{inv}(x);$   
             $y = \text{invpd}(x);$

INPUT      $x$              $N \times N$  matrix or  $K$ -dimensional array where the last two dimensions are  $N \times N$ .

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  $4 \times 4$  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:

**trap 1**, return error code

<b>inv</b>	<b>invpd</b>
50	20

**trap 0**, terminate with error message

<b>inv</b>	<b>invpd</b>
Matrix singular	Matrix not positive definite



## invswp

---

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;  
  
          bols = 1.017201  0.484244
```

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.

## invswp

**PURPOSE**    Computes a generalized sweep inverse.

**FORMAT**     $y = \text{invswp}(x);$

**INPUT**       $x$              $N \times N$  matrix.

**OUTPUT**     $y$              $N \times N$  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, Appendix C.

**EXAMPLE** `let x[3,3] = 1 2 3 4 5 6 7 8 9;`  
`y = invswp(x);`

```
      -1.6666667    0.66666667  0.0000000
y =   1.3333333   -0.33333333  0.0000000
      0.0000000    0.0000000  0.0000000
```

i

**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.

## iscplx

---

OUTPUT    *y*            scalar, 1 if *x* is complex, 0 if it is real.

EXAMPLE    *x* = { 1, 2i, 3 };  
              *y* = iscplx(*x*);

*y* = 1.0000000

SEE ALSO    **hasimag, iscplx**

## iscplx

PURPOSE    Returns whether a data set is complex or real.

FORMAT    *y* = **iscplx**(*fh*);

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* = **isden**(*x*);

INPUT      *x*            N×K matrix or N-dimensional array.

OUTPUT     $y$             scalar, 1 if  $x$  contains a denormal, 0 if it does not.

EXAMPLE     $x = \{ 1, \exp(-724.5), 3 \};$   
                $y = \text{isden}(x);$

$y = 1.0000000$

SEE ALSO    **denToZero**

## isinfnanmiss

PURPOSE    Returns true if the argument contains an infinity, NaN, or missing value.

FORMAT     $y = \text{isinfnanmiss}(x);$

INPUT     $x$              $N \times K$  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     $x$              $N \times K$  matrix.

## keep (dataloop)

---

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 6 3 4 };  
              *y* = ismiss(*x*);

*y* = 0.0000000

SEE ALSO    **scalmiss**, **miss**, **missrv**

## 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)**



**PURPOSE** Returns the ASCII value of the next key available in the keyboard buffer.

**FORMAT**  $y = \mathbf{key};$

**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.

Here are the values returned if the key pressed is not a standard ASCII character in the range of 1-255:

1015	SHIFT+TAB
1016-1025	ALT+Q, W, E, R, T, Y, U, I, O, P
1030-1038	ALT+A, S, D, F, G, H, J, K, L
1044-1050	ALT+Z, X, C, V, B, N, M
1059-1068	F1-F10
1071	HOME
1072	CURSOR UP
1073	PAGE UP
1075	CURSOR LEFT
1077	CURSOR RIGHT
1079	END
1080	CURSOR DOWN
1081	PAGE DOWN
1082	INSERT

1083	DELETE
1084-1093	SHIFT+F1-F10
1094-1103	CTRL+F1-F10
1104-1113	ALT+F1-F10
1114	CTRL+PRINT SCREEN
1115	CTRL+CURSOR LEFT
1116	CTRL+CURSOR RIGHT
1117	CTRL+END
1118	CTRL+PAGE DOWN
1119	CTRL+HOME
1120-1131	ALT+1,2,3,4,5,6,7,8,9,0,-,=
1132	CTRL+PAGE UP

```
EXAMPLE  format /rds 1,0;
          kk = 0;
          do until kk =\,= 27;
              kk = key;
              if kk =\,= 0;
                  continue;
              elseif kk =\,= vals(" ");
                  print "space \\" kk;
              elseif kk =\,= vals("\r");
                  print "carriage return \\" kk;
              elseif kk >= vals("0") and kk <= vals("9");
                  print "digit \\" kk chrs(kk);
              elseif vals(upper(chrs(kk))) >= vals("A") and
                  vals(upper(chrs(kk))) <= vals("Z");
                  print "alpha \\" kk chrs(kk);
              else;
                  print "\\\" kk;
              endif;
          endo;
```

This is an example of a loop that processes keyboard input. This loop will continue until the escape key (ASCII 27) is pressed.

---

SEE ALSO    **vals, chrs, upper, lower, con, cons**

**keyav**

PURPOSE    Check if keystroke is available.

FORMAT     $x = \mathbf{keyav};$

OUTPUT     $x$             scalar, value of key or 0 if no key is available.

SEE ALSO    **keyw, key**

**keyw****k**

PURPOSE    Waits for and gets a key.

FORMAT     $k = \mathbf{keyw};$

OUTPUT     $k$             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 **key** for a table of return values for extended and function keys.

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*);

**INPUT** *name* literal, name of the keyword. This name will be a global symbol.

*str* 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 12.

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)

PURPOSE Lags variables a specified number of periods.

FORMAT **lag** *nv1* = *var1:p1* [[*nv2* = *var2:p2...*]];

INPUT *var* name of the variable to lag.  
*p* scalar constant, number of periods to lag.

OUTPUT *nv* 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 dataloop, and is then automatically deleted.

## lag1

---

### lag1

PURPOSE	Lags a matrix by one time period for time series analysis.	
FORMAT	$y = \text{lag1}(x);$	
INPUT	$x$	N×K matrix.
OUTPUT	$y$	N×K matrix, $x$ lagged 1 period.
REMARKS	<b>lag1</b> lags $x$ by one time period, so the first observations of $y$ are missing.	
SOURCE	lag.src	
SEE ALSO	<b>lag1</b>	

### lag1

PURPOSE	Lags a matrix a specified number of time periods for time series analysis.	
FORMAT	$y = \text{lag1}(x, t);$	
INPUT	$x$	N×K matrix.
	$t$	scalar, number of time periods.
OUTPUT	$y$	N×K matrix, $x$ lagged $t$ periods.
REMARKS	If $t$ is positive, <b>lag1</b> lags $x$ back $t$ time periods, so the first $t$ observations of $y$ are missing. If $t$ is negative, <b>lag1</b> lags $x$ forward $t$ time periods, so the last $t$ observations of $y$ are missing.	

SOURCE `lag.src`

SEE ALSO **lag1**

## lapeighb

**PURPOSE** Computes eigenvalues only of a real symmetric or complex Hermitian matrix selected by bounds.

**FORMAT** `ve = lapeighb(x,vl,vu);`

**INPUT**

<i>x</i>	N×N matrix, real symmetric or complex Hermitian.
<i>vl</i>	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 $abstol + EPS \cdot \max( a ,  b )$ , where $EPS$ is machine precision. If <i>abstol</i> is less than or equal to zero, then $EPS \cdot \ T\ $ will be used in its place, where $T$ is the tridiagonal matrix obtained by reducing the input matrix to tridiagonal form.

**OUTPUT** *ve* M×1 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 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 DYESVX and ZHEEVX. Further documentation of these functions may be found in the LAPACK User's Guide.

## lapeighi

---

```
EXAMPLE  x = { 5   2   1,
               2   6   2,
               1   2   9 };

          vl = 5;
          vu = 10;
          ve = lapeighi(x,vl,vu,0);
          print ve;

          6.0000
```

SEE ALSO **lapeighb**, **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);`

<b>INPUT</b>	<i>x</i>	N×N 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 $[a,b]$ of width less than or equal to $abstol + EPS \cdot \max( a , b )$ , where $EPS$ is machine precision. If <i>abstol</i> is less than or equal to zero, then $EPS \cdot \ 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 *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 DYESVX and ZHEEVX. Further documentation of these functions may be found in the LAPACK User's Guide.

EXAMPLE  $x = \begin{Bmatrix} 5 & 2 & 1, \\ 2 & 6 & 2, \\ 1 & 2 & 9 \end{Bmatrix};$

```

il = 2;
iu = 3;
ve = lapeighi(x,il,iu,0);
print ve;

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  $\{ ve, va \} = \mathbf{lapeighvb}(x, vl, vu, abstol);$

INPUT *x*  $N \times N$  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* M×1 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.

*va* N×M matrix, eigenvectors.

REMARKS **lapeighvb** computes eigenvalues and eigenvectors which are found on the half open interval  $[vl, vu]$ . **lapeighvb** is based on the LAPACK drivers DYESVX and ZHEEVX. Further documentation of these functions may be found in the LAPACK User's Guide.

EXAMPLE  $x = \begin{Bmatrix} 5 & 2 & 1 \\ 2 & 6 & 2 \\ 1 & 2 & 9 \end{Bmatrix};$

```

vl = 5;
vu = 10;
{ ve, va } = lapeighvb(x, vl, vu, 0);
print ve;

6.000000000

print va;

-0.57735027
0.000000000
0.000000000

```

SEE ALSO **lapeighvb**

## 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>	N×N 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-il</i> +1)×1 vector, eigenvalues.
<i>va</i>	N×( <i>iu-il</i> +1) matrix, eigenvectors.

**REMARKS** **lapeighvi** computes *iu-il*+1 eigenvalues and eigenvectors given a range of indices, i.e., the *i*<sup>th</sup> to *j*<sup>th</sup> 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 DYESVX 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 };
```

## lapgeig

---

```
il = 2;
iu = 3;
{ ve,va } = lapeighvi(x,il,iu,0);
print ve;

        6.000000000
       10.60555128

print va;

      -0.57735027      -0.57735027
      -0.31970025      -0.49079864
       0.000000000       0.000000000
```

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** *A*        N×N matrix, real or complex general matrix.  
*B*        N×N matrix, real or complex general matrix.

**OUTPUT** *va1*      N×1 vector, numerator of eigenvalues.  
*va2*      N×1 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 DGEGV and ZGEGV.

SEE ALSO **lapgeig, lapgeigh**

## lapgeigh

**PURPOSE** Computes generalized eigenvalues for a pair of real symmetric or Hermitian matrices.

**FORMAT** `ve = lapgeigh(A,B);`

**INPUT** *A*         $N \times N$  matrix, real or complex symmetric or Hermitian matrix.  
*B*         $N \times N$  matrix, real or complex positive definite symmetric or Hermitian matrix.

**OUTPUT** *ve*         $N \times 1$  vector, eigenvalues.

**REMARKS** *ve* is the vector of eigenvalues of the solution of the generalized symmetric eigenproblem of the form  $Ax = \lambda Bx$ .

**EXAMPLE**  $A = \begin{Bmatrix} 3 & 4 & 5, \\ 2 & 5 & 2, \\ 3 & 2 & 4 \end{Bmatrix};$

$B = \begin{Bmatrix} 4 & 2 & 2, \\ 2 & 6 & 1, \\ 2 & 1 & 8 \end{Bmatrix};$

`ve = lapgeigh(A,B);`

`print ve;`

## lapgeighv

---

-0.70051730  
0.48661989  
1.27818313

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* } = **lapgeighv**(*A*, *B*);

**INPUT** *A* N×N matrix, real or complex symmetric or Hermitian matrix.  
*B* N×N matrix, real or complex positive definite symmetric or Hermitian matrix.

**OUTPUT** *ve* N×1 vector, eigenvalues.  
*va* N×N matrix, eigenvectors.

**REMARKS** *ve* and *va* are the eigenvalues and eigenvectors of the solution of the generalized symmetric eigenproblem of the form  $Ax = \lambda Bx$ . Equivalently, *va* diagonalizes  $U'^{-1}AU^{-1}$  in the following way

$$vaU'^{-1}AY^{-1}va' = ve$$

where  $B = U'U$ . This procedure calls the LAPACK routines DSYGV and ZHEGV.

**EXAMPLE**  $A = \begin{bmatrix} 3 & 4 & 5 \end{bmatrix}$ ,

```

      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** { *val*, *va2*, *lve*, *rve* } = **lapgeigv**(*A*, *B*);

**INPUT** *A*        N×N matrix, real or complex general matrix.  
*B*        N×N matrix, real or complex general matrix.

**OUTPUT** *val*        N×1 vector, numerator of eigenvalues.

## lapgsvdcst

---

*va2*      N×1 vector, denominator of eigenvalues.  
*lve*      N×N left eigenvectors.  
*rve*      N×N 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  $U'^{-1}AU^{-1}$  where  $B = U'U$ , that is,

$$lve \ U'^{-1}AU \ lve' = w$$

and

$$rve'U'^{-1}AU^{-1}rve = w$$

This procedure calls the LAPACK routines DGEGV and ZGEGV.

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* } = **lapgsvdcst**(*A,B*);

INPUT    *A*      M×N matrix.



	$B$	$P \times N$ 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.
	$U$	$M \times M$ 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 **lapgsvdcs** and **lapgsvdcs**). Letting  $K+L$  = 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,  $Z = [0R]$ , and if  $M-K-L \geq 0$

$$D_1 = \begin{array}{cc} & \begin{array}{cc} K & L \end{array} \\ \begin{array}{c} K \\ L \\ M-K-L \end{array} & \begin{bmatrix} I & 0 \\ 0 & C \\ 0 & 0 \end{bmatrix} \end{array}$$

$$D_2 = \begin{array}{cc} & \begin{array}{cc} K & L \end{array} \\ \begin{array}{c} P \\ P-L \end{array} & \begin{bmatrix} 0 & S \\ 0 & 0 \end{bmatrix} \end{array}$$

$$\begin{bmatrix} 0 & R \end{bmatrix} = \begin{array}{cc} & \begin{array}{cc} N-K-L & K & L \end{array} \\ \begin{array}{c} K \\ L \end{array} & \begin{bmatrix} 0 & R_{11} & R_{12} \\ 0 & 0 & R_{22} \end{bmatrix} \end{array}$$

or if  $M-K-L < 0$

$$D1 = \begin{array}{cc} & \begin{array}{ccc} K & M-K & K+L-M \end{array} \\ \begin{array}{c} K \\ M-K \end{array} & \begin{bmatrix} I & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix} \end{array}$$

$$D2 = \begin{array}{cc} & \begin{array}{ccc} K & M-K & K+L-M \end{array} \\ \begin{array}{c} M-K \\ K+L-M \\ P-L \end{array} & \begin{bmatrix} 0 & S & 0 \\ 0 & 0 & I \\ 0 & 0 & 0 \end{bmatrix} \end{array}$$

$$\begin{bmatrix} 0 & R \end{bmatrix} = \begin{array}{cc} & \begin{array}{cccc} N-K-L & K & M-K & K+L-M \end{array} \\ \begin{array}{c} K \\ M-K \\ K+L-M \end{array} & \begin{bmatrix} 0 & R11 & R12 & R13 \\ 0 & 0 & R22 & R23 \\ 0 & 0 & 0 & R33 \end{bmatrix} \end{array}$$

(2) Form the matrix

$$X = Q \begin{bmatrix} I & 0 \\ 0 & R^{-1} \end{bmatrix}$$

then

$$A = U'^{-1} E_1 X$$

$$B = V'^{-1} E_2 X^{-1}$$

where  $E1 = \begin{bmatrix} 0 & D1 \end{bmatrix}$  and  $E2 = \begin{bmatrix} 0 & D2 \end{bmatrix}$ .

(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 **lapgsvds**, **lapgsvdst**

## lapgsvds

**PURPOSE** Compute the generalized singular value decomposition of a pair of real or complex general matrices.

**FORMAT** { *C,S,R* } = **lapgsvds**(*A,B*);

**INPUT** *A*        *M*×*N* real or complex matrix.  
           *B*        *P*×*N* real or complex matrix.

**OUTPUT** *C*        *L*×1 vector, singular values for *A*.  
           *S*        *L*×1 vector, singular values for *B*.  
           *R*        (*K*+*L*)×(*K*+*L*) upper triangular matrix.

**REMARKS** (1) The generalized singular value decomposition of *A* and *B* is

$$U'AQ = D_1Z$$

$$V'BQ = D_2Z$$

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*)×(*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,  $Z = [0R]$ , and if  $M-K-L \geq 0$

$$D1 = \begin{array}{cc} & \begin{array}{cc} K & L \end{array} \\ \begin{array}{c} K \\ L \\ M-K-L \end{array} & \begin{bmatrix} I & 0 \\ 0 & C \\ 0 & 0 \end{bmatrix} \end{array}$$

$$D2 = \begin{array}{cc} & \begin{array}{cc} K & L \end{array} \\ \begin{array}{c} P \\ P-L \end{array} & \begin{bmatrix} 0 & S \\ 0 & 0 \end{bmatrix} \end{array}$$

$$[0R] = \begin{array}{cc} & \begin{array}{cc} N-K-L & \begin{array}{cc} K & L \end{array} \end{array} \\ \begin{array}{c} K \\ L \end{array} & \begin{bmatrix} 0 & R11 & R12 \\ 0 & 0 & R22 \end{bmatrix} \end{array}$$

or if  $M-K-L < 0$

$$D1 = \begin{array}{cc} & \begin{array}{ccc} K & M-K & K+L-M \end{array} \\ \begin{array}{c} K \\ M-K \end{array} & \begin{bmatrix} I & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix} \end{array}$$

$$D2 = \begin{array}{cc} & \begin{array}{ccc} K & M-K & K+L-M \end{array} \\ \begin{array}{c} M-K \\ K+L-M \\ P-L \end{array} & \begin{bmatrix} 0 & S & 0 \\ 0 & 0 & I \\ 0 & 0 & 0 \end{bmatrix} \end{array}$$

$$[0R] = \begin{array}{cc} & \begin{array}{ccc} N-K-L & \begin{array}{ccc} K & M-K & K+L-M \end{array} \end{array} \\ \begin{array}{c} K \\ M-K \\ K+L-M \end{array} & \begin{bmatrix} 0 & R11 & R12 & R13 \\ 0 & 0 & R22 & R23 \\ 0 & 0 & 0 & R33 \end{bmatrix} \end{array}$$

(2) Form the matrix

$$X = Q \begin{bmatrix} I & 0 \\ 0 & R^{-1} \end{bmatrix}$$

then

$$A = U'^{-1} E_1 X$$

$$B = V'^{-1} E_2 X^{-1}$$

where  $E_1 = \begin{bmatrix} 0 & D_1 \end{bmatrix}$  and  $E_2 = \begin{bmatrix} 0 & D_2 \end{bmatrix}$ .

(3) The generalized singular value decomposition of  $A$  and  $B$  implicitly produces the singular value decomposition of  $AB^{-1}$ :

$$AB^{-1} = U D_1 D_2^{-1} V'$$

This procedure calls the LAPACK routines DGGSD and ZGGSD.

SEE ALSO **lapgsvdcst**, **lapgsvdst**

**lapgsvdst**

**PURPOSE** Compute the generalized singular value decomposition of a pair of real or complex general matrices.

**FORMAT** {  $D_1, D_2, Z, U, V, Q$  } = **lapgsvdst**( $A, B$ );

## lapgsvdst

---

INPUT	$A$	$M \times N$ matrix.
	$B$	$P \times N$ 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	

$$U' A Q = D1 Z$$

$$V' B Q = D2 Z$$

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,  $D1$  and  $D2$  are  $M \times (K+L)$  and  $P \times (K+L)$  matrices with entries on the diagonal,  $Z = [0R]$ , and if  $M-K-L \geq 0$

$$D1 = \begin{array}{cc} & \begin{array}{cc} K & L \end{array} \\ \begin{array}{c} K \\ L \\ M-K-L \end{array} & \begin{bmatrix} I & 0 \\ 0 & C \\ 0 & 0 \end{bmatrix} \end{array}$$

$$D2 = \begin{array}{cc} & \begin{array}{cc} K & L \end{array} \\ \begin{array}{c} P \\ P-L \end{array} & \begin{bmatrix} 0 & S \\ 0 & 0 \end{bmatrix} \end{array}$$

$$\begin{bmatrix} 0 & R \end{bmatrix} = \begin{matrix} & N-K-L & K & L \\ K & \begin{bmatrix} 0 & R11 & R12 \end{bmatrix} \\ L & \begin{bmatrix} 0 & 0 & R22 \end{bmatrix} \end{matrix}$$

or if  $M-K-L < 0$

$$D1 = \begin{matrix} & K & M-K & K+L-M \\ K & \begin{bmatrix} I & 0 & 0 \end{bmatrix} \\ M-K & \begin{bmatrix} 0 & 0 & 0 \end{bmatrix} \end{matrix}$$

$$D2 = \begin{matrix} & K & M-K & K+L-M \\ M-K & \begin{bmatrix} 0 & S & 0 \end{bmatrix} \\ K+L-M & \begin{bmatrix} 0 & 0 & I \end{bmatrix} \\ P-L & \begin{bmatrix} 0 & 0 & 0 \end{bmatrix} \end{matrix}$$

$$\begin{bmatrix} 0 & R \end{bmatrix} = \begin{matrix} & N-K-L & K & M-K & K+L-M \\ K & \begin{bmatrix} 0 & R11 & R12 & R13 \end{bmatrix} \\ M-K & \begin{bmatrix} 0 & 0 & R22 & R23 \end{bmatrix} \\ K+L-M & \begin{bmatrix} 0 & 0 & 0 & R33 \end{bmatrix} \end{matrix}$$

(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  $E1 = \begin{bmatrix} 0 & D1 \end{bmatrix}$  and  $E2 = \begin{bmatrix} 0 & D2 \end{bmatrix}$ .

(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 **lapgsvds, lapgsvdcst**

## lapgschur

**PURPOSE** Compute the generalized Schur form of a pair of real or complex general matrices.

**FORMAT**  $\{ sa, sb, q, z \} = \text{lapgschur}(A, B);$

**INPUT**  $A$   $N \times N$  matrix, real or complex general matrix.  
 $B$   $N \times N$  matrix, real or complex general matrix.

**OUTPUT**  $sa$   $N \times N$  matrix, Schur form of  $A$ .  
 $sb$   $N \times N$  matrix, Schur form of  $B$ .  
 $q$   $N \times N$  matrix, left Schur vectors.  
 $z$   $N \times N$  matrix, right Schur vectors.

**REMARKS** The pair of matrices  $A$  and  $B$  are in generalized real Schur form when  $B$  is upper triangular with non-negative diagonal, and  $A$  is block upper triangular with  $1 \times 1$  and  $2 \times 2$  blocks. The  $1 \times 1$  blocks correspond to real generalized eigenvalues and the  $2 \times 2$  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  $2 \times 2$  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 that reduce  $A$  and  $B$  to Schur form:

$$sa = q'Az$$

$$sb = q'Bz$$

This procedure calls the LAPACK routines DGEYS and ZGEYS.

SOURCE    lapschur.src

lapsvdcusv

PURPOSE    Computes the singular value decomposition of a real or complex rectangular matrix, returns compact  $u$  and  $v$ .

FORMAT    {  $u, s, v$  } = **lapsvdcusv**( $x$ );

INPUT     $x$              $M \times N$  matrix, real or complex rectangular matrix.

OUTPUT     $u$              $M \times \min(M,N)$  matrix, left singular vectors.  
              $s$              $\min(M,N) \times N$  matrix, singular values.  
              $v$              $N \times N$  matrix, right singular values.

REMARKS    **lapsvdcusv** 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. **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     $x = \{ \begin{array}{ccc} 2.143 & 4.345 & 6.124, \\ 1.244 & 5.124 & 3.412, \end{array} \begin{array}{ccc} 0.235 & 5.657 & 8.214 \end{array} \};$

$\{ u, s, v \} = \text{lapsvdcusv}(x);$

print s;

-0.55531277	0.049048431	0.83019394
-0.43090168	0.83684123	-0.33766923
-0.71130266	-0.54524400	-0.44357356

print s;

13.895868	0.00000000	0.00000000
0.00000000	2.1893939	0.00000000
0.00000000	0.00000000	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, lapsvdcusv**

## lapsvds

**PURPOSE** Computes the singular values of a real or complex rectangular matrix

**FORMAT**  $s = \text{lapsvds}(x);$

**INPUT**  $x$   $M \times N$  matrix, real or complex rectangular matrix.

**OUTPUT**  $s$   $\min(M,N) \times 1$  vector, singular values.

**REMARKS** **lapsvd** computes the singular values of a real or complex rectangular matrix. The SVD is

$$x = usv'$$

where  $v$  is the matrix of right singular vectors. For the computation of the singular vectors, see **lapsvdcusv** and **lapsvducsv**.

**lapsvd** is based on the LAPACK drivers DGESVD and ZGESVD. Further documentation of these functions may be found in the LAPACK User's Guide.

**EXAMPLE**  $x = \{ \begin{matrix} 2.143 & 4.345 & 6.124, \\ 1.244 & 5.124 & 3.412, \\ 0.235 & 5.657 & 8.214 \end{matrix} \};$

```
va = lapsvd(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 };
```

## lapsvdusv

---

```
ve = lapsvds(xi);  
print ve;  
  
10.352877 4.0190557 2.3801546
```

SEE ALSO **lapsvdcusv**, **lapsvdusv**

## lapsvdusv

**PURPOSE** Computes the singular value decomposition a real or complex rectangular matrix.

**FORMAT**  $\{ u, s, v \} = \mathbf{lapsvdusv}(x);$

**INPUT**  $x$   $M \times N$  matrix, real or complex rectangular matrix.

**OUTPUT**  $u$   $M \times M$  matrix, left singular vectors.  
 $s$   $M \times N$  matrix, singular values.  
 $v$   $N \times N$  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**  $x = \{ 2.143 \ 4.345 \ 6.124, \dots \}$

```

1.244 5.124 3.412,
0.235 5.657 8.214 };

{ u,s,v } = lapsvdusv(x);

print u;

-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**

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*;

**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:  $\sim$  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;
```

$$x = \begin{pmatrix} 1 & 2 \\ 3 & 4 \end{pmatrix}$$

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 [10.6.2](#), 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";
```

## let

---

```
      age
x =   pay
      sex
```

EXAMPLE    `let x;`

```
x = 0
```

```
let x = { 1 2 3, 4 5 6, 7 8 9 };
```

```
      1 2 3
x =   4 5 6
      7 8 9
```

```
let x[3,3] = 1 2 3 4 5 6 7 8 9;
```

```
      1 2 3
x =   4 5 6
      7 8 9
```

```
let x[3,3] = 1;
```

```
      1 1 1
x =   1 1 1
      1 1 1
```

```
let x[3,3];
```

```
      0 0 0
x =   0 0 0
      0 0 0
```



---

```
let x = 1 2 3 4 5 6 7 8 9;
```

```
      1
      2
      3
      4
x =   5
      6
      7
      8
      9
```

```
let x = dog cat;
```

```
x =  DOG
     CAT
```

```
let x = "dog" "cat";
```

```
x =  dog
     cat
```

```
let string x = { "Median Income", "Country" };
```

```
x =  Median Income
     Country
```

SEE ALSO    **con, cons, declare, load**

## lib

**PURPOSE** Builds and updates library files.

**FORMAT** **lib** *library* [*file*] [*-flag -flag...*];

**INPUT** *library* literal, name of library.

*file* optional literal, name of source file to be updated or added.

*flags* optional literal preceded by ‘-’, controls operation of library update.  
To control handling of path information on source filenames:

**-addpath** (default) add paths to entries without paths  
and expand relative paths.

**-gausspath** reset all paths using a normal file search.

**-leavepath** leave all path information untouched.

**-nopath** drop all path information.

To specify a library update or a complete library build:

**-update** (default) update the symbol information for  
the specified file only.

**-build** update the symbol information for every  
library entry by compiling the actual source  
file.

**-delete** delete a file from the library.

**-list** 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. The

directory will usually be on a RAM disk. 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.

**REMARKS** 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** [**-l**] *lib1* [, *lib2*, *lib3*, *lib4*...];

**library**;

**REMARKS** If no arguments are given, the list of current libraries will be printed out.

The **-l** option will produce a listing of libraries, files, and symbols for all active libraries. This file will reside in the directory defined by the **lib\_path**

configuration variable. The file will have a unique name beginning with `liblst_`.

For more information about the library system, see **LIBRARIES**, Chapter 18.

The default extension for library files is `.lcg`.

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 = dog(x);
```

is encountered in a program, **dog** 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 <code>dog.g</code>
<code>autodelete off</code>	return <b>Undefined symbol</b> error message

If **dog** calls **cat** and **cat** calls **bird** 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 **dog** in exactly the same sequence as the autoloader. The file containing **dog** will be displayed in the window, and you can scroll up and down and look at the code and comments.

Library files are simple ASCII files that you can create with a text editor. Here is an example:

```
/*
```

```
** 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. 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
```

## #lineson, #linesoff

---

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 source browser or help facility 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.

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  $/$  operator is accurate to within approximately  $4\text{e-}13$ .

EXAMPLE     $b = \{ 2, 3, 4 \};$

## listwise (dataloop)

---

```
a = { 10 2 3, 6 14 2, 1 1 9 };  
x = linsolve(b,A);
```

```
print x
```

```
0.045863309  
0.13399281  
0.42446043
```

SEE ALSO **qrsol**, **qrtsol**, **solpd**, **cholsol**

## 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	$y$	$N \times K$ matrix or $N$ -dimensional array containing the natural log values of the elements of $x$ .
REMARKS	<p><b>ln</b> is defined for <math>x \neq 0</math>.</p> <p>If <math>x</math> is negative, complex results are returned.</p> <p>You can turn the generation of complex numbers for negative inputs on or off in the <b>GAUSS</b> configuration file, and with the <b>sysstate</b> function, case 8. If you turn it off, <b>ln</b> will generate an error for negative inputs.</p> <p>If <math>x</math> is already complex, the complex number state doesn't matter; <b>ln</b> will compute a complex result.</p> <p><math>x</math> can be any expression that returns a matrix.</p>	
EXAMPLE	<p><math>y = \ln(16);</math></p> <p><math>y = 2.7725887</math></p>	
SEE ALSO	<b>log</b>	

lncdfbvn

PURPOSE	Computes natural log of bivariate Normal cumulative distribution function.	
FORMAT	$y = \text{lncdfbvn}(x1, x2, r);$	
INPUT	$x1$	$N \times K$ matrix, abscissae.
	$x2$	$L \times M$ matrix, abscissae.

## lncdfbvn2

---

	$r$	P×Q matrix, correlations.
OUTPUT	$y$	$\max(N,L,P) \times \max(K,M,Q)$ matrix, $\ln Pr(X < x1, X < x2 r)$ .
REMARKS	$x1$ , $x2$ , and $r$ must be E×E conformable.	
SOURCE	lncdfn.src	
SEE ALSO	<b>cdfbvn</b> , <b>lncdfmvn</b>	

## lncdfbvn2

**PURPOSE** Returns natural log of standardized bivariate Normal cumulative distribution function of a bounded rectangle.

**FORMAT**  $y = \text{lncdfbvn2}(h, dh, k, dk, r);$

**INPUT**

$h$	N×1 vector, upper limits of integration for variable 1.
$dh$	N×1 vector, increments for variable 1.
$k$	N×1 vector, upper limits of integration for variable 2.
$dk$	N×1 vector, increments for variable 2.
$r$	N×1 vector, correlation coefficients between the two variables.

**OUTPUT**  $y$  N×1 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 N×1 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**( $y$ )±*err*. The size of the error depends on the input arguments. If **trap 2** is set, a warning message is displayed when *err* ≥ **exp**( $y$ )/100.

For an estimate of the actual error, see **cdfbvn2e**.

#### 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:
0.0000e+000 +/- 2.8e-060
-INF
```

## lncdfmvn

---

SEE ALSO **cdfbvn2, cdfbvn2e**

### lncdfmvn

**PURPOSE** Computes natural log of multivariate Normal cumulative distribution function.

**FORMAT**  $y = \text{lncdfmvn}(x, r);$

**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.

**FORMAT**  $y = \text{lncdfn}(x);$

**INPUT**  $x$   $N \times K$  matrix or N-dimensional array, abscissae.

**OUTPUT**  $y$   $N \times K$  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  $x$   $M \times N$  matrix, abscissae.  
 $r$   $K \times L$  matrix,  $E \times E$  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:

$$\begin{array}{llll} |x| \leq 1 & \text{and} & dx \leq 1 & \pm 1e - 14 \\ 1 < |x| < 37 & \text{and} & |dx| < 1/|x| & \pm 1e - 13 \\ \min(x, x + dx) > -37 & \text{and} & y > -690 & \pm 1e - 11 \text{ or better} \end{array}$$

A relative error of  $\pm 1e-14$  implies that the answer is accurate to better than  $\pm 1$  in the  $14^{th}$  digit after the decimal point.

EXAMPLE `print lncdfn2(-10,29);`  
`-7.6198530241605269e-24`  
`print lncdfn2(0,1);`  
`-1.0748623268620716e+00`  
`print lncdfn2(5,1);`  
`-1.5068446096529453e+01`

## lncdfnc

---

SOURCE    `lncdfn.src`

SEE ALSO    `cdfn2`

### lncdfnc

PURPOSE    Computes natural log of complement of Normal cumulative distribution function.

FORMAT    `y = lncdfnc(x);`

INPUT    `x`             $N \times K$  matrix, abscissae.

OUTPUT    `y`             $N \times K$  matrix,  $\ln(1 - \Pr(X < x))$ .

SOURCE    `lncdfn.src`

### lnfact

PURPOSE    Computes the natural log of the factorial function and can be used to compute log gamma.

FORMAT    `y = lnfact(x);`

INPUT    `x`             $N \times K$  matrix or  $N$ -dimensional array, all elements must be positive.

OUTPUT    `y`             $N \times K$  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.

For  $x \geq 1$ , 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:

$$\text{lngamma}(x) = \text{lnfact}(x-1);$$

EXAMPLE    `let x = 100 500 1000;`  
              `y = lnfact(x);`

```

                363.739375560
y =  2611.33045846
      5912.12817849

```

SOURCE    `lnfact.src`

SEE ALSO    **gamma**

TECHNICAL    For  $x > 1$ , Stirling's formula is used.  
 NOTES        For  $0 < x \leq 1$ , **ln(gamma(x+1))** is used.

PURPOSE    Computes multivariate Normal log-probabilities.

## Inpdfmvt

---

FORMAT     $z = \text{lnpdfmvt}(x, s);$

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`

## Inpdfmvt

PURPOSE   Computes multivariate Student's  $t$  log-probabilities.

FORMAT     $z = \text{lnpdfmvt}(x, s, nu);$

INPUT     $x$              $N \times K$  matrix, data.  
           $s$              $K \times K$  matrix, covariance matrix.  
           $nu$           scalar, degrees of freedom.

OUTPUT    $z$              $N \times 1$  vector, log-probabilities.

SOURCE   `lnpdfn.src`

SEE ALSO   **lnpdft**



## Inpdfn

**PURPOSE** Computes standard Normal log-probabilities.

**FORMAT**  $z = \text{lnpdfn}(x);$

**INPUT**  $x$   $N \times K$  matrix or N-dimensional array, data.

**OUTPUT**  $z$   $N \times K$  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:

$$z = -\ln(\text{sqrt}(2*\text{pi})) - x.*x/2;$$

For multivariate log-probabilities, see **lnpdfmvn**.

**EXAMPLE**  $x = \{-2, -1, 0, 1, 2\};$   
 $z = \text{lnpdfn}(x);$

$$z = \begin{matrix} -2.9189385 \\ -1.4189385 \\ -0.91893853 \\ -1.4189385 \\ -2.9189385 \end{matrix}$$

## Inpdfn

**PURPOSE** Computes Student's t log-probabilities.

## load, loadf, loadk, loadm, loadp, loads

---

FORMAT     $z = \text{lnpdf}(x, nu);$

INPUT     $x$              $N \times K$  matrix, data.  
           $nu$            scalar, degrees of freedom.

OUTPUT    $z$              $N \times K$  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 **lnpdfmvt**.

SEE ALSO   **lnpdfmvt**

## 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:

<b>load, loadm</b>	matrix
<b>loads</b>	string
<b>loadf</b>	function ( <b>fn</b> )
<b>loadk</b>	keyword ( <b>keyword</b> )
<b>loadp</b>	procedure ( <b>proc</b> )

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:

<b>load</b>	.fmt - matrix file or delimited ASCII file
<b>loadm</b>	.fmt - matrix file or delimited ASCII file
<b>loads</b>	.fst - string file
<b>loadf</b>	.fcg - user-defined function ( <b>fn</b> ) file
<b>loadk</b>	.fcg - user-defined keyword ( <b>keyword</b> ) file
<b>loadp</b>	.fcg - user-defined procedure ( <b>proc</b> ) 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 corrmatrix; endp;
loadp corrmatrix;
y = corrmatrix;

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.

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  $N \times 1$  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  $N \times 1$  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  $N \times 1$  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, loadf, loadk, loadm, loadp, loads

---

**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**

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 **loadadd, 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

---

```
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.

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**

## load

**PURPOSE** Loads a data set.

**FORMAT** `y = load(dataset)`;

**INPUT** *dataset* string, name of data set.

**OUTPUT** *y* N×K matrix of data.

**REMARKS** The data set must not be larger than a single **GAUSS** matrix.

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*.

**SOURCE** `saveload.src`

## loadstruct

---

GLOBALS    `__maxvec`

### loadstruct

PURPOSE    Loads a structure into memory from a file on the disk.

FORMAT    `{ instance,retcode } = loadstruct(file_name,structure_type);`

INPUT    *file\_name*   string, name of file containing structure.  
          *structure\_type*   string, structure type.

OUTPUT    *instance*    instance of the structure.  
          *retcode*    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

PURPOSE    Load a previously saved graphic panel configuration.

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	<code>pwindow.src</code>	
GLOBALS	<code>_pwindmx</code>	
SEE ALSO	<code>savewind</code>	

**local**

PURPOSE	Declare variables that are to exist only inside a procedure.
FORMAT	<b>local</b> <i>x</i> , <i>y</i> , <i>f</i> : <b>proc</b> ;
REMARKS	<p>The statement above would place the names <i>x</i>, <i>y</i>, and <i>f</i> in the local symbol table for the current procedure being compiled. This statement is legal only between the <b>proc</b> statement and the <b>endp</b> statement of a procedure definition.</p> <p>These symbols cannot be accessed outside of the procedure.</p> <p>The symbol <i>f</i> 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.</p> <p>See PROCEDURES AND KEYWORDS, Chapter 12.</p>
SEE ALSO	<b>proc</b>

## loess

---

### locate

PURPOSE    Positions the cursor in the window.

FORMAT    **locate** *m*, *n*;

PORTABILITY    **Windows** only

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>	N×1 vector, dependent variable.
	<i>indvars</i>	N×K matrix, independent variables.
GLOBAL INPUT	<b>_loess_Span</b>	scalar, degree of smoothing. Must be greater than 2/N. Default = .67777.
	<b>_loess_NumEval</b>	scalar, number of points in <i>ys</i> and <i>xs</i> . Default = 50.
	<b>_loess_Degree</b>	scalar, if 2, quadratic fit, otherwise linear. Default = 1.
	<b>_loess_WgtType</b>	scalar, type of weights. If 1, robust, symmetric weights, otherwise Gaussian. Default = 1.
	<b>__output</b>	scalar, if 1, iteration information and results are printed, otherwise nothing is printed.
OUTPUT	<i>yhat</i>	N×1 vector, predicted <i>depvar</i> given <i>indvars</i> .
	<i>ys</i>	<b>_loess_numEval</b> ×1 vector, ordinate values given abscissae values in <i>xs</i> .
	<i>xs</i>	<b>_loess_numEval</b> ×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	loess.src	

loessmt

PURPOSE	Computes coefficients of locally weighted regression.	
INCLUDE	loessmt.sdf	
FORMAT	{ <i>yhat</i> , <i>ys</i> , <i>xs</i> } = <b>loessmt</b> ( <i>lc0</i> , <i>depvar</i> , <i>indvars</i> );	
INPUT	<i>lc0</i>	an instance of a <b>loessmtControl</b> structure, containing the following members:

## loessmtControlCreate

---

	<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, symmetric weights, otherwise Gaussian. Default = 1.
	<i>lc0.output</i>	scalar, if 1, iteration information and results are printed, otherwise nothing is printed.
	<i>depvar</i>	$N \times 1$ vector, dependent variable.
	<i>indvars</i>	$N \times K$ matrix, independent variables.
OUTPUT	<i>yhat</i>	$N \times 1$ vector, predicted <i>depvar</i> given <i>indvars</i> .
	<i>ys</i>	<i>lc0.numEval</i> $\times 1$ vector, ordinate values given abscissae values in <i>xs</i> .
	<i>xs</i>	<i>lc0.numEval</i> $\times 1$ 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	<code>loessmt.src</code>	
SEE ALSO	<b>loessmtControlCreate</b>	

## loessmtControlCreate

PURPOSE	Creates default <b>loessmtControl</b> structure.
INCLUDE	<code>loessmt.sdf</code>
FORMAT	<code>c = loessmtControlCreate;</code>

---

OUTPUT	<i>c</i>	instance of a <b>loessmtControl</b> structure with members set to default values.
SOURCE	<code>loessmt.src</code>	
SEE ALSO	<b>loessmt</b>	

## log

PURPOSE    Computes the  $\log_{10}$  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 10 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    `x = round(rndu(3,3)*10+1);`  
               `y = log(x);`

## loglog

---

```
          4.0000000000  2.0000000000  1.0000000000
x = 10.0000000000  4.0000000000  8.0000000000
          7.0000000000  2.0000000000  6.0000000000

          0.6020599913  0.3010299957  0.3010299957
y = 1.0000000000  0.6020599913  0.9030899870
          0.8450980400  0.3010299957  0.7781512504
```

SEE ALSO [ln](#)

## loglog

**PURPOSE**    Graphs X vs. Y using log coordinates.

**LIBRARY**    pgraph

**FORMAT**    **loglog(x,y);**

**INPUT**    *x*            N×1 or N×M matrix. Each column contains the X values for a particular line.

*y*            N×1 or N×M 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.

**LIBRARY**    pgraph

**FORMAT**    **logx(x,y);**

**INPUT**    *x*            N×1 or N×M matrix. Each column contains the X values for a particular line.

*y*            N×1 or N×M matrix. Each column contains the Y 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.

**LIBRARY**    pgraph

**FORMAT**    **logy(x,y);**

**INPUT**    *x*            N×1 or N×M matrix. Each column represents the X values for a particular line.

*y*            N×1 or N×M 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 <b>loopnextindex</b> succeeds.
<i>i</i>	M×1 vector of indices into an array, where M≤N.
<i>o</i>	N×1 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** 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**

```
orders = { 2,3,4,5,6,7 };
a = arrayalloc(orders,0);
ind = { 1,1,1,1 };

loopni:

    setarray a, ind, rndn(6,7);
```

```
loopnextindex loopni, ind, orders;
```

This example sets each 6×7 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, rndn(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 6×7 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 N×K matrix of character data to be converted to lowercase.

**OUTPUT** `y` string or N×K 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

## lowmat, lowmat1

---

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

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`            N×N matrix.

**OUTPUT**    `L`            N×N 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.

EXAMPLE    `x = { 1   2 -1,`  
              `2   3 -2,`  
              `1 -2   1 };`

```
L = lowmat(x);
L1 = lowmat1(x);
```

The resulting matrices are

$$L = \begin{pmatrix} 1 & 0 & 0 \\ 2 & 3 & 0 \\ 1 & -2 & 1 \end{pmatrix}$$

$$L1 = \begin{pmatrix} 1 & 0 & 0 \\ 2 & 1 & 0 \\ 1 & -2 & 1 \end{pmatrix}$$

SOURCE `diag.src`

SEE ALSO `upmat`, `upmat1`, `diag`, `diagrv`, `croust`, `croutp`

PURPOSE Computes the solution of  $Lx = b$  where  $L$  is a lower triangular matrix.

FORMAT  $x = \text{ltrisol}(b, L);$

INPUT  $b$   $P \times K$  matrix.  
 $L$   $P \times P$  lower triangular matrix.

OUTPUT  $x$   $P \times K$  matrix, soluion 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  $L*x[:,i] = b[:,i]$ .

**PURPOSE** Computes the LU decomposition of a square matrix with partial (row) pivoting, such that:  $X = LU$ .

**FORMAT**  $\{ l, u \} = \text{lu}(x);$

**INPUT**  $x$   $N \times N$  square nonsingular matrix.

**OUTPUT**  $l$   $N \times N$  “scrambled” lower triangular matrix. This is a lower triangular matrix that has been reordered based on the row pivoting.

$u$   $N \times N$  upper triangular matrix.

**EXAMPLE** `rndseed 13;  
format /rd 10,4;  
x = complex(rndn(3,3),rndn(3,3));  
{ l,u } = lu(x);  
x2 = l*u;`

$$x = \begin{bmatrix} 0.1523 + 0.7685i & -0.8957 + 0.0342i & 2.4353 + 2.7736i \\ -1.1953 + 1.2187i & 1.2118 + 0.2571i & -0.0446 - 1.7768i \\ 0.8038 + 1.3668i & 1.2950 - 1.6929i & 1.6267 + 0.2844i \end{bmatrix}$$

$$l = \begin{bmatrix} 0.2589 - 0.3789i & -1.2417 - 0.5225i & 1.0000 \\ 1.0000 & 0.0000 & 0.0000 \\ 0.2419 - 0.8968i & 1.0000 & 0.0000 \end{bmatrix}$$

$$u = \begin{bmatrix} -1.1953 + 1.2187i & 1.2118 + 0.2571i & -0.0446 - 1.7768i \\ 0.0000 & 0.7713 - 0.6683i & 3.2309 + 0.6742i \\ 0.0000 & 0.0000 & 6.7795 + 5.7420i \end{bmatrix}$$

$$\mathbf{x2} = \begin{pmatrix} 0.1523 + 0.7685i & -0.8957 + 0.0342i & 2.4353 + 2.7736i \\ -1.1953 + 1.2187i & 1.2118 + 0.2571i & -0.0446 - 1.7768i \\ 0.8038 + 1.3668i & 1.2950 - 1.6929i & 1.6267 + 0.2844i \end{pmatrix}$$

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**  $b$  P×K matrix.  
 $L$  P×P lower triangular matrix.  
 $U$  P×P upper triangular matrix.

**OUTPUT**  $x$  P×K matrix, solution of  $LUx = b$ .

**REMARKS** If  $b$  has more than one column, each column is solved for separately, i.e., **lusol** solves  $LUx[:,i] = b[:,i]$ .

machEpsilon

**PURPOSE** Returns the smallest number such that  $1 + eps > 1$ .

**FORMAT**  $eps = \text{machEpsilon};$

**OUTPUT**  $eps$  scalar, machine epsilon.

## make (dataloop)

---

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*;

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 \$ sex = lower(sex);

SEE ALSO    **vector (dataloop)**

### makevars

PURPOSE    Creates separate global vectors from the columns of a matrix.

FORMAT    **makevars**(*x*, *vnames*, *xnames*);



INPUT	<i>x</i>	N×K matrix whose columns will be converted into individual vectors.
	<i>vnames</i>	string or M×1 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 K×1 character vector containing names to be associated with the columns of the matrix <i>x</i> .
REMARKS	<p>If <i>xnames</i> = 0, the prefix X will be used to create names. Therefore, if there are 9 columns in <i>x</i>, the names will be X1-X9, if there are 10, they will be X01-X10, and so on.</p> <p>If <i>xnames</i> or <i>vnames</i> is a string, the individual names must be separated by spaces or commas:</p>	

```
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**.

```
let x[3,3] = 101 35 50000
              102 29 13000
```

## makewind

---

```
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.

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 GRAPHIC PANELS, Section 24.3.

**SOURCE** `pwindow.src`

**SEE ALSO** **window, endwind, setwind, getwind, begwind, nextwind**

## margin

**PURPOSE** Sets the margins for the current graph's graphic panel.

**LIBRARY** `pgraph`

**FORMAT** **margin(*l,r,t,b*);**

<b>INPUT</b>	<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.

## matalloc

---

All input inch values for this procedure are based on a full size window of 9×6.855 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      *r*            scalar, rows.  
            *c*            scalar, columns.

OUTPUT    *y*             $r \times c$  matrix.

REMARKS    The contents are unspecified. This function is used to allocate a matrix that will be 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**      $y = \text{matinit}(r, c, v);$

**INPUT**        $r$             scalar, rows.  
                  $c$             scalar, columns.  
                  $v$             scalar, value to initialize.

**OUTPUT**       $y$              $r \times c$  matrix with each element equal to the value of  $v$ .

**SEE ALSO**     **matalloc, ones, zeros, eye**

## mattoarray

**PURPOSE**     Converts a matrix to a type array.

**FORMAT**      $y = \text{mattoarray}(x);$

**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**      $x = 5 * \text{ones}(2, 3);$   
                  $y = \text{mattoarray}(x);$   
  
                  $y$  will be a  $2 \times 3$  array of fives.

## maxc

---

SEE ALSO **arraytomat**

### maxc

**PURPOSE** Returns a column vector containing the largest element in each column of a matrix.

**FORMAT**  $y = \text{maxc}(x);$

**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**:

$$y = \text{maxc}(\text{maxc}(x));$$

**EXAMPLE**  $x = \text{rndn}(4,2);$   
 $y = \text{maxc}(x);$

$$x = \begin{bmatrix} -2.124474 & 1.376765 \\ 0.348110 & 1.172391 \\ -0.027064 & 0.796867 \\ 1.421940 & -0.351313 \end{bmatrix}$$

$$y = \begin{matrix} 1.421940 \\ 1.376765 \end{matrix}$$

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.

**FORMAT**  $y = \text{maxindc}(x);$

**INPUT**  $x$   $N \times K$  matrix.

**OUTPUT**  $y$   $K \times 1$  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**.

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**  $x = \text{round}(\text{rndn}(4,4)*5);$   
 $y = \text{maxc}(x);$   
 $z = \text{maxindc}(x);$

## maxv

---

$$\mathbf{x} = \begin{bmatrix} 1 & -11 & 0 & 5 \\ 0 & 0 & -2 & -6 \\ -8 & 0 & 3 & 2 \\ -11 & 5 & -4 & 5 \end{bmatrix}$$

$$\mathbf{y} = \begin{bmatrix} 1 \\ 5 \\ 3 \\ 5 \end{bmatrix}$$

$$\mathbf{z} = \begin{bmatrix} 1 \\ 4 \\ 3 \\ 1 \end{bmatrix}$$

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**  $z = \mathbf{maxv}(x,y);$

<b>GLOBAL</b>	$x$	N×K matrix
<b>INPUT</b>	$y$	N×K matrix

**OUTPUT**  $z$  A N×K 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 x = rndn(10,10);
        y = rndn(10,10);
```

```
        z = maxv(x,y);
\end{verbatim}
```

```
\See      \commandname{minv}
```

```
\endcr\CR{maxvec}
\index{maxvec#@\commandname{maxvec}}
```

```
\Purpose
```

Returns maximum vector length allowed.

```
\Format  \textcode{\substitute{y} = maxvec;}
```

```
\GInput
```

```
\begin{commandlist}
```

```
\item[\_\_maxvec] scalar, maximum vector length allowed.
```

```
\end{commandlist}
```

```
\Output
```

```
\begin{argumentlist}
```

```
\item[y] scalar, maximum vector length.
```

```
\end{argumentlist}
```

```
\Remarks
```

`\commandname{maxvec}` returns the value in the global scalar `\commandname{\_\_maxvec}`, which can be reset in the calling program.

`\commandname{maxvec}` is called by `\rtl{}` functions and applications when determining how many rows can be read from a data set in one call to `\commandname{readr}`.

## maxbytes

---

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
\begin{verbatim}
    y = maxvec;
    print y;

                20000.000
```

SOURCE    system.src

## maxbytes

PURPOSE    Returns maximum memory to be used.

FORMAT    `y = maxbytes;`

GLOBAL    `__maxbytes`    scalar, maximum memory to be used.  
INPUT

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;
```

```
1000000000.000
```

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.
$n$	scalar, highest order.
$alpha$	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  $y$ :

$$y[i, :] = I_\alpha(x[i]) \quad I_{\alpha+1}(x[i]) \quad \cdots \quad I_{\alpha+n-1}(x[i])$$

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:

$$\text{mbesseli0}(x) = \exp(-x) * \text{mbesseli0}(x)$$

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  $\pi$  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.

$\kappa$  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);
    retp(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)]);
    retp(rows(dev)*ln(mbesseli0(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));
```

## meanc

---

SOURCE    `ribes1.src`

### meanc

PURPOSE    Computes the mean of every column of a matrix.

FORMAT    `y = meanc(x);`

INPUT      `x`             $N \times K$  matrix.

OUTPUT    `y`             $K \times 1$  matrix containing the mean of every column of  $x$ .

EXAMPLE    `x = meanc(rndu(2000,4));`

```

           0.492446
           0.503543
x =        0.502905
           0.509283
```

In this example, 4 columns of uniform random numbers are generated in a matrix, and the mean is computed for each column.

SEE ALSO    `stdc`

### median

PURPOSE    Computes the medians of the columns of a matrix.

FORMAT    `m = median(x);`

INPUT	$x$	$N \times K$ matrix.
OUTPUT	$m$	$K \times 1$ vector containing the medians of the respective columns of $x$ .
EXAMPLE	$x = \begin{Bmatrix} 8 & 4, \\ 6 & 8, \\ 3 & 7 \end{Bmatrix};$ $y = \text{median}(x);$ $y = \begin{Bmatrix} 6.0000000 \\ 7.0000000 \end{Bmatrix}$	
SOURCE	median.src	

mergeby

PURPOSE	Merges two sorted files by a common variable.		
FORMAT	<b>mergeby</b> ( <i>infile1</i> , <i>infile2</i> , <i>outfile</i> , <i>keytyp</i> );		
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:		

## mergevar

---

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 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**     $x = \text{mergevar}(vnames);$

**INPUT**     $vnames$     string or  $K \times 1$  column vector containing the names of  $K$  global matrices.

**OUTPUT**     $x$      $N \times M$  matrix that contains the concatenated matrices, where  $M$  is the sum of the columns in the  $K$  matrices specified in  $vnames$ .

**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    `let vnames = age pay sex;  
              x = mergevar(vnames);`

The matrices **age**, **pay** and **sex** will be concatenated horizontally to create **x**.

SOURCE    `vars.src`

SEE ALSO    **makevars**

## 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 = rndn(4,2);  
              y = minc(x);`

## minindc

---

$$\mathbf{x} = \begin{array}{rr} -1.061321 & -0.729026 \\ -0.021965 & 0.184246 \\ 1.843242 & -1.847015 \\ 1.977621 & -0.532307 \end{array}$$
$$\mathbf{y} = \begin{array}{r} -1.061321 \\ -1.847015 \end{array}$$

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**  $y = \text{minindc}(x);$

**INPUT**  $x$   $N \times K$  matrix.

**OUTPUT**  $y$   $K \times 1$  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**.

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**  $x = \text{round}(\text{rndn}(5,4)*5);$

```
y = minc(x);
z = minindc(x);
```

```
      -5   6  -4  -1
      2  -2   1   3
x =   6   0   1  -7
     -6   0   8  -4
      7  -4   8   3
```

```
      -6
      -4
y =   -4
      -7
```

```
      4
      5
z =   1
      3
```

SEE ALSO **maxindc**, **minc**, **maxc**

## 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**  $y = \text{miss}(x, v);$   
 $y = \text{missrv}(x, v);$

**INPUT**  $x$   $N \times K$  matrix.

## miss, missrv

---

	$v$	$L \times M$ matrix, $E \times E$ conformable with $x$ .
OUTPUT	$y$	$\max(N,L)$ by $\max(K,M)$ matrix.
REMARKS	For <b>miss</b> , elements in $x$ that are equal to the corresponding elements in $v$ will be replaced with the <b>GAUSS</b> 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.

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**, **issmiss**, **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.

EXAMPLE     $v = -1 \sim 4 \sim 5;$   
               $y = \text{miss}(x,v);$

In this example, **x** must have 3 columns. All -1's in the first column will be changed to missings, along with all 4's in the second column and 5's in the third column.

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,e);`

**INPUT** *x* N×K matrix.  
*e* N×K logical matrix (matrix of 0's and 1's) that serves as a “mask” for *x*; the 1's in *e* correspond to the values in *x* that are to be converted into missing values.

**OUTPUT** *y* N×K 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.

## minv

---

This function is like **miss**, but is more general in that a range of values can be converted into missings.

EXAMPLE    `x = rndu(3,2);`  
             `/* logical expression */`  
             `e = (x .> .10) .and (x .< .20);`  
             `y = missex(x,e);`

A 3×2 matrix of uniform random numbers is created. All values in the interval (0.10, 0.20) are converted to missing.

SOURCE    `datatran.src`

SEE ALSO    **miss**, **missrv**

## 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	<i>x</i>	N×K matrix
INPUT	<i>y</i>	N×K matrix

OUTPUT    *z*            A N×K matrix whose values are the minimum of each element from the arguments *x* and *y*.

REMARKS    **maxv** works for sparse matrices as well as arrays.

EXAMPLE    `x = rndn(10,10);`  
             `y = rndn(10,10);`

```
z = minv(x,y);
\end{vebatim}
```

```
\See      \commandname{maxv}
```

```
\endcr\CR{moment}
\index{moment#@\commandname{moment}}
```

```
\Purpose
```

Computes a cross-product\index{cross-product} matrix. This is the same as  $\text{\substitute{x}}^{\prime}\text{\substitute{x}}$ .

```
\Format    \textcode{\substitute{y} = moment(\substitute{x},\substitute{d})}
```

```
\Input
```

```
    \begin{argumentlist}
    \item[x]  N$\times$K matrix or M-dimensional array where
the last two dimensions are N$\times$K.
    \item[d]  scalar, controls handling of missing values.
```

```
\begin{List}
```

```
\item[0] missing values will not be checked for.
This is the fastest option.
```

```
\item[1] ‘‘listwise deletion\index{deletion}\index{listwise dele}
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 \substitute{x} contains missing
values, then \textcode{moment(\substitute{x},1)} will return a s
```

```
\item[2] ‘‘pairwise deletion\index{pairwise deletion}’’ is used.
Any element of \substitute{x} that is missing is excluded from
the computation of the moment matrix\index{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.
    \end{List}
\end{argumentlist}

\Output
    \begin{argumentlist}
    \item[y]  $K \times K$  matrix or M-dimensional array where
the last two dimensions are  $K \times K$ , the cross-product
of  $\text{\textcode{\substitute{x}}}$ .
    \end{argumentlist}

\Remarks

The fact that the moment
matrix is symmetric is taken into account to cut
execution time almost in half.

If  $\text{\textcode{\substitute{x}}}$  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  $\text{\textcode{\substitute{x}}}$ . In other words, for a
 $10 \times 4 \times 4$  array  $\text{\textcode{\substitute{x}}}$ , the resulting array
 $\text{\textcode{\substitute{y}}}$  will contain the cross-products of each of the 10
 $4 \times 4$  arrays contained in  $\text{\textcode{\substitute{x}}}$ , so
 $\text{\textcode{\substitute{y}}[\text{\textcode{\substitute{n}}},..]} = \text{\textcode{\substitute{x}}[\text{\textcode{\substitute{n}}}]}$ ,
for  $1 \leq n \leq 10$ .

If there is no missing data then  $\text{\textcode{\substitute{d}}} = 0$  should be used
because it will be faster.

The  $\text{\textcode{\commandname{/}}}$  operator (matrix division) will automatically
form a moment matrix (performing pairwise
deletions if  $\text{\textcode{\textcode{trap 2}}}$  is set) and will compute the  $\text{\textcode{\commandname{ol}}}$ 
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  $\text{\textcode{\commandname{/}}}$  operator is

```



used.

```
\Example
\begin{verbatim}
    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**       *dataset*     string, name of data set.

*vars*        K×1 character vector, names of variables

                 - or -

                 K×1 numeric vector, indices of columns.

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**     **\_\_con**            scalar, default 1.

**INPUT**        **1**        a constant term will be added.

**0**        no constant term will be added.

**\_\_miss**       scalar, default 0.

## movingave

---

- 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 *m*.

**\_\_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, 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*

M×M matrix, where  $M = K + \text{__con}$ , 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

EXAMPLE 

```
z = { age, pay, sex };  
m = momentd("freq",z);
```

SOURCE `momentd.src`

## movingave

PURPOSE Computes moving average of a series.

FORMAT `y = movingave(x,d);`

INPUT	$x$	$N \times K$ matrix.
	$d$	scalar, order of moving average.
OUTPUT	$y$	$N \times K$ matrix, filtered series. The first $d-1$ rows of $x$ are set to missing values.
REMARKS	<b>movingave</b> is essentially a smoothing time series filter. The moving average as performed by column and thus it treats the $N \times K$ matrix as $K$ time series of length $N$ .	
SEE ALSO	<b>movingaveWgt</b> , <b>movingaveExpwgt</b>	

## 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	<b>movingaveExpwgt</b> is smoothing time series filter using exponential weights. The moving average as performed by column and thus it treats the $N \times K$ matrix as $K$ time series of length $N$ .	
SEE ALSO	<b>movingaveWgt</b> , <b>movingave</b>	

## movingaveWgt

PURPOSE	Computes weighted moving average of a series	
FORMAT	$y = \text{movingaveWgt}(x, d, w);$	
INPUT	$x$	$N \times K$ matrix.
	$d$	scalar, order of moving average.
	$w$	$d \times 1$ vector, weights.
OUTPUT	$y$	$N \times K$ matrix, filtered series. The first $d-1$ rows of $x$ are set to missing values.
REMARKS	<b>movingaveWgt</b> is essentially a smoothing time series filter with weights. The moving average as performed by column and thus it treats the $N \times K$ matrix as $K$ time series of length $N$ .	
SEE ALSO	<b>movingave</b> , <b>movingaveExpwgt</b>	

## msym

PURPOSE	Allows the user to set the symbol that <b>GAUSS</b> uses when missing values are converted to ASCII and vice versa.	
FORMAT	<b>msym</b> <i>str</i> ;	
INPUT	<i>str</i>	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.

SEE ALSO    **print, printfm**

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** *[[nos]]* *[[, mps]]*;

INPUT    *nos*       scalar, which indicates the maximum number of global symbols allowed.

*mps*       scalar, which indicates the number of bytes of main program space to be allocated. The second argument is obsolete, and included only for backwards compatibility. Memory is dynamically allocated.

REMARKS    Procedures, user-defined functions, and global matrices, strings, and string arrays are all global symbols.

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

## nextindex

---

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*                        M×1 vector of indices into an array, where M≤N.  
             *o*                        N×1 vector of orders of an N-dimensional array.

OUTPUT    *ni*                        M×1 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    `a = ones(2520,1);`  
             `a = areshape(a,3|4|5|6|7);`  
             `orders = getorders(a);`

```
ind = { 2,3,5 };
ind = nextindex(ind,orders);
```

```
      2
ind = 4
      1
```

n

In this example, **nextindex** incremented **ind** to index the next 6×7 subarray in array **a**.

SEE ALSO **previousindex, loopnextindex, walkindex**

nextn, nextnevn

PURPOSE	Returns allowable matrix dimensions for computing FFT's.	
FORMAT	$n = \text{nextn}(n0);$ $n = \text{nextnevn}(n0);$	
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. $n \geq n0$ .
REMARKS	<b>nextn</b> and <b>nextnevn</b> 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:	

$$2^p \times 3^q \times 5^r \times 7^s, \quad \begin{matrix} p,q,r \text{ nonnegative integers} \\ s = 0 \text{ or } 1 \end{matrix}$$

## nextwind

---

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 Function	Vector Length	Matrix Rows	Matrix Columns
<b>fftn</b>	<b>nextn</b>	<b>nextn</b>	<b>nextn</b>
<b>rfftn</b>	<b>nextnevn</b>	<b>nextn</b>	<b>nextnevn</b>
<b>rfftnp</b>	<b>nextnevn</b>	<b>nextn</b>	<b>nextnevn</b>

EXAMPLE    `n = nextn(456);`

`n = 480.00000`

SOURCE    `optim.src`

SEE ALSO    **fftn**, **optn**, **optnevn**, **rfftn**, **rfftnp**

## nextwind

PURPOSE    Set the current graphic panel to the next available graphic panel.

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 GRAPHIC PANELS, Section 24.3.

SOURCE     pwindow.src

SEE ALSO     **endwind, begwind, setwind, getwind, makewind, window**

null

PURPOSE     Computes an orthonormal basis for the (right) null space of a matrix.

FORMAT      $b = \text{null}(x);$

INPUT      $x$              $N \times M$  matrix.

OUTPUT      $b$              $M \times K$  matrix, where  $K$  is the nullity of  $x$ , such that:

$$x * b = 0 \qquad (N \times K \text{ matrix of zeros})$$

and

$$b' b = I \qquad (M \times M \text{ identity matrix})$$

The error returns are returned in  $b$ :

error code	reason
1	there is no null space
2	$b$ is too large to return in a single matrix

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  $x$ .

## null1

---

EXAMPLE    `let x[2,4] = 2 1 3 -1  
                              3 5 1 2;`

`b = null(x);  
z = x*b;  
i = b'b;`

SOURCE    `null.src`

GLOBALS    `_qrdc, _qrs1`

## null1

PURPOSE    Computes an orthonormal basis for the (right) null space of a matrix.

FORMAT    `nu = null1(x,dataset);`

INPUT      *x*            N×M matrix.  
            *dataset*    string, the name of a data set **null1** will write.

OUTPUT    *nu*            scalar, the nullity of *x*.

REMARKS    **null1** computes an M×K matrix *b*, where K is the nullity of *x*, such that:

$$x * b = 0 \quad (\text{N} \times \text{K matrix of zeros})$$

and

$$b'b = I \quad (\text{M} \times \text{M identity matrix})$$

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.

SOURCE    `null.src`

GLOBALS    `_qrdc, _qrs1`

n

numCombinations

PURPOSE    Computes number of combinations of  $n$  things taken  $k$  at a time.

FORMAT    `y = numCombinations( $n,k$ );`

INPUT       $n$             scalar.  
              $k$             scalar.

OUTPUT     $y$             scalar, number of combinations of  $n$  things take  $k$  at a time.

EXAMPLE    `y = numCombinations(25,5);`  
  
             `print y;`  
  
             53130.0000

SEE ALSO    `combineate, combined`

ols

PURPOSE    Computes a least squares regression.

---

FORMAT    { *vnam*,*m*,*b*,*stb*,*vc*,*stderr*,*sigma*,*cx*,*rsq*,*resid*,*dwstat* } =  
**ols**(*dataset*,*depvar*,*indvars*);

INPUT    *dataset*    string, name of data set or null string.  
                       If *dataset* is a null string, the procedure assumes that the actual data  
                       has been passed in the next two arguments.

*depvar*    If *dataset* 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 *dataset* is a null string or 0:  
                               N×1 vector, the dependent variable.

*indvars*    If *dataset* contains a string:  
                               K×1 character vector, names of independent variables  
                               - or -  
                               K×1 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:  
                               N×K matrix, the independent variables.

GLOBAL    Defaults are provided for the following global input variables, so they can be  
 INPUT    ignored unless you need control over the other options provided by this  
           procedure.

**\_\_altnam**        character vector, default 0.  
                               This can be a (K+1)×1 or (K+2)×1 character vector of  
                               alternate variable names for the output. If **\_\_con** is 1, this  
                               must be (K+2)×1. The name of the dependent variable is the  
                               last element.

**\_\_con**            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  $m$ .

**\_\_miss**

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.

**\_\_output**

scalar, default 1.

**1** print the statistics.

**0** do not print statistics.

**\_\_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 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.

**\_olsres**

scalar, default 0.

**1** compute residuals (*resid*) and Durbin-Watson statistic (*dwstat*).

**0** *resid* = 0, *dwstat* = 0.

OUTPUT    *vnam*     $(K+2) \times 1$  or  $(K+1) \times 1$  character vector, the variable names used in the regression. If a constant term is used, this vector will be  $(K+2) \times 1$ , and the first name will be "CONSTANT". The last name will be the name of the dependent variable.

o

*m* M×M 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	<i>indvars</i>	<i>depvar</i>
(constant)	(independent variables)	(dependent variable)

A constant term is always used in computing *m*.

*b* D×1 vector, the least squares estimates of parameters  
Error handling is controlled by the low order bit of the trap flag.

<b>trap 0</b>	terminate with error message
<b>trap 1</b>	return scalar error code in <i>b</i>
<b>30</b>	system singular
<b>31</b>	system underdetermined
<b>32</b>	same number of columns as rows
<b>33</b>	too many missings
<b>34</b>	file not found
<b>35</b>	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.

*stb* K×1 vector, the standardized coefficients.

*vc* D×D matrix, the variance-covariance matrix of estimates.

*stderr* D×1 vector, the standard errors of the estimated parameters.

*sigma* scalar, standard deviation of residual.

*cx* (K+1)×(K+1) matrix, correlation matrix of variables with the dependent variable as the last column.

*rsq* scalar, R square, coefficient of determination.

*resid* 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 N×1 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  $N \times 1$  vector of residuals.

*dwstat* scalar, Durbin-Watson statistic.

**REMARKS** 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**.

**EXAMPLE**

```
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;
```

## olsmt

---

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.

SOURCE    `ols.src`

SEE ALSO    **olsqr**

## olsmt

PURPOSE    Computes a least squares regression.

FORMAT    `oout = olsmt(oc0,dataset,depvar,indvars);`

INPUT	<i>oc0</i>	instance of an <b>olsmtControl</b> structure containing the following members:
	<i>oc0.altnam</i>	character vector, default 0. This can be a $(K+1) \times 1$ or $(K+2) \times 1$ character vector of alternate variable names for the output. If <i>oc0.con</i> is 1, this must be $(K+2) \times 1$ . The name of the dependent variable is the last element.
	<i>oc0.con</i>	scalar, default 1. <b>1</b> a constant term will be added, $D = K+1$ . <b>0</b> no constant term will be added, $D = K$ . A constant term will always be used in constructing the moment matrix <i>m</i> .
	<i>oc0.miss</i>	scalar, default 0. <b>0</b> 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.

*oc0.vpad*

scalar, default 1.

If 0, internally created variable names are not padded to the same length (e.g. “X1, X2,..., X10”).

If 1, they are padded with zeros to the same length (e.g., “X01, X02,..., X10”).

*oc0.output*

scalar, default 1.

- 1 print the statistics.
- 0 do not print statistics.

*oc0.res*

scalar, default 0.

- 1 compute residuals (*resid*) and Durbin-Watson statistic (*dwstat*).
- 0 *oout.resid* = 0, *oout.dwstat* = 0.

*oc0.rnam*

string, default “\_olsmtres”.

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> .
	<i>oc0.maxvec</i>	scalar, default 20000. The largest number of elements allowed in any one matrix.
	<i>oc0.fcmtol</i>	scalar, default 1e-12. Tolerance used to fuzz the comparison operations to allow for round off error.
<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: N×1 vector, the dependent variable.	
<i>indvars</i>	If <i>dataset</i> contains a string: K×1 character vector, names of independent variables - or - K×1 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 <i>dataset</i> is a null string or 0: N×K matrix, the independent variables.	
OUTPUT	<i>oout</i>	instance of an <b>olsmtOut</b> structure containing the following members:  <i>oout.vnam</i> (K+2)×1 or (K+1)×1 character vector, the variable names used in the regression. If a constant term is used, this vector will be (K+2)×1, and the first name will be

	“CONSTANT”. The last name will be the name of the dependent variable.																
<i>oout.m</i>	M×M 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: <table><tr><td>1.0</td><td><i>indvars</i></td><td><i>depvar</i></td></tr><tr><td>constant</td><td>independent variables</td><td>dependent variables</td></tr></table> A constant term is always used in computing $m$ .	1.0	<i>indvars</i>	<i>depvar</i>	constant	independent variables	dependent variables										
1.0	<i>indvars</i>	<i>depvar</i>															
constant	independent variables	dependent variables															
<i>oout.b</i>	D×1 vector, the least squares estimates of parameters Error handling is controlled by the low order bit of the trap flag. <table><tr><td><b>trap 0</b></td><td>terminate with error message</td></tr><tr><td><b>trap 1</b></td><td>return scalar error code in <math>b</math></td></tr><tr><td><b>30</b></td><td>system singular</td></tr><tr><td><b>31</b></td><td>system underdetermined</td></tr><tr><td><b>32</b></td><td>same number of columns as rows</td></tr><tr><td><b>33</b></td><td>too many missings</td></tr><tr><td><b>34</b></td><td>file not found</td></tr><tr><td><b>35</b></td><td>no variance in an independent variable</td></tr></table> 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.	<b>trap 0</b>	terminate with error message	<b>trap 1</b>	return scalar error code in $b$	<b>30</b>	system singular	<b>31</b>	system underdetermined	<b>32</b>	same number of columns as rows	<b>33</b>	too many missings	<b>34</b>	file not found	<b>35</b>	no variance in an independent variable
<b>trap 0</b>	terminate with error message																
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<b>30</b>	system singular																
<b>31</b>	system underdetermined																
<b>32</b>	same number of columns as rows																
<b>33</b>	too many missings																
<b>34</b>	file not found																
<b>35</b>	no variance in an independent variable																
<i>oout.stb</i>	K×1 vector, the standardized coefficients.																
<i>oout.vc</i>	D×D matrix, the variance-covariance matrix of estimates.																

<code>oout.st derr</code>	D×1 vector, the standard errors of the estimated parameters.
<code>oout.sigma</code>	scalar, standard deviation of residual.
<code>oout.cx</code>	(K+1)×(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, <code>oout.resid = y - x * oout.b</code> . 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 N×1 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 N×1 vector of residuals.
<code>oout.dwstat</code>	scalar, Durbin-Watson statistic.

REMARKS 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**.

EXAMPLE

```
#include olsmt.sdf

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**.

```

#include olsmt.sdf

struct olsmtControl oc0;
struct olsmtOut oOut;
oc0 = olsmtControlCreate;

data = "olsdat";
depvar = { score };
indvars = { region,age,marstat };
oc0.res = 1;
output file = lpt1 on;
oOut = olsmt(oc0,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.

## olsmtControlCreate

---

SOURCE    `olsmt.src`

SEE ALSO    `olsmtControlCreate`, `olsqrmt`

### 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.

SOURCE    `olsmt.src`

SEE ALSO    `olsmt`

### olsqr

PURPOSE    Computes OLS coefficients using QR decomposition.

FORMAT    `b = olsqr(y,x);`

INPUT	<code>y</code>	N×1 vector containing dependent variable.
	<code>x</code>	N×P matrix containing independent variables.

GLOBAL INPUT	<code>_olsqtol</code>	scalar, the tolerance for testing if diagonal elements are approaching zero. The default value is $10^{-14}$ .
-----------------	-----------------------	--

OUTPUT	<i>b</i>	P×1 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	<p>This provides an alternative to <i>y/x</i> for computing least squares coefficients.</p> <p>This procedure is slower than the <i>/</i> operator. However, for near singular matrices it may produce better results.</p> <p><b>olsqr</b> handles matrices that do not have full rank by returning zeros for the coefficients that cannot be estimated.</p>	
SOURCE	olsqr.src	
SEE ALSO	<b>ols</b> , <b>olsqr2</b> , <b>orth</b> , <b>qqr</b>	

olsqr2

PURPOSE	Computes OLS coefficients, residuals, and predicted values using the QR decomposition.	
FORMAT	{ <i>b,r,p</i> } = <b>olsqr2</b> ( <i>y,x</i> );	
INPUT	<i>y</i>	N×1 vector containing dependent variable.
	<i>x</i>	N×P matrix containing independent variables.
GLOBAL INPUT	<b>_olsqtol</b>	scalar, the tolerance for testing if diagonal elements are approaching zero. The default value is 10 <sup>-14</sup> .
OUTPUT	<i>b</i>	P×1 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.
	<i>r</i>	P×1 vector of residuals. ( <i>r</i> = <i>y</i> - <i>x</i> * <i>b</i> )

## olsqrmt

---

$p$              $P \times 1$  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.

SOURCE    `olsqr.src`

SEE ALSO    **olsqr**, **orth**, **qqr**

## olsqrmt

PURPOSE    Computes OLS coefficients using QR decomposition.

FORMAT     $b = \text{olsqrmt}(y, x, tol);$

INPUT       $y$              $N \times 1$  vector containing dependent variable.  
              $x$              $N \times P$  matrix containing independent variables.  
              $tol$           scalar, the tolerance for testing if diagonal elements are approaching zero. The default value is  $10^{-14}$ .

OUTPUT     $b$              $P \times 1$  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**

o

ones

PURPOSE    Creates a matrix of ones.

FORMAT    `y = ones(r,c);`

INPUT    *r*            scalar, number of rows.  
          *c*            scalar, number of columns.

OUTPUT    *y*             $r \times c$  matrix of ones.

REMARKS    Noninteger arguments will be truncated to an integer.

EXAMPLE    `x = ones(3,2);`

```
           1.000000  1.000000
x = 1.000000  1.000000
     1.000000  1.000000
```

SEE ALSO    **zeros, eye**

## open

**PURPOSE** Opens an existing **GAUSS** data file.

**FORMAT** **open** *fh=filename* **[[for mode]]** **[[w32]]** **[[varindxi** *[[offs]]* **]]**;

**INPUT** *filename* literal or ^string.

*filename* 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.

*mode* literal, the modes supported with the optional **for** subcommand are:

- |               |   |
|---------------|---|
| <b>read</b>   | 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 <b>writer</b> function is disabled for files opened in this way. This is the only mode available for matrix files ( <b>.fmt</b> ), which are always written in one piece with the <b>save</b> command. |
| <b>append</b> | 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 <b>writer</b> function will add data to the end of the file without overwriting any of the existing data in the file. The <b>readr</b> function is disabled for files opened in this way. This mode is used to add additional rows to the end of a file.   |
| <b>update</b> | 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 **-w32** flag is an optimization for Windows. It is ignored on all other platforms. **GAUSS** 7.0 uses 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. 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.

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 **varindxi** 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    *fh*            scalar, file handle.  
*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 **open** 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 = c:dat1 varindx;  
open q2 = c: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;
i = 1; rf = rowsf(fx);
sampsiz = round(rf*0.1);
rndsmpx = zeros(sampsiz,colsf(fx));
do until i > sampsiz;
    r = ceil(rndu(1,1)*rf);
    call seekr(fx,r);
    rndsmpx[i,.] = readr(fx,1);
    i = i+1;
endo;
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  $n = \text{optn}(n0);$

## optn, optnevn

---

$n = \text{optnevn}(n0);$

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.  $n \geq n0$ .

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:

$$2^p \times 3^q \times 5^r \times 7^s, \quad p, q, r \text{ nonnegative integers} \\ s=0 \text{ or } 1$$

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 33600×1 vector can compute as much as 20% faster than a 32768×1 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,  $2^{15}$ . **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 Function	Vector Length	Matrix Rows	Matrix Columns
<b>fftn</b>	<b>optn</b>	<b>optn</b>	<b>optn</b>
<b>rfftn</b>	<b>optnevn</b>	<b>optn</b>	<b>optnevn</b>
<b>rfftnp</b>	<b>optnevn</b>	<b>optn</b>	<b>optnevn</b>



EXAMPLE `n = optn(231);`

`n = 240.00000`

SEE ALSO `fftn, nextn, nextnevn, rfftn, rfftnp`

o

## orth

PURPOSE Computes an orthonormal basis for the column space of a matrix.

FORMAT `y = orth(x);`

INPUT `x`  $N \times K$  matrix.

GLOBAL INPUT `_orthtol` scalar, the tolerance for testing if diagonal elements are approaching zero. The default is  $1.0e-14$ .

OUTPUT `y`  $N \times L$  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);`

$$y = \begin{bmatrix} -0.58123819 & -0.81373347 \\ -0.81373347 & 0.58123819 \end{bmatrix}$$

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]] [[on|off|reset]];**

**INPUT** *filename* literal or ^string.

The **file=filename** 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 **output.out**.

**on, off, reset** literal, mode flag:

**on** opens the auxiliary output file or device and causes the results of all **print** 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.

**off** closes the auxiliary output file and turns off the auxiliary output.

**reset** similar to the **on** 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);;  
enddo;  
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**

o

outtyp (dataloop)

PURPOSE    Specifies the precision of the output data set.

FORMAT    **outtyp** *num\_constant*;

INPUT      *num\_constant*    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 precison 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      *n*                scalar, width of auxilary output.

## pacf

---

- 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 80 columns.
- EXAMPLE**    `outwidth 132;`
- This statement will change the auxiliary output width to 132 columns.
- SEE ALSO**    **output, print**

## pacf

- PURPOSE**    Computes sample partial autocorrelations.
- FORMAT**    `rkk = pacf(y,k,d);`
- INPUT**       *y*            N×1 vector, data.  
              *k*            scalar, maximum number of partial autocorrelations to compute.  
              *d*            scalar, order of differencing.
- OUTPUT**    *rkk*           K×1 vector, sample partial autocorrelations.
- EXAMPLE**    `proc pacf(y,k,d);`  
              `local a,l,j,r,t;`  
              `r = acf(y,k,d);`

```

a = zeros(k,k);
a[1,1] = r[1];

t = 1;
l = 2;

do while l le k;
    a[l,1] = (r[l]-a[l-1,1:t]*rev(r[1:l-1]))/
              (1-a[l-1,1:t]*r[1:t]);
    j = 1;

    do while j <= t;
        a[l,j] = a[l-1,j] - a[l,1]*a[l-1,l-j];
        j = j+1;
    endo;

    t = t+1;
    l = l+1;
endo;

retp(diag(a));
endp;

```

SOURCE    tsutil.src

**packedToSp**

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

**FORMAT**     $y = \text{packedToSp}(r, c, p);$

## packedToSp

---

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 complex.

The format for  $p$  is as follows:

If  $p$  is  $N \times 3$ :

Column 1	Column 2	Column 3
non-zero values	row indices	column indices

If  $p$  is  $N \times 4$ :

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   `sparse matrix y;`  
              `p = { 1 2 4, 2 5 1, 3 8 9, 4 13 5 };`  
              `y = packedToSp(15,10,p);`

This example creates a  $15 \times 10$  sparse matrix **y**, containing the following non-zero values:



Non-zero value	Index
1	(2,4)
2	(5,1)
3	(8,9)
4	(13,5)

SEE ALSO **spCreate, denseToSp**

p

packr

- PURPOSE

Deletes the rows of a matrix that contain any missing values.
- FORMAT

`y = packr(x);`
- INPUT

`x`

N×K matrix.
- OUTPUT

`y`

L×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 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

```
x = miss(ceil(rndu(3,3)*10),1);
y = packr(x);
```

$$\mathbf{x} = \begin{matrix} & . & 9 & 10 \\ 4 & 2 & & . \\ 3 & 4 & 9 & \end{matrix}$$
$$\mathbf{y} = \begin{matrix} 3 & 4 & 9 \end{matrix}$$

## parse

---

In this example, the matrix **x** is formed with random integers and missing values. **packr** is used to delete rows with missing values.

```
open fp = mydata;
obs = 0;
sum = 0;
do until eof(fp);
    x = packr(readr(fp,100));
    if not scalmiss(x);
        obs = obs+rows(x);
        sum = sum+sumc(x);
    endif;
endo;
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*);

**INPUT**       *str*            string consisting of a series of tokens and/or delimiters.  
              *delim*        N×K character matrix of delimiters that might be found in *str*.

OUTPUT	<i>tok</i>	M×1 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 <i>str</i> must be 8 characters or less in size. If they are longer, the contents of <i>tok</i> is unpredictable.	
SEE ALSO	<b>token</b>	

p

pause

PURPOSE	Pauses for a specified number of seconds.	
FORMAT	<b>pause</b> ( <i>sec</i> );	
INPUT	<i>sec</i>	scalar, seconds to pause.
SOURCE	pause.src	
SEE ALSO	<b>wait</b>	

pdfn

PURPOSE	Computes the standard Normal (scalar) probability density function.	
FORMAT	$y = \text{pdfn}(x);$	
INPUT	$x$	N×K matrix.
OUTPUT	$y$	N×K matrix containing the standard Normal probability density function of $x$ .

## pi

---

**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 = rndn(2,2);`  
`y = pdfn(x);`

```
x = -1.828915    0.514485  
    -0.550219   -0.275229
```

```
y = 0.074915    0.349488  
    0.342903    0.384115
```

## pi

**PURPOSE** Returns the mathematical constant  $\pi$ .

**FORMAT** `y = pi;`

**OUTPUT** *y* scalar, the value of  $\pi$ .

**EXAMPLE** `format /rdn 16,14;`  
`print pi;`

```
3.14159265358979
```

pinv

p

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 = pinv(x);`

INPUT       `x`            N×M matrix.

GLOBAL     `_svdtol`       scalar, any singular values less than `_svdtol` are treated as  
INPUT       zero in determining the rank of the input matrix. The default value for `_svdtol` is 1.0e-13.

OUTPUT     `y`            M×N matrix that satisfies the 4 Moore-Penrose conditions:  
                  $xyx = x$   
                  $xyx = y$   
                  $xy$  is symmetric  
                  $yx$  is symmetric

GLOBAL     `_svderr`       scalar, if not all of the singular values can be computed  
OUTPUT     `_svderr` will be nonzero.

EXAMPLE     `x = { 6 5 4, 2 7 5 };`  
             `y = pinv(x);`

                 0.22017139   -0.16348055  
`y =`   -0.052076467   0.13447594  
         -0.015161503   0.077125906

SOURCE     `svd.src`

## pinvmt

### 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** *x* N×M matrix.  
*tol* scalar, any singular values less than *tol* are treated as zero in determining the rank of the input matrix.

**OUTPUT** *y* M×N matrix that satisfies the 4 Moore-Penrose conditions:  
 $xyx = x$   
 $xyx = y$   
*xy* is symmetric  
*yx* is symmetric  
*err* scalar, if not all of the singular values can be computed *err* will be nonzero.

**EXAMPLE** `x = { 6 5 4, 2 7 5 };`  
`tol = 1e-13;`  
`{ y, err } = pinvmt(x, tol);`

$$y = \begin{pmatrix} 0.22017139 & -0.16348055 \\ -0.052076467 & 0.13447594 \\ -0.015161503 & 0.077125906 \end{pmatrix}$$

`err = 0`

**SOURCE** `svdmt.src`

PURPOSE	Graph data using polar coordinates.		
LIBRARY	pgraph		
FORMAT	<b>polar</b> ( <i>radius</i> , <i>theta</i> );		
INPUT	<i>radius</i>	N×1 or N×M matrix. Each column contains the magnitude for a particular line.	
	<i>theta</i>	N×1 or N×M matrix. Each column represents the angle values for a particular line.	
SOURCE	polar.src		
SEE ALSO	<b>xy, logx, logy, loglog, scale, xtics, ytics</b>		

PURPOSE	Computes the characteristic polynomial of a square matrix.		
FORMAT	$c = \mathbf{polychar}(x);$		
INPUT	$x$	N×N matrix.	
OUTPUT	$c$	(N+1)×1 vector of coefficients of the N <sup>th</sup> 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 ( $c[1]=1$ ).		

## polyeval

---

SOURCE    `poly.src`

SEE ALSO    **polymake**, **polymult**, **polyroot**, **polyeval**

### polyeval

PURPOSE    Evaluates polynomials. Can either be one or more scalar polynomials or a single matrix polynomial.

FORMAT    `y = polyeval(x,c);`

INPUT    `x`         $1 \times K$  or  $N \times N$ ; that is,  $x$  can either represent  $K$  separate scalar values at which to evaluate the (scalar) polynomial(s), or it can represent a single  $N \times N$  matrix.

`c`         $(P+1) \times K$  or  $(P+1) \times 1$  matrix of coefficients of polynomials to evaluate. If  $x$  is  $1 \times K$ , then  $c$  must be  $(P+1) \times K$ . If  $x$  is  $N \times N$ ,  $c$  must be  $(P+1) \times 1$ . That is, if  $x$  is a matrix, it can only be evaluated at a single set of coefficients.

OUTPUT    `y`         $K \times 1$  vector (if  $c$  is  $(P+1) \times K$ ) or  $N \times N$  matrix (if  $c$  is  $(P+1) \times 1$  and  $x$  is  $N \times N$ ):

$$y = (c[1,.] * x^p + c[2,.] * x^{(p-1)} + \dots + c[p+1,.]');$$

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    `x = 2;`  
             `let c = 1 1 0 1 1;`  
             `y = polyeval(x,c);`

The result is 27. Note that this is the decimal value of the binary number 11011.



```
y = polyeval(x,1|zeros(n,1));
```

This will raise the matrix  $x$  to the  $n^{th}$  power (e.g:  $x*x*x*x*\dots*x$ ).

SOURCE poly.src

SEE ALSO polymake, polychar, polymult, polyroot

p

polyint

PURPOSE Calculates an  $N^{th}$  order polynomial interpolation.

FORMAT `y = polyint(xa,ya,x);`

INPUT    *xa*            N×1 vector,  $X$  values.  
          *ya*            N×1 vector,  $Y$  values.  
          *x*            scalar,  $X$  value to solve for.

GLOBAL    **\_poldeg**            scalar, the degree of polynomial required, default 6.  
INPUT

OUTPUT    *y*            result of interpolation or extrapolation.

GLOBAL    **\_polerr**            scalar, interpolation error.  
OUTPUT

REMARKS    Calculates an  $N^{th}$  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 determine an up to  $N^{th}$  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.

## polymake

---

SOURCE    `polyint.src`

TECHNICAL    Press, W.P., B.P. Flannery, S.A. Teukolsky, and W.T. Vetterling. *Numerical*  
NOTES        *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`            N×1 vector containing roots of the desired polynomial.

OUTPUT    `c`            (N+1)×1 vector containing the coefficients of the N<sup>th</sup> order  
polynomial with roots `r`:

$$p(z) = c[1] * z^n + c[2] * z^{(n-1)} + \dots + c[n] * z + c[n+1];$$

REMARKS    The coefficient of  $z^n$  is set to unity (`c[1]=1`).

EXAMPLE    `r = { 2, 1, 3 };`  
              `c = polymake(r);`

```
          1.0000000
      c = -6.0000000
          11.0000000
          -6.0000000
```

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  $y = \text{polymat}(x,p);$

INPUT  $x$   $N \times K$  matrix.  
 $p$  scalar, positive integer.

OUTPUT  $y$   $N \times (p \times K)$  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  $r = \text{polymroot}(c);$

INPUT  $c$   $(N+1) \times K \times K$  matrix of coefficients of an  $N^{th}$  order polynomial of rank  $K$ .

## polymult

---

OUTPUT      $r$              K\*N vector containing the roots of the determinantal equation.

REMARKS      $c$  is constructed of N+1 K×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  $\det(A2 * t^2 + A1 * t + A0) = 0$  where:

$$A2 = \begin{pmatrix} 1 & 2 \\ 2 & 1 \end{pmatrix}$$

$$A1 = \begin{pmatrix} 5 & 8 \\ 10 & 7 \end{pmatrix}$$

$$A0 = \begin{pmatrix} 3 & 4 \\ 6 & 5 \end{pmatrix}$$

```
a2 = { 1 2, 2 1 };  
a1 = { 5 8, 10 7 };  
a0 = { 3 4, 6 5 };
```

```
print polymroot(a2|a1|a0);
```

```
-4.3027756  
-.69722436  
-2.6180340  
-.38196601
```

polymult

p

PURPOSE Multiplies polynomials.

FORMAT  $c = \text{polymult}(c1, c2);$

INPUT  $c1$  (D1+1)×1 vector containing the coefficients of the first polynomial.  
 $c2$  (D2+1)×1 vector containing the coefficients of the second polynomial.

OUTPUT  $c$  (D1+D2)×1 vector containing the coefficients of the product of the two polynomials.

EXAMPLE  $c1 = \{ 2, 1 \};$   
 $c2 = \{ 2, 0, 1 \};$   
 $c = \text{polymult}(c1, c2);$

$c =$   
4.0000000  
2.0000000  
2.0000000  
1.0000000

SOURCE poly.src

SEE ALSO **polymake, polychar, polyroot, polyeval**

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  $5 * x^3 + 6 * x + 3$  would be:  $c1=5| 0| 6| 3$ . (Note that zeros must be explicitly given if there are powers of  $x$  missing.)

## pop

---

### polyroot

PURPOSE	Computes the roots of a polynomial given the coefficients.		
FORMAT	$y = \text{polyroot}(c);$		
INPUT	$c$	(N+1)×1 vector of coefficients of an N <sup>th</sup> order polynomial: $p(z) = c[1] * z^n + c[2] * z^{(n-1)} + \dots + c[n] * z + c[n + 1]$	
OUTPUT	$y$	N×1 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		
SEE ALSO	<b>polymake</b> , <b>polychar</b> , <b>polymult</b> , <b>polyeval</b>		

### pop

PURPOSE	Provides access to a last-in, first-out stack for matrices.		
FORMAT	<b>pop</b> $b$ ; <b>pop</b> $a$ ;		
REMARKS	This is used with <b>gosub</b> , <b>goto</b> , and <b>return</b> 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**'ed 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;
```

$$c = z \qquad b = y \qquad a = x$$

Note that there must be a separate **pop** statement for each matrix popped.

SEE ALSO **gosub, goto, return**

pqgwin

PURPOSE Sets the graphics viewer mode.

## previousindex

---

LIBRARY	pgraph
FORMAT	<b>pqgwin one;</b>  <b>pqgwin many;</b>
REMARKS	If you call <b>pqgwin one</b> , only a single viewer will be used. If you call <b>pqgwin many</b> , a new viewer will be used for each graph.  <b>pqgwin manual</b> and <b>pqgwin auto</b> are supported for backwards compatibility, <b>manual=one</b> , <b>auto=many</b> .
EXAMPLE	pqgwin many;
SOURCE	pgraph.src
SEE ALSO	<b>setvwrmode</b>

## previousindex

PURPOSE	Returns the index of the previous element or subarray in an array.		
FORMAT	$pi = \text{previousindex}(i,o);$		
INPUT	$i$	M×1 vector of indices into an array, where M≤N.	
	$o$	N×1 vector of orders of an N-dimensional array.	
OUTPUT	$pi$	M×1 vector of indices, the index of the previous element or subarray in the array corresponding to $o$ .	
REMARKS	<b>previousindex</b> 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);`

                 2  
`ind = 2`  
                 5

p

In this example, **previousindex** decremented **ind** to index the previous 6×7 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    *x*            N×K data matrix, N>K, full rank.  
          *j*            scalar, number of principal components to be computed (*j*≤K).

OUTPUT   *p*            N×J matrix of the first *j* principal components of *x* in descending order of amount of variance explained.  
          *v*            J×1 vector of fractions of variance explained.  
          *a*            J×K matrix of factor loadings, such that  $x = p*a$  +error.

REMARKS    Adapted from a program written by Mico Loretan.

## print

---

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** **[/flush]** **[/typ]** **[/fmted]** **[/mf]** **[/jnt]** *list\_of\_expressions***[:,];**

**INPUT** */typ* literal, symbol type flag.

**/mat, /sa, /str** Indicate which symbol types you are setting the output format for: matrices and arrays (**/mat**), string arrays (**/sa**), and/or strings (**/str**). You can specify more than one */typ* flag; the format will be set for all types indicated. If no */typ* flag is listed, **print** assumes **/mat**.

*/fmted* literal, enable formatting flag.

**/on, /off** Enable/disable formatting. 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 1×1 string array.

*/mf* literal, matrix format. It controls the way rows of a matrix are separated from one another. The possibilities are:

**/m0** no delimiters before or after rows when printing out matrices.

<b>/m1</b> or <b>/mb1</b>	print 1 carriage return/line feed pair before each row of a matrix with more than 1 row.
<b>/m2</b> or <b>/mb2</b>	print 2 carriage return/line feed pairs before each row of a matrix with more than 1 row.
<b>/m3</b> or <b>/mb3</b>	print “Row 1”, “Row 2”...before each row of a matrix with more than one row.
<b>/ma1</b>	print 1 carriage return/line feed pair after each row of a matrix with more than 1 row.
<b>/ma2</b>	print 2 carriage return/line feed pairs after each row of a matrix with more than 1 row.
<b>/a1</b>	print 1 carriage return/line feed pair after each row of a matrix.
<b>/a2</b>	print 2 carriage return/line feed pairs after each row of a matrix.
<b>/b1</b>	print 1 carriage return/line feed pair before each row of a matrix.
<b>/b2</b>	print 2 carriage return/line feed pairs before each row of a matrix.
<b>/b3</b>	print “Row 1”, “Row 2”... before each row of a matrix.
<b>/jnt</b>	literal, controls justification, notation, and the trailing character.

### **Right-Justified**

<b>/rd</b>	Signed decimal number in the form <code>[[ - ]]#####.####</code> , where <code>####</code> 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.
<b>/re</b>	Signed number in the form <code>[[ - ]]#.##E±###</code> , where <code>#</code> is one decimal digit, <code>##</code> is one or more decimal digits depending on the precision, and <code>###</code> is three decimal digits. If precision is 0, the form will be <code>[[ - ]]#E±###</code> with no decimal point printed.

<b>/ro</b>	This will give a format like <b>/rd</b> or <b>/re</b> depending on which is most compact for the number being printed. A format like <b>/re</b> will be used only if the exponent value is less than -4 or greater than the precision. If a <b>/re</b> format is used, a decimal point will always appear. The precision signifies the number of significant digits displayed.
<b>/rz</b>	This will give a format like <b>/rd</b> or <b>/re</b> depending on which is most compact for the number being printed. A format like <b>/re</b> will be used only if the exponent value is less than -4 or greater than the precision. If a <b>/re</b> 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

<b>/ld</b>	Signed decimal number in the form <code>[[ - ]#####.###</code> , where <code>####</code> 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.
<b>/le</b>	Signed number in the form <code>[[ - ]#.##E±###</code> , where <code>#</code> is one decimal digit, <code>##</code> is one or more decimal digits depending on the precision, and <code>###</code> is three decimal digits. If precision is 0, the form will be <code>[[ - ]#E±###</code> with no decimal point printed. If the number is positive, a space character will replace the leading minus sign.
<b>/lo</b>	This will give a format like <b>/ld</b> or <b>/le</b> depending on which is most compact for the number being printed. A format like <b>/le</b> will be used only if the exponent value is less than -4 or greater than the

	precision. If a <b>/le</b> 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.
<b>/lz</b>	This will give a format like <b>/ld</b> or <b>/le</b> depending on which is most compact for the number being printed. A format like <b>/le</b> will be used only if the exponent value is less than -4 or greater than the precision. If a <b>/le</b> 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:

	<code>format /rdn 1,3;</code>
<b>s</b>	The number will be followed immediately by a space character. This is the default.
<b>c</b>	The number will be followed immediately by a comma.
<b>t</b>	The number will be followed immediately by a tab character.
<b>n</b>	No trailing character.

The default when **GAUSS** is first started is:

	<code>format /m1 /ro 16,8;</code>
<b>::</b>	Double semicolons following a <b>print</b> statement will suppress the final carriage return/line feed.

*list\_of\_expressions* any **GAUSS** expressions that produce matrices, arrays, strings, or string arrays and/or names of variables to print, separated by spaces.

**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>\b</b>	backspace (ASCII 8)
<b>\e</b>	escape (ASCII 27)
<b>\f</b>	form feed (ASCII 12)
<b>\g</b>	beep (ASCII 7)
<b>\l</b>	line feed (ASCII 10)
<b>\r</b>	carriage return (ASCII 13)
<b>\t</b>	tab (ASCII 9)
<b>\###</b>	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;
```

EXAMPLE

```
x = rndn(3,3);
format /rd 16,8;
print x;
```

```
format /re 12,2;
print x;
print /rd/m3 x;
```

0.14357994	-1.39272762	-0.91942414
0.51061645	-0.02332207	-0.02511298
-1.54675893	-1.04988540	0.07992059

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

Row 1

0.14	-1.39	-0.92
------	-------	-------

Row 2

## printdos

---

```
          0.51      -0.02      -0.03
Row 3
      -1.55      -1.05      0.08
```

In this example, a 3×3 random matrix is printed using 3 different formats. Notice that in the last statement, the format is overridden in the **print** statement itself but the field and precision remain the same.

```
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** `printdos "\27[7m"; /* set for reverse video */`



---

```
printdos "\27[0m"; /* set for normal text */
```

SEE ALSO **print**, **printfm**, **screen**

## printfm

p

**PURPOSE** Prints a matrix using a different format for each column of the matrix.

**FORMAT**  $y = \text{printfm}(x, \text{mask}, \text{fmt});$

**INPUT**

$x$	$N \times K$ matrix which is to be printed and which may contain both character and numeric data.
$\text{mask}$	$L \times M$ matrix, $E \times E$ conformable with $x$ , containing ones and zeros, which is used to specify whether the particular row, column, or element is to be printed as a character (0) or numeric (1) value.
$\text{fmt}$	$K \times 3$ or $1 \times 3$ matrix where each row specifies the format for the respective column of $x$ .

**OUTPUT**  $y$  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  $\text{mask}$  is 0, then that element of  $x$  is printed as a character string of up to 8 characters. If  $\text{mask}$  contains a 1, then that element of  $x$  is assumed to be a double precision floating point number.

The contents of  $\text{fmt}$  are as follows:

<b>[K,1]</b>	format string,	a string 8 characters maximum.
<b>[K,2]</b>	field width,	a number < 80.
<b>[K,3]</b>	precision,	a number < 17.

The format strings correspond to the **format** slash commands as follows:

```
/rdn    ‘‘*.1f’’  
/ren    ‘‘*.1E’’  
/ron    ‘‘#*.1G’’  
/rzn    ‘‘*.1G’’  
  
/ldn    ‘‘- *.1f’’  
/len    ‘‘- *.1E’’  
/lon    ‘‘-# *.1G’’  
/lzn    ‘‘- *.1G’’
```

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

‘‘*.1f,’’	right-justified decimal followed by a comma.
‘‘-*.1s ’’	left-justified string followed by a space.
‘‘*.1f’’	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*.1f’’	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
"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);

```

p

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`  $N \times K$  matrix which is to be printed.

`mask` scalar, 1 if `x` is numeric or 0 if `x` is character.

- or -

$1 \times K$  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** `x = rndn(5,4);`  
`call printfmt(x,1);`

**SOURCE** `gauss.src`

**GLOBALS** `__fmtcv, __fmtnv`

**SEE ALSO** `formatcv, formatnv`

**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*);

**INPUT** *nrets* constant, number of objects returned by the procedure. If *nrets* is not explicitly given, the default is 1. Legal values are 0 to 1023. The **retp** statement is used to return values from a procedure.

*name* literal, name of the procedure. This name will be a global symbol.

*arglist* 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 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 */
```

## prodc

---

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 12.

SEE ALSO **keyword, call, endp, local, retp**

---

**prodc**

PURPOSE	Computes the products of all elements in each column of a matrix.	
FORMAT	$y = \text{prodc}(x);$	
INPUT	$x$	$N \times K$ matrix.
OUTPUT	$y$	$K \times 1$ matrix containing the products of all elements in each column of $x$ .
REMARKS	<p>To find the products of the elements in each row of a matrix, transpose before applying <b>prodc</b>. If <math>x</math> is complex, use the bookkeeping transpose (<math>\cdot'</math>).</p> <p>To find the products of all of the elements in a matrix, use the <b>vecr</b> function before applying <b>prodc</b>.</p>	
EXAMPLE	<pre>let x[3,3] = 1 2 3               4 5 6               7 8 9;  y = prodc(x);        28 y =   80      162</pre>	
SEE ALSO	<b>sumc</b> , <b>meanc</b> , <b>stdc</b>	

p

## putarray

PURPOSE	Puts a contiguous subarray into an N-dimensional array and returns the resulting array.	
FORMAT	$y = \text{putarray}(a, loc, src);$	

## putf

---

INPUT	<i>a</i>	N-dimensional array.
	<i>loc</i>	M×1 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.
OUTPUT	<i>y</i>	N-dimensional array.
REMARKS	<p>If <i>loc</i> is an N×1 vector, then <i>src</i> must be a scalar. If <i>loc</i> is an [N-1]×1 vector, then <i>src</i> must be a 1-dimensional array or a 1×L vector, where L is the size of the fastest moving dimension of the array. If <i>loc</i> is an [N-2]×1 vector, then <i>src</i> must be a K×L matrix, or a K×L 2-dimensional array, where K is the size of the second fastest moving dimension.</p> <p>Otherwise, if <i>loc</i> is an M×1 vector, then <i>src</i> must be an [N-M]-dimensional array, whose dimensions are the same size as the corresponding dimensions of array <i>a</i>.</p>	
EXAMPLE	<pre>a = arrayalloc(2 3 4 5 6,0); src = arrayinit(4 5 6,5); loc = { 2,1 }; a = putarray(a,loc,src);</pre> <p>This example sets the contiguous 4×5×6 subarray of <b>a</b> beginning at [2,1,1,1,1] to the array <b>src</b>, in which each element is set to the specified value 5.</p>	
SEE ALSO	<b>setarray</b>	

## putf

PURPOSE	Writes the contents of a string to a file.
FORMAT	<i>ret</i> = <b>putf</b> ( <i>filename, str, start, len, mode, append</i> );



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.
	<b>0</b>	normal return
	<b>1</b>	null file name
	<b>2</b>	file open error
	<b>3</b>	file write error
	<b>4</b>	output string too long
	<b>5</b>	null output string, or illegal <i>mode</i> value
	<b>6</b>	illegal <i>append</i> value
	<b>16</b>	(1) append specified but file did not exist; file was created (warning only)
REMARKS	If <i>mode</i> is set to (1) binary, a string of length <i>len</i> will be written to <i>filename</i> . If <i>mode</i> is set to (0) ASCII, the string will be output up to length <i>len</i> or until <b>putf</b> encounters a ^Z (ASCII 26) in <i>str</i> . The ^Z will not be written to <i>filename</i> .	
	If <i>append</i> is set to (0) overwrite, the current contents of <i>filename</i> will be destroyed. If <i>append</i> is set to (1) append, <i>filename</i> will be created if it does not already exist.	
	If an error occurs, <b>putf</b> will either return an error code or terminate the program with an error message, depending on the <b>trap</b> state. If bit 2 (the 4's bit) of the trap flag is 0, <b>putf</b> will terminate with an error message. If bit 2 of the trap flag is 1, <b>putf</b> will return an error code. The value of the trap flag can be tested with <b>trapchk</b> .	
SOURCE	putf.src	
SEE ALSO	<b>getf</b>	

## putvals

### putvals

**PURPOSE** Inserts values into a matrix or N-dimensional array.

**FORMAT** `y = putvals(x,inds,vals);`

**INPUT** *x*  $M \times K$  matrix or N-dimensional array.

*inds*  $L \times D$  matrix of indices, specifying where the new values are to be inserted, where  $D$  is the number of dimensions in *x*.

*vals*  $L \times 1$  vector, new values to insert.

**OUTPUT** *y*  $M \times K$  matrix or N-dimensional array, copy of *x* containing the new values in *vals*.

**REMARKS** If *x* is a vector, *inds* should be an  $L \times 1$  vector. If *x* is a matrix, *inds* should be an  $L \times 2$  matrix. Otherwise if *x* is an N-dimensional array, *inds* should be an  $L \times N$  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);
```

---

	1.0000	0.3616	0.6032	-0.3974
	0.7644	-1.8509	-0.2703	2.0000
y =	0.7886	3.0000	-1.4998	4.0000
	0.6639	-0.7972	1.2713	0.1896
	0.6303	0.7879	5.0000	-0.5419

p

## pvCreate

**PURPOSE** Returns an initialized an instance of structure of type **PV**.

**FORMAT** *p1* = **pvCreate**;

**OUTPUT** *p1* an instance of structure of type **PV**

**EXAMPLE** struct PV p1;  
p1 = pvCreate;

**SOURCE** pv.src

## pvGetIndex

**PURPOSE** Gets row indices of a matrix in a parameter vector.

**FORMAT** *id* = **pvGetIndex**(*p1*,*nm1*);

**INPUT** *p1* an instance of structure of type **PV**.  
*nm1* name or row number of matrix.

## pvGetParNames

---

OUTPUT	<i>id</i>	K×1 vector, row indices of matrix described by <i>nmI</i> in parameter vector.
SOURCE	<code>pv.src</code>	

## pvGetParNames

PURPOSE	Generates names for parameter vector stored in structure of type <b>PV</b> .	
INCLUDE	<code>pv.sdf</code>	
FORMAT	<code>s = pvGetParNames(pI);</code>	
INPUT	<i>pI</i>	an instance of structure of type <b>PV</b> .
OUTPUT	<i>s</i>	K×1 string array, names of parameters.
REMARKS	If the vector in the structure of type <b>PV</b> 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	<pre>#include pv.sdf struct PV p1;  p1 = pvCreate;  x = { 1 2,       3 4 };  mask = { 1 0,          0 1 };</pre>	

```
p1 = pvPackm(p1,x,"P",mask);
```

```
print pvGetParNames(p1);
```

```
P[1,1]
```

```
P[2,2]
```

SOURCE pv.src

p

## pvGetParVector

PURPOSE Retrieves parameter vector from structure of type **PV**.

INCLUDE pv.sdf

FORMAT  $p = \text{pvGetParVector}(p1);$

INPUT  $p1$  an instance of structure of type **PV**.

OUTPUT  $p$   $K \times 1$  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 

```
#include pv.sdf
struct PV p1;
```

```
p1 = pvCreate;
```

```
x = { 1 2,
      3 4 };
```

```
mask = { 1 0,
```

## pvLength

---

```
      0 1 };  
  
p1 = pvPackm(p1,x,"X",mask);  
  
print pvUnpack(p1,1);
```

```
1.000  2.000  
3.000  4.000
```

```
print pvGetParVector(p1);
```

```
1.000  
4.000
```

SOURCE    pv.src

## pvLength

PURPOSE    Returns the length of a parameter vector.

FORMAT     $n = \mathbf{pvLength}(p1);$

INPUT     $p1$             an instance of structure of type **PV**.

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**(*pl*);

INPUT       *pl*             an instance of structure of type **PV**.

OUTPUT      *n*             K×1 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     *pl* = **pvPack**(*pl*,*x*,*nm*);

INPUT       *pl*             an instance of structure of type **PV**.  
               *x*             M×N matrix or N-dimensional array.  
               *nm*           string, name of matrix/array.

OUTPUT      *pl*             an instance of structure of type **PV**.

EXAMPLE     **#include pv.sdf**  
               **y = rndn(100,1);**  
               **x = rndn(100,5);**

## pvPacki

---

```
struct PV p1;  
p1 = pvCreate;  
p1 = pvPack(p1,x,"Y");  
p1 = pvPack(p1,y,"X");
```

These matrices can be extracted using the **pvUnpack** command:

```
y = pvUnpack(p1,"Y");  
x = pvUnpack(p1,"X");
```

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 *pl* = **pvPacki**(*pl*,*x*,*nm*,*i*);

INPUT	<i>pl</i>	an instance of structure of type <b>PV</b> .
	<i>x</i>	M×N 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 *pl* an instance of structure of type **PV**.

EXAMPLE **#include pv.sdf**



```

y = rndn(100,1);
x = rndn(100,5);
struct PV p1;
p1 = pvCreate;
p1 = pvPacki(p1,y,"Y",1);
p1 = pvPacki(p1,x,"X",2);

```

These matrices can be extracted using the **pvUnpack** command:

```

y = pvUnpack(p1,1);
x = pvUnpack(p1,2);

```

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**

<i>p1</i>	an instance of structure of type <b>PV</b> .
<i>x</i>	M×N matrix or N-dimensional array.
<i>nm</i>	string, name of matrix/array or N-dimensional array.
<i>mask</i>	M×N 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);

      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

**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 <b>PV</b> .
<i>x</i>	M×N matrix or N-dimensional array.
<i>nm</i>	string, matrix or array name.
<i>mask</i>	M×N 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** *p1* 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,
```

## pvPacks

---

```
      3 4 };

mask = { 1 0,
        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);`

<b>INPUT</b>	<i>p1</i>	an instance of structure of type <b>PV</b> .
	<i>x</i>	M×M symmetric matrix.
	<i>nm</i>	string, matrix name.

OUTPUT *p1* 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");

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**

<code>p1</code>	an instance of structure of type <b>PV</b> .
<code>x</code>	M×M symmetric matrix.
<code>nm</code>	string, matrix name.
<code>i</code>	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
```

SEE ALSO **pvPacks**, **pvUnpack**

## pvPacksm

p

**PURPOSE** Packs symmetric matrix into a structure of type **PV** with a mask.

**INCLUDE** `pv.sdf`

**FORMAT** `pl = pvPacksm(pl,x,nm,mask);`

**INPUT**

<i>pl</i>	an instance of structure of type <b>PV</b> .
<i>x</i>	M×M symmetric matrix.
<i>nm</i>	string, matrix name.
<i>mask</i>	M×M matrix, mask matrix of zeros and ones.

**OUTPUT** *pl* 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

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**

**p**

**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**    *p1*            an instance of structure of type **PV**.  
              *x*             M×M symmetric matrix.  
              *nm*            string, matrix name.  
              *mask*          M×M matrix, symmetric mask matrix of zeros and ones.  
              *i*             scalar, index of matrix in lookup table.

**OUTPUT**    *p1*            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

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);
```

```

10.000  2.000  12.000
 2.000  11.000  5.000
12.000  5.000  13.000

```

SEE ALSO **pvPacksm**, **pvUnpack**

p

## pvPutParVector

**PURPOSE** Inserts parameter vector into structure of type **PV**.

**INCLUDE** `pv.sdf`

**FORMAT** `p1 = pvPutParVector(p1,p);`

**INPUT** *p1* an instance of structure of type **PV**.  
*p* K×1 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 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,

```

## pvTest

---

```
      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
```

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**.

FORMAT `x = pvUnpack(pl,m);`

INPUT *pl* an instance of structure of type **PV**.  
*m* string, name of matrix, or integer, index of matrix.

OUTPUT *x* M×N general matrix or M×M symmetric matrix or N-dimensional array.

SOURCE `pv.src`

## QNewton

PURPOSE Optimizes a function using the BFGS descent algorithm.

FORMAT `{ x,f,g,ret } = QNewton(&fct,start);`

INPUT *&fct* 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.  
*start* K×1 vector, start values.

GLOBAL INPUT `_qn_RelGradTol` scalar, convergence tolerance for relative gradient of estimated coefficients. Default = 1e-5.

**\_qn\_GradProc** 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  $K \times 1$  vector of parameter values, and a single output argument, a  $K \times 1$  vector of gradients of the function with respect to the parameters evaluated at the vector of parameter values. If **\_qn\_GradProc** is 0, **QNewton** uses **gradp**.

**\_qn\_MaxIters** scalar, maximum number of iterations. Default = 1e+5. Termination can be forced by pressing C on the keyboard.

**\_qn\_PrintIters** scalar, if 1, print iteration information. Default = 0. Can be toggled during iterations by pressing P on the keyboard.

**\_qn\_ParNames**  $K \times 1$  vector, labels for parameters.

**\_qn\_PrintResults** scalar, if 1, results are printed.

OUTPUT    *x*             $K \times 1$  vector, coefficients at the minimum of the function.

*f*            scalar, value of function at minimum.

*g*             $K \times 1$  vector, gradient at the minimum of the function.

*ret*          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
*/

z = loadadd("tobit"); /* get data */
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(invdpd(h)));

output off;

/*
**  log-likelihood proc
*/

proc lpr(b);
    local s,m,u;
    s = b[4];
    if s <= 1e-4;
        retp(error(0));
    endif;
    m = z[.,2:4]*b[1:3,.];
    u = z[.,1] ./= 0;
    retp(-sumc(u.*lnpdfn2(z[.,1]-m,s) +
        (1-u).*(ln(cdfnc(m/sqrt(s))))));
endproc;

```

q

## QNewtonmt

---

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.

INCLUDE qnewtonmt.sdf

FORMAT *out* = **QNewtonmt**(&*fct*,*par*,*data*,*c*);

INPUT &*fct*      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 a **DS** structure containing data, if any. And, one output argument, the value of the function evaluated at the input vector of parameter values.

*par*      an instance of a **PV** structure. The *par* instance is passed to the user-provided procedure pointed to by &*fct*. *par* is constructed using the **pvPack** functions.



*data* an array of instances of a **DS** structure. This array is passed to the user-provided pointed by *&fct* to be used in the objective function. **QNewtonmt** does not look at this structure. Each instance contains the the following members which can be set in whatever way that is convenient for computing the objective function:

*data[i].dataMatrix* N×K matrix, data matrix.  
*data[i].dataArray* N×K×L... array, data array.  
*data[i].vnames* string array, variable names (optional).  
*data[i].dsname* string, data name (optional).  
*data[i].type* scalar, type of data (optional).

*c* 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 *c*, the members are:

*c.CovType* scalar, if 1, ML covariance matrix, else if 2, QML covariance matrix is computed. Default is 0, no covariance matrix.  
*c.GradProc* 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.  
*c.MaxIters* scalar, maximum number of iterations. Default = 1e+5.  
*c.MaxTries* scalar, maximum number of attempts in random search. Default = 100.  
*c.relGradTol* scalar, convergence tolerance for gradient of estimated coefficients. Default = 1e-5. When this criterion has been satisfied **QNewtonmt** exits the iterations.  
*c.randRadius* scalar, If zero, no random search is attempted. If nonzero, it is the radius of the random search. Default = .001.  
*c.output* 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 <b>QNewtonmtOut</b> structure. For an instance named <i>out</i> , the members are:	
		<i>out.par</i>	instance of a <b>PV</b> 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: <ul style="list-style-type: none"> <li><b>0</b> normal convergence.</li> <li><b>1</b> forced exit.</li> <li><b>2</b> maximum number of iterations exceeded.</li> <li><b>3</b> function calculation failed.</li> <li><b>4</b> gradient calculation failed.</li> <li><b>5</b> Hessian calculation failed.</li> <li><b>6</b> line search failed.</li> <li><b>7</b> error with constraints.</li> <li><b>8</b> function complex.</li> </ul>
		<i>out.moment</i>	K×K matrix, covariance matrix of parameters, if <i>c.covType</i> > 0.
		<i>out.hessian</i>	K×K matrix, matrix of second derivatives of objective function with respect to parameters.
REMARKS	<p>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.</p> <p>These functions have one input argument that is an instance of type struct <b>PV</b> and a second argument that is an instance of type struct <b>DS</b>. On input to the call to <b>QNewtonmt</b>, the first argument contains starting values for the parameters and the second argument any required data. The data are passed in a separate argument because the structure in the first argument will be copied as it is passed through procedure calls which would be very costly if it contained large</p>		

data matrices. Since **QNewtonmt** makes no changes to the second argument it will be passed by pointer thus saving time because its contents aren't copied.

The **PV** structures 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 Micherlitz(struct PV par1, struct DS data1);
    local p0,e,s2,x,y;
    p0 = pvUnpack(par1,"parameters");
    y = data1.dataMatrix[.,1];
    x = data1.dataMatrix[.,2];
    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 first and second columns of a data matrix stored in a single **DS** structure. Alternatively these two columns of data can be entered into a vector of **DS** structures one for each column of data:

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** The following is a complete example for estimating the parameters of the Micherlitz equation in data on the parameters and where an optional gradient procedure has been provided.

q

```
#include QNewtonmt.sdf

struct DS d0;
d0 = dsCreate;

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);
d0.dataMatrix = y~x;

struct QNewtonmtControl c0;
c0 = QNewtonmtControlCreate;
c0.output = 1; /* print results */
c0.covType = 1; /* compute moment matrix */
               /* of parameters */

struct PV par1;
par1 = pvCreate;
par1 = pvPack(par1,1|1|0,"parameters");

struct QNewtonmt out1;
out1 = QNewtonmt(&Micherlitz,par1,d0,c0);
```

SOURCE qnewtonmt.src

SEE ALSO [QNewtonmtControlCreate](#), [QNewtonmtOutCreate](#)

## QNewtonmtControlCreate

**PURPOSE** Creates default **QNewtonmtControl** structure.

**INCLUDE** `qnewtonmt.sdf`

**FORMAT** `c = QNewtonmtControlCreate;`

**OUTPUT** `c` instance of **QNewtonmtControl** structure with members set to default values.

**SOURCE** `qnewtonmt.src`

**SEE ALSO** [QNewtonmt](#)

q

## QNewtonmtOutCreate

**PURPOSE** Creates default **QNewtonmtOut** structure.

**FORMAT** `c = QNewtonmtOutCreate;`

**OUTPUT** `c` instance of **QNewtonmtOut** structure with members set to default values.

**SOURCE** `qnewtonmt.src`

**SEE ALSO** [QNewtonmt](#)

## QNewtonSet

**PURPOSE**     Resets global variables used by **QNewton** to default values.

**FORMAT**     **QNewtonSet;**

**SOURCE**     `qnewton.src`

## QProg

**PURPOSE**     Solves the quadratic programming problem.

**FORMAT**      $\{ x, u1, u2, u3, u4, u5 \} = \text{QProg}(start, q, r, a, b, c, d, bnds);$

<b>INPUT</b>	<i>start</i>	K×1 vector, start values.
	<i>q</i>	K×K matrix, symmetric model matrix.
	<i>r</i>	K×1 vector, model constant vector.
	<i>a</i>	M×K matrix, equality constraint coefficient matrix, or scalar 0, no equality constraints.
	<i>b</i>	M×1 vector, equality constraint constant vector, or scalar 0, will be expanded to M×1 vector of zeros.
	<i>c</i>	N×K matrix, inequality constraint coefficient matrix, or scalar 0, no inequality constraints.
	<i>d</i>	N×1 vector, inequality constraint constant vector, or scalar 0, will be expanded to N×1 vector of zeros.
	<i>bnds</i>	K×2 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 ±1e200.

GLOBAL INPUT **\_qprog\_maxit** scalar, maximum number of iterations. Default = 1000.

OUTPUT *x*  $K \times 1$  vector, coefficients at the minimum of the function.  
*u1*  $M \times 1$  vector, Lagrangian coefficients of equality constraints.  
*u2*  $N \times 1$  vector, Lagrangian coefficients of inequality constraints.  
*u3*  $K \times 1$  vector, Lagrangian coefficients of lower bounds.  
*u4*  $K \times 1$  vector, Lagrangian coefficients of upper bounds.  
*ret* 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 **QProg** solves the standard quadratic programming problem:

$$\min \frac{1}{2} x' Q x - x' R$$

subject to constraints,

$$A x = B$$

$$C x \geq D$$

and bounds,

$$x_{low} \leq x \leq x_{up}$$

SOURCE `qprog.src`

## QProgmt

PURPOSE Solves the quadratic programming problem.

INCLUDE `qprogmt.sdf`

FORMAT `qOut = QProgmt(qIn);`

INPUT	<i>qIn</i>	instance of a <b>qprogmtIn</b> structure containing the following members:
	<i>qIn.start</i>	K×1 vector, start values.
	<i>qIn.q</i>	K×K matrix, symmetric model matrix.
	<i>qIn.r</i>	K×1 vector, model constant vector.
	<i>qIn.a</i>	M×K matrix, equality constraint coefficient matrix, or scalar 0, no equality constraints.
	<i>qIn.b</i>	M×1 vector, equality constraint constant vector, or scalar 0, will be expanded to M×1 vector of zeros.
	<i>qIn.c</i>	N×K matrix, inequality constraint coefficient matrix, or scalar 0, no inequality constraints.
	<i>qIn.d</i>	N×1 vector, inequality constraint constant vector, or scalar 0, will be expanded to N×1 vector of zeros.
	<i>qIn.bounds</i>	K×2 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 ±1e200.
	<i>maxit</i>	scalar, maximum number of iterations. Default = 1000.

OUTPUT	<i>qOut</i>	instance of a <b>qprogmtOut</b> structure containing the following members:
--------	-------------	---



<i>qOut.x</i>	K×1 vector, coefficients at the minimum of the function.										
<i>qOut.lagrange</i>	instance of a <b>qprogMTLagrange</b> structure containing the following members: <table> <tr> <td><i>qOut.lagrange.lineq</i></td><td>M×1 vector, Lagrangian coefficients of equality constraints.</td></tr> <tr> <td><i>qOut.lagrange.linineq</i></td><td>N×1 vector, Lagrangian coefficients of inequality constraints.</td></tr> <tr> <td><i>qOut.lagrange.bounds</i></td><td>K×2 matrix, Lagrangian coefficients of bounds, the first column contains the lower bounds and the second the upper bounds.</td></tr> </table>	<i>qOut.lagrange.lineq</i>	M×1 vector, Lagrangian coefficients of equality constraints.	<i>qOut.lagrange.linineq</i>	N×1 vector, Lagrangian coefficients of inequality constraints.	<i>qOut.lagrange.bounds</i>	K×2 matrix, Lagrangian coefficients of bounds, the first column contains the lower bounds and the second the upper bounds.				
<i>qOut.lagrange.lineq</i>	M×1 vector, Lagrangian coefficients of equality constraints.										
<i>qOut.lagrange.linineq</i>	N×1 vector, Lagrangian coefficients of inequality constraints.										
<i>qOut.lagrange.bounds</i>	K×2 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. <table> <tr> <td><b>0</b></td><td>successful termination</td></tr> <tr> <td><b>1</b></td><td>max iterations exceeded</td></tr> <tr> <td><b>2</b></td><td>machine accuracy is insufficient to maintain decreasing function values</td></tr> <tr> <td><b>3</b></td><td>model matrices not conformable</td></tr> <tr> <td><b>&lt;0</b></td><td>active constraints inconsistent</td></tr> </table>	<b>0</b>	successful termination	<b>1</b>	max iterations exceeded	<b>2</b>	machine accuracy is insufficient to maintain decreasing function values	<b>3</b>	model matrices not conformable	<b>&lt;0</b>	active constraints inconsistent
<b>0</b>	successful termination										
<b>1</b>	max iterations exceeded										
<b>2</b>	machine accuracy is insufficient to maintain decreasing function values										
<b>3</b>	model matrices not conformable										
<b>&lt;0</b>	active constraints inconsistent										

q

REMARKS **QProgmt** solves the standard quadratic programming problem:

$$\min \frac{1}{2}x'Qx - x'R$$

subject to constraints,

$$Ax = B$$

## QProgmtInCreate

---

$$Cx \geq D$$

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`

FORMAT    `s = QProgmtInCreate;`

OUTPUT    `s`            instance of structure of type **QProgmtInCreate**.

SOURCE    `qprogmt.src`

SEE ALSO    **QProgmt**

### qqr

**PURPOSE** Computes the orthogonal-triangular (QR) decomposition of a matrix  $X$ , such that:

$$X = Q_1 R$$

**FORMAT** {  $q1, r$  } = **qqr**( $x$ );

**INPUT**  $x$   $N \times P$  matrix.

**OUTPUT**  $q1$   $N \times K$  unitary matrix,  $K = \min(N, P)$ .  
 $r$   $K \times P$  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**:

```
 $r = \text{cholup}(\text{zeros}(\text{cols}(x), \text{cols}(x)), x);$ 
```

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 **qyr**. 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:

$$Q'X = [ R_1 \ R_2 ]$$

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 { *ql*, *r*, *e* } = **qqre**(*x*);

INPUT *x* N×P matrix.

OUTPUT *ql* N×K unitary matrix, K = min(N,P).

*r* K×P upper triangular matrix.

*e* P×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  $X[:, E]$ .

If you want only the  $R$  matrix, see **qre**. 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  $X_1$  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

$$X_2 = X_1 A$$

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:

$$Q'X = [ R_1 \ R_2 ]$$

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

$$b = \text{qrsol}(Q'Y, RI) | \text{zeros}(N-P, 1);$$

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**, **qtyre**, **olsqr**

q

## qqrep

PURPOSE Computes the orthogonal-triangular (QR) decomposition of a matrix  $X$ , such that:

$$X[., E] = Q_1 R$$

FORMAT `{ ql, r, e } = qqrep(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    *ql*        N×K unitary matrix, K = min(N,P).  
              *r*         K×P upper triangular matrix.  
              *e*         P×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  $X[., E]$ .

**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

q

PURPOSE    Computes the orthogonal-triangular (QR) decomposition of a matrix  $X$ , such that:

$$X = Q_1 R$$

FORMAT    `r = qr(x);`

INPUT       $x$              $N \times P$  matrix.

OUTPUT     $r$              $K \times P$  upper triangular matrix,  $K = \min(N,P)$ .

REMARKS    `qr` is the same as `qqr` but doesn't return the  $Q_1$  matrix. If  $Q_1$  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  $Q_1$  matrix because in most cases it is not required and can be very large. If you need the  $Q_1$  matrix, see the function **qqr**. If you need the entire  $Q$  matrix, call **qyr** 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 **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**, **qtyre**

**qre**

PURPOSE     Computes the orthogonal-triangular (QR) decomposition of a matrix  $X$ , such that:

$$X[:, E] = Q_1 R$$

FORMAT      $\{ r, e \} = \text{qre}(x);$

INPUT        $x$               $N \times P$  matrix.

OUTPUT       $r$               $K \times P$  upper triangular matrix,  $K = \min(N, P)$ .  
               $e$               $P \times 1$  permutation vector.

REMARKS     **qre** is the same as **qqre** but doesn't return the  $Q_1$  matrix. If  $Q_1$  is not wanted, **qre** 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  $Q_1$  has  $P$  columns, then

$$X[:, E] = Q_1 R$$

is the QR decomposition of  $X[:, E]$ .

**qre** 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 **qqre**. If you need the entire  $Q$  matrix, call **qyre** 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 **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:

$$R = \begin{bmatrix} R_{11} & R_{12} \\ 0 & 0 \end{bmatrix}$$

where  $R_{11}$  is  $M \times M$  and  $R_{12}$  is  $M \times (P-M)$ . Then

$$A = R_{11}^{-1} R_{12}$$

and

$$X_2 = X_1 A$$

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:

$$Q'X = [ R_1 \ R_2 ]$$

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

$$b = \text{qrsol}(Q'Y, RI) | \text{zeros}(N-P, 1);$$

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 `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×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×P upper triangular matrix,  $K = \min(N,P)$ .  
             *e*      P×1 permutation vector.

REMARKS    **qrep** is the same as **qqrep** but doesn't return the  $Q_1$  matrix. If  $Q_1$  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

$$Q = [ Q_1 \ Q_2 ]$$

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*.

q

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 = qrsol(b,R);`

INPUT       $b$             P×L matrix.  
             $R$             P×P upper triangular matrix.

OUTPUT     $x$             P×L matrix.

REMARKS    **qrsol** applies a backsolve to  $Rx = b$  to solve for  $x$ . Generally  $R$  will be the  $R$  matrix from a QR factorization. **qrsol** may be used, however, in any situation where  $R$  is upper triangular.

SOURCE    `qrsol.src`

SEE ALSO    **qqr**, **qr**, **qtyr**, **qrtsol**

## qrtsol

PURPOSE	Computes the solution of $R'x = b$ where $R$ is an upper triangular matrix.	
FORMAT	$x = \mathbf{qrtsol}(b, R);$	
INPUT	$b$	P×L matrix.
	$R$	P×P upper triangular matrix.
OUTPUT	$x$	P×L matrix.
REMARKS	<p><b>qrtsol</b> applies a forward solve to <math>R'x = b</math> to solve for <math>x</math>. Generally <math>R</math> will be the <math>R</math> matrix from a QR factorization. <b>qrtsol</b> may be used, however, in any situation where <math>R</math> is upper triangular. If <math>R</math> is lower triangular, transpose before calling <b>qrtsol</b>.</p> <p>If <math>R</math> is not transposed, use <b>qrsol</b>.</p>	
SOURCE	qrsol.src	
SEE ALSO	<b>qqr, qr, qtyr, qrsol</b>	

## qtyr

PURPOSE	Computes the orthogonal-triangular (QR) decomposition of a matrix $X$ and returns $Q'Y$ and $R$ .	
FORMAT	$\{ \mathit{qty}, r \} = \mathbf{qtyr}(y, x);$	
INPUT	$y$	N×L matrix.



---

	$x$	N×P matrix.
OUTPUT	$qty$	N×L unitary matrix.
	$r$	K×P upper triangular matrix, K = min(N,P).
REMARKS	Given X, there is an orthogonal matrix $Q$ such that $Q'X$ is zero below its diagonal, i.e.,	

$$Q'X = \begin{bmatrix} R \\ 0 \end{bmatrix}$$

q

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  $Q'_1Y$  where  $Y$  is an N×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.  $Q'_1Y$  will be a submatrix of  $Q'Y$ . In particular,

$$G = Q'_1Y = qty[1 : P, .]$$

and  $Q'_2Y$  is the remaining submatrix:

$$H = Q'_2Y = qty[P + 1 : N, .]$$

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

$$s_j = \sum_{i=1}^{N-P} H_{i,j}, \quad j = 1, 2, \dots, L$$

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 superior numerical 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),.];
b = qrsol(q1ty,r);      /* LS coefficients */
s2 = sumc(q2ty^2);      /* residual sums of squares */
```

**SOURCE** qtyr.src

**SEE ALSO** **qqr**, **qtyre**, **qtyrep**, **olsqr**

qtyre

**PURPOSE**     Computes the orthogonal-triangular (QR) decomposition of a matrix  $X$  and returns  $Q'Y$  and  $R$ .

**FORMAT**     {  $qty, r, e$  } = **qtyre**( $y, x$ );

**INPUT**        $y$              $N \times L$  matrix.  
               $x$              $N \times P$  matrix.

**OUTPUT**      $qty$            $N \times L$  unitary matrix.  
               $r$              $K \times P$  upper triangular matrix,  $K = \min(N, P)$ .  
               $e$              $P \times 1$  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  $X[., E]$ .

q

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

$$X_2 = X_1 A$$

that is,  $A$  is an  $M \times (P-M)$  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,

$$Q_1'Y = qty[1 : P, .]$$

and  $Q_2'Y$  is the remaining submatrix:

$$Q_2'Y = qty[P + 1 : N, .]$$

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**:

$$b = \text{qrsol}(C, A);$$

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

$$b = \text{qrsol}(Q'Y, RI) | \text{zeros}(N-P, 1);$$

SOURCE    `qtyr.src`

SEE ALSO    **qqr, qre, qtyr**

## qtyrep

**PURPOSE**    Computes the orthogonal-triangular (QR) decomposition of a matrix  $X$  using a pivot vector and returns  $Q'Y$  and  $R$ .

**FORMAT**    `{ qty, r, e } = qtyrep(y, x, 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**     $qty$              $N \times L$  unitary matrix.

$r$              $K \times P$  upper triangular matrix,  $K = \min(N, P)$ .

$e$              $P \times 1$  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  $X[:, E]$ .

**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*.

```
EXAMPLE  y = { 4 7 2,
               5 9 1,
               6 3 3 };
          x = { 12 9 5,
               4 3 5,
               4 2 7 };
          pvt = { 11, 10, 3 };

          { qty,r,e } = qtyrep(y,x,pvt);
```

q

## quantile

---

```
      -6.9347609      -9.9498744      -3.0151134
qty =  4.0998891  3.5527137e - 15    2.1929640
      3.4785054      6.3245553    0.3162277
```

```
      -13.266499      -9.6483630      -8.1408063
r =   0.0000000  -0.95346259    4.7673129
      0.0000000    0.0000000    3.1622777
```

```
      1.0000000
e =   2.0000000
      3.0000000
```

SOURCE    `qtyr.src`

SEE ALSO    `qrep`, `qtyre`

## quantile

PURPOSE    Computes quantiles from data in a matrix, given specified probabilities.

FORMAT    `y = quantile(x,e)`

INPUT    *x*            N×K matrix of data.  
          *e*            L×1 vector, **quantile** levels or probabilities.

OUTPUT    *y*            L×K matrix, quantiles.

REMARKS    **quantile** will not succeed if N\*minc(e) is less than 1, or N\*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.



EXAMPLE    `rndseed 345567;`

```
x = rndn(1000,4); /* data */
e = { .025, .5, .975 }; /* quantile levels */
y = quantile(x,e);
```

```
print "medians";
print y[2,.];
print;
print "95 percentiles";
print y[1,.];
print y[3,.];
```

produces:

medians

-0.0020	-0.0408	-0.0380	-0.0247
---------	---------	---------	---------

95 percentiles

-1.8677	-1.9894	-2.1474	-1.8747
---------	---------	---------	---------

1.9687	2.0899	1.8576	2.0545
--------	--------	--------	--------

SOURCE    `quantile.src`

q

quantiled

PURPOSE    Computes quantiles from data in a data set, given specified probabilities.

FORMAT    `y = quantiled(dataset,e,var);`

INPUT    *dataset*    string, data set name, or N×M matrix of data.

## quantiled

---

*e*             $L \times 1$  vector, quantile levels or probabilities.  
*var*           $K \times 1$  vector or scalar zero. If  $K \times 1$ , character vector of labels selected for analysis, or numeric vector of column numbers in data set of variables selected for analysis. If scalar zero, all columns are selected.  
If dataset is a matrix *var* cannot be a character vector.

OUTPUT     *y*             $L \times K$  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.

Example:

```
y = quantiled("tobit",e,0);  
  
print "medians";  
print y[2,.];  
print;  
print "95 percentiles";  
print y[1,.];  
print y[3,.];
```

produces:

medians

0.0000	1.0000	-0.0021	-0.1228
--------	--------	---------	---------

95 percentiles

-1.1198	1.0000	-1.8139	-2.3143
2.3066	1.0000	1.4590	1.6954

---

SOURCE    `quantile.src`

**qyr**

PURPOSE    Computes the orthogonal-triangular (QR) decomposition of a matrix  $X$  and returns  $QY$  and  $R$ .

FORMAT    `{  $qy, r$  } = qyr(y, x);`

INPUT       $y$              $N \times L$  matrix.  
               $x$              $N \times P$  matrix.

OUTPUT     $qy$              $N \times L$  unitary matrix.  
               $r$              $K \times P$  upper triangular matrix,  $K = \min(N, P)$ .

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

$$X = Q_1 R$$

**q**

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**.

EXAMPLE     $x = \{ 1 \ 11, \ 7 \ 3, \ 2 \ 1 \};$   
                $y = \{ 2 \ 6, \ 5 \ 10, \ 4 \ 3 \};$   
                $\{ qy, \ r \} = \text{qyr}(y, x);$

$$qy = \begin{matrix} & 4.6288991 & 9.0506281 \\ -3.6692823 & & -7.8788202 \\ 3.1795692 & & 1.0051489 \end{matrix}$$

$$r = \begin{matrix} -7.3484692 & -4.6268140 \\ 0.0000000 & 10.468648 \end{matrix}$$

SOURCE    `qyr.src`

SEE ALSO    **qqr, qyre, qyrep, olsqr**

## 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       $y$              $N \times L$  matrix.

	$x$	$N \times P$ matrix.
OUTPUT	$qy$	$N \times L$ unitary matrix.
	$r$	$K \times P$ upper triangular matrix, $K = \min(N, P)$ .
	$e$	$P \times 1$ 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  $X[:, E]$ .

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  $N \times L$  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:

$$Q'X[:, E] = [ R_1 \ R_2 ]$$

## qyrep

---

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  $R_2$ .

EXAMPLE     $x = \{ 1 \ 11, \ 7 \ 3, \ 2 \ 1 \};$   
               $y = \{ 2 \ 6, \ 5 \ 10, \ 4 \ 3 \};$   
               $\{ qy, \ r, \ e \} = qyre(y,x);$

$$qy = \begin{array}{rr} -0.59422765 & -3.0456088 \\ -6.2442636 & -11.647846 \\ 2.3782485 & -0.22790230 \end{array}$$
$$r = \begin{array}{rr} -11.445523 & -2.9705938 \\ 0.0000000 & -6.7212776 \end{array}$$
$$e = \begin{array}{r} 2.0000000 \\ 1.0000000 \end{array}$$

SOURCE    `qyr.src`

SEE ALSO    **qqr**, **qre**, **qyr**

## qyrep

PURPOSE    Computes the orthogonal-triangular (QR) decomposition of a matrix  $X$  using a pivot vector and returns  $QY$  and  $R$ .

FORMAT     $\{ qy, r, e \} = \mathbf{qyrep}(y, x, pvt);$

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,  $K = \min(N, P)$ .  
             *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

## rank

---

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:

$$Q'X[:, E] = [ R_1 \ R_2 ]$$

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  $R_2$ .

SOURCE    `qyr.src`

SEE ALSO    **qr, qqrep, qrep, qtyrep**

## rank

PURPOSE    Computes the rank of a matrix, using the singular value decomposition.

FORMAT     $k = \text{rank}(x);$

INPUT       $x$              $N \times P$  matrix.

GLOBAL    **\_svdtol**            scalar, the tolerance used in determining if any of the  
INPUT                            singular values are effectively 0. The default value is  $10e^{-13}$ .  
This can be changed before calling the procedure.



OUTPUT	<i>k</i>	an estimate of the rank of <i>x</i> . This equals the number of singular values of <i>x</i> that exceed a prespecified tolerance in absolute value.
GLOBAL OUTPUT	<code>_svderr</code>	scalar, if not all of the singular values can be computed <code>_svderr</code> will be nonzero.
SOURCE	<code>svd.src</code>	

rankindx

r

PURPOSE	Returns the vector of ranks of a vector.	
FORMAT	<code>y = rankindx(x,flag);</code>	
INPUT	<i>x</i>	N×1 vector.
	<i>flag</i>	scalar, 1 for numeric data or 0 for character data.
OUTPUT	<i>y</i>	N×1 vector containing the ranks of <i>x</i> . 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 <i>y</i> from N+1).
REMARKS	<b>rankindx</b> assigns different ranks to elements that have equal values (ties). Missing values are assigned the lowest ranks.	
EXAMPLE	<pre>let x = 12 4 15 7 8; r = rankindx(x,1);</pre>	

```
      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(fl,r);`

**INPUT** *fl* scalar, file handle of an open file.  
*r* scalar, number of rows to read.

**OUTPUT** *y* N×K matrix, the data read from the file.

**REMARKS** The first time a **readr** statement is encountered, the first *r* rows will be read. The next time it is encountered, the next *r* rows will be read in, and so on. If the end of the data set is reached before *r* 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

r

PURPOSE Returns the real part of  $x$ .

FORMAT  $zr = \text{real}(x);$

INPUT  $x$   $N \times K$  matrix or  $N$ -dimensional array.

OUTPUT  $zr$   $N \times K$  matrix or  $N$ -dimensional array, the real part of  $x$ .

REMARKS If  $x$  is not complex,  $zr$  will be equal to  $x$ .

EXAMPLE  $x = \{ \begin{matrix} 1 & 11, \\ 7i & 3, \\ 2+i & 1 \end{matrix} \};$

$zr = \text{real}(x);$

```

          1.0000000  11.0000000
zr = 0.0000000  3.0000000
      2.0000000  1.0000000

```

SEE ALSO **complex, imag**

## recode

---

### 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**

<i>x</i>	N×1 vector to be recoded (changed).
<i>e</i>	N×K matrix of 1's and 0's.
<i>v</i>	K×1 vector containing the new values to be assigned to the recoded variable.

**OUTPUT** *y* N×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 };  
  
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);  
e = e1~e2~e3~e4;  
  
v = { 1,
```

```

2,
3,
4 };

y = recode(x,e,v);

```

```

20
45
x = 32
63
29

```

```

0 0 0 0
0 0 1 0
e = 0 1 0 0
0 0 0 0
1 0 0 0

```

```

1
2
v = 3
4

```

```

20
3
y = 2
63
1

```

SOURCE    datatran.src

SEE ALSO    **code**, **substute**

**r**

## 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**  
          *val\_1* **for** *expression\_1*,  
          *val\_2* **for** *expression\_2*,  
          .  
          .  
          .  
          *val\_n* **for** *expression\_n*;

**INPUT**   *var*           literal, the new variable name.  
          *val*           scalar, value to be used if corresponding expression is TRUE.  
          *expression* 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)**

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:

$$y[t, .] = x[t, .] + a[1, .] * y[t - 1, .] + \dots + a[P, .] * y[t - P, .], t = P + 1, \dots, N$$

and

$$y[t, .] = y0[t, .], t = 1, \dots, P$$

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:

$$\begin{aligned} y[1,t] &= \text{rho}[1,1]*y[1,t-1] + x[t,.*]b + e[1,t] \\ y[2,t] &= \text{rho}[2,1]*y[2,t-1] + x[t,.*]b + e[2,t] \end{aligned}$$

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**, **recserrc**

## 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**  $x$   $N \times K$  or  $1 \times K$  matrix  
 $z$   $N \times K$  or  $1 \times K$  matrix.

**OUTPUT**  $y$   $N \times K$  matrix in which each column is a series generated by a recursion of the form:

$$\begin{aligned} y(1) &= x(1) + z(1) \\ y(t) &= y(t-1) * x(t) + z(t), t = 2, \dots, N \end{aligned}$$

**REMARKS** The following **GAUSS** code could be used to emulate **recsercp** when the number of rows in  $x$  and  $z$  is the same:

```
n = rows(x); /* assume here that rows(z) */
              /* is also n */
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;
```

r

## recserrc

---

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     $c1 = c[1, .];$   
               $n = \text{rows}(c) - 1;$   
               $y = \text{recsercp}(x, \text{trim}(c ./ c1, 1, 0));$   
               $p = c1 .* y[n, .];$

If  $\mathbf{x}$  is a scalar and  $\mathbf{c}$  is an  $(N+1) \times 1$  vector, the result  $\mathbf{p}$  will contain the value of the polynomial whose coefficients are given in  $\mathbf{c}$ . That is:

$$p = c[1, .] \cdot x^n + c[2, .] \cdot x^{(n-1)} + \dots + c[n+1, .];$$

Note that both  $\mathbf{x}$  and  $\mathbf{c}$  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  $\mathbf{x} = 2$ , and if  $\mathbf{c}$  contains the digits of the binary representation of a number, then  $\mathbf{p}$  will be the decimal representation of that number.

SEE ALSO    **recserar**, **recserrc**

## recserrc

PURPOSE    Computes a recursive series involving division.

FORMAT     $y = \text{recserrc}(x, z);$

INPUT	$x$	$1 \times K$ or $K \times 1$ vector.
	$z$	$N \times K$ matrix.

**OUTPUT**     $y$              $N \times K$  matrix in which each column is a series generated by a recursion of the form:

$$\begin{aligned} y[1] &= x \bmod z[1], & x &= \text{trunc}(x/z[1]) \\ y[2] &= x \bmod z[2], & x &= \text{trunc}(x/z[2]) \\ y[3] &= x \bmod z[3], & x &= \text{trunc}(x/z[3]) \\ &\vdots \\ y[n] &= x \bmod z[n] \end{aligned}$$

**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`

rerun

**PURPOSE**    Displays the most recently created graphics file.

## reshape

---

LIBRARY    pgraph

FORMAT    **rerun**;

REMARKS    **rerun** is used by the **endwind** function.

SOURCE    pcart.src

GLOBALS    **\_pcmdlin, \_pnotify, \_psilent, \_ptek, \_pzoom**

## reshape

PURPOSE    Reshapes a matrix.

FORMAT     $y = \text{reshape}(x, r, c);$

INPUT     $x$              $N \times K$  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.

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     $y = \text{reshape}(x, 2, 6);$

If  $x = \begin{matrix} 1 & 2 & 3 & 4 \\ 5 & 6 & 7 & 8 \\ 9 & 10 & 11 & 12 \end{matrix}$  then  $y = \begin{matrix} 1 & 2 & 3 & 4 & 5 & 6 \\ 7 & 8 & 9 & 10 & 11 & 12 \end{matrix}$

If  $x = \begin{matrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{matrix}$  then  $y = \begin{matrix} 1 & 2 & 3 & 4 & 5 & 6 \\ 7 & 8 & 9 & 1 & 2 & 3 \end{matrix}$

If  $x = \begin{matrix} 1 & 2 & 3 & 4 & 5 \\ 6 & 7 & 8 & 9 & 10 \\ 11 & 12 & 13 & 14 & 15 \end{matrix}$  then  $y = \begin{matrix} 1 & 2 & 3 & 4 & 5 & 6 \\ 7 & 8 & 9 & 10 & 11 & 12 \end{matrix}$

If  $x = \begin{matrix} 1 & 2 \\ 3 & 4 \end{matrix}$  then  $y = \begin{matrix} 1 & 2 & 3 & 4 & 1 & 2 \\ 3 & 4 & 1 & 2 & 3 & 4 \end{matrix}$

If  $x = 1$  then  $y = \begin{matrix} 1 & 1 & 1 & 1 & 1 & 1 \\ 1 & 1 & 1 & 1 & 1 & 1 \end{matrix}$

SEE ALSO **submat, vec**

**retp**

**PURPOSE** Returns from a procedure or keyword.

**FORMAT** **retp;**

**retp**( $x, y, \dots$ );

**REMARKS** For more details, see PROCEDURES AND KEYWORDS, Chapter 12.

## return

---

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,...);**

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  $x$   $N \times K$  matrix.

OUTPUT  $y$   $N \times K$  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  $x = \text{round}(\text{rndn}(5,3)*10);$   
 $y = \text{rev}(x);$

$$x = \begin{bmatrix} 10 & 7 & 8 \\ 7 & 4 & -9 \\ -11 & 0 & -3 \\ 3 & 18 & 0 \\ 9 & -1 & 20 \end{bmatrix}$$

$$y = \begin{bmatrix} 9 & -1 & 20 \\ 3 & 18 & 0 \\ -11 & 0 & -3 \\ 7 & 4 & -9 \\ 10 & 7 & 8 \end{bmatrix}$$

SEE ALSO **sortc**

rfft

PURPOSE Computes a real 1- or 2-D Fast Fourier transform.

FORMAT  $y = \text{rfft}(x);$

INPUT  $x$   $N \times K$  real 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**     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.

**EXAMPLE**      $x = \{ 6 \ 9, \ 8 \ 1 \};$   
                    $y = \text{rfft}(x);$

$$y = \begin{matrix} 6.0000000 & 1.0000000 \\ 1.5000000 & -2.5000000 \end{matrix}$$

**SEE ALSO**     **rfffti**, **fft**, **ffti**, **fftm**, **fftni**

## **rffti**

**PURPOSE**     Computes inverse real 1- or 2-D Fast Fourier transform.

**FORMAT**      $y = \text{rffti}(x);$

**INPUT**         $x$              $N \times K$  matrix.

**OUTPUT**        $y$              $L \times M$  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**.

**EXAMPLE**      $x = \{ 6 \ 1, \ 1.5 \ -2.5 \};$



---

```
y = rffti(x);
```

```

y =  6.0000000  9.0000000
    8.0000000  1.0000000

```

SEE ALSO **rfft**, **fft**, **ffti**, **fftm**, **fftm**

## rfftip

r

**PURPOSE** Computes an inverse real 1- or 2-D FFT. Takes a packed format FFT as input.

**FORMAT** `y = rfftip(x);`

**INPUT** `x` N×K matrix or K-length vector.

**OUTPUT** `y` L×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:

$$2^p \times 3^q \times 5^r \times 7^s, \quad \begin{array}{l} p, q, r \geq 0 \\ s = 0 \text{ or } 1 \end{array}$$

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

**rffftip** 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 **fffti**. Note, however, that **fffti** expects a full FFT, including negative frequency information, for input.

Do not pass **rffftip** the output from **rfft** or **rfftn**—it will return incorrect results. Use **rfffti** with those routines.

SEE ALSO **fft**, **ffti**, **fftm**, **fftmi**, **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 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:

$$2^p \times 3^q \times 5^r \times 7^s, \quad \begin{array}{l} p, q, r \geq 0 \text{ for rows of matrix} \\ p > 0, q, r \geq 0 \text{ for columns of matrix} \\ p > 0, q, r \geq 0 \text{ for length of vector} \\ s = 0 \text{ or } 1 \text{ for all dimensions} \end{array}$$

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 33600×1 vector can compute as much as 20 percent faster than a 32768×1 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  $1/(L \times M)$ .

SEE ALSO **fft, ffti, fftm, fftmi, fftn, rfft, rffti, rfftip, rfftnp, rfftp**

---

PURPOSE	Computes a real 1- or 2-D FFT. Returns the results in a packed format.	
FORMAT	$y = \mathbf{rfftnp}(x);$	
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	<p>For 1-D FFT's, <b>rfftnp</b> returns the positive frequencies in ascending order in the first <math>M/2</math> elements, and the Nyquist frequency in the last element. For 2-D FFT's, <b>rfftnp</b> 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 <math>M/2</math> 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 <b>rfft</b> and <b>rfftn</b> for routines that return the negative frequencies as well.)</p>	

**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:

$$2^p \times 3^q \times 5^r \times 7^s, \quad \begin{array}{l} p, q, r \geq 0 \quad \text{for rows of matrix} \\ p > 0, \quad q, r \geq 0 \quad \text{for columns of matrix} \\ p > 0, \quad q, r \geq 0 \quad \text{for length of vector} \\ s = 0 \text{ or } 1 \quad \text{for all dimensions} \end{array}$$

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 33600×1 vector can compute as much as 20 percent faster than a 32768×1 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 **rfftnp**. 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 **rfftnp** would pad a matrix or vector.)

**rfftnp** scales the computed FFT by  $1/(L \times M)$ .

SEE ALSO    **fft, ffti, fftm, fftmi, fftn, rfft, rffti, rfftip, rfftn, rfftp**

rfftp

PURPOSE    Computes a real 1- or 2-D FFT. Returns the results in a packed format.

FORMAT     $y = \text{rfftp}(x);$

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 powers of 2 greater than or equal to  $N$  and  $K$ , respectively.

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, **rfftp** 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**, **rfft**, **rffti**, **rfftp**, **rfftn**, **rfftnp**

## rndbeta

**PURPOSE**    Computes pseudo-random numbers with beta distribution.

**FORMAT**     $x = \text{rndbeta}(r, c, a, b);$

**INPUT**     $r$             scalar, number of rows of resulting matrix.  
               $c$             scalar, number of columns of resulting matrix.  
               $a$              $M \times N$  matrix,  $E \times E$  conformable with  $r \times c$  resulting matrix, shape parameters for beta distribution.  
               $b$              $K \times L$  matrix,  $E \times E$  conformable with  $r \times c$  resulting matrix, shape parameters for beta distribution.

**OUTPUT**     $x$              $r \times c$  matrix, beta distributed pseudo-random numbers.

**REMARKS**    The properties of the pseudo-random numbers in  $x$  are:

$$\begin{aligned} E(x) &= a/(a+b) \\ Var(x) &= a*b/((a+b+1)*(a+b)^2) \\ x &> 0 \\ x &< 1 \\ a &> 0 \\ b &> 0 \end{aligned}$$

SOURCE **random.src**

**r**

**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** *c* scalar, constant for the random number generator.  
*a* scalar, multiplier for the random number generator.  
*seed* scalar, initial seed for the random number generator.

Parameter default values and ranges:

<i>seed</i>	time(0)	$0 < seed < 2^{32}$
<i>a</i>	1664525	$0 < a < 2^{32}$
<i>c</i>	1013904223	$0 \leq c < 2^{32}$

**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.

**SEE ALSO** **rndu**, **rndn**, **rndi**, **rndLCi**, **rndKMi**



**rndgam**

**PURPOSE**     Computes pseudo-random numbers with gamma distribution.

**FORMAT**      $x = \text{rndgam}(r,c,alpha);$

**INPUT**      $r$             scalar, number of rows of resulting matrix.  
               $c$             scalar, number of columns of resulting matrix.  
               $alpha$          $M \times N$  matrix,  $E \times E$  conformable with  $r \times c$  resulting matrix, shape  
                                 parameters for gamma distribution.

**OUTPUT**      $x$              $r \times c$  matrix, gamma distributed pseudo-random numbers.

**REMARKS**     The properties of the pseudo-random numbers in  $x$  are:

$$\begin{aligned} E(x) &= alpha \\ Var(x) &= alpha \\ x &> 0 \\ alpha &> 0 \end{aligned}$$

To generate **gamma(alpha,theta)** pseudo-random numbers where  $theta$  is a scale parameter, multiply the result of **rndgam** by  $theta$ . Thus:

$$z = theta * \text{rndgam}(1,1,alpha);$$

has the properties

$$\begin{aligned}E(z) &= \alpha \times \theta \\ \text{Var}(z) &= \alpha \times \theta^2 \\ z &> 0 \\ \alpha &> 0 \\ \theta &> 0\end{aligned}$$

SOURCE    `random.src`

## rndi

PURPOSE    Returns a matrix of random integers,  $0 \leq y < 2^{32}$ .

FORMAT    `y = rndi(r,c);`

INPUT      *r*            scalar, row dimension.  
            *c*            scalar, column dimension.

OUTPUT    *y*             $r \times c$  matrix of random integers between 0 and  $2^{32}-1$ , inclusive.

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.

Each seed is generated from the preceding seed, using the formula

$$\text{new\_seed} = (((a * \text{seed}) \% 2^{32}) + c) \% 2^{32}$$

where **%** is the mod operator. The new seeds are the values returned. The multiplicative constant and the additive constant may be changed using **rndmult** and **rndcon** respectively.

SEE ALSO    **rndu, rndn, rndcon, rndmult**

**rndKmbeta**

**r**

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*×*c* matrix, or *r*×1 vector, or 1×*c* vector, or scalar, first shape argument for beta distribution.

*b*            *r*×*c* matrix, or *r*×1 vector, or 1×*c* vector, or scalar, second shape argument for beta distribution.

*state*       scalar or 500×1 vector.

**Scalar case:**

*state* = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

**500×1 vector case:**

*state* = the state vector returned from a previous call to one of the **rndKM** random number functions.

OUTPUT    *x*            *r*×*c* matrix, beta distributed random numbers.

*newstate*    500×1 vector, the updated state.

## rndKMgam

---

REMARKS     The properties of the pseudo-random numbers in  $x$  are:

$$E(x) = \frac{a}{a+b}, Var(x) = \frac{(a*b)}{(a+b+1)*(a+b)^2}$$

$$0 < x < 1, a > 0, b > 0$$

$r$  and  $c$  will be truncated to integers if necessary.

SOURCE     `randkm.src`

TECHNICAL     **rndKMbeta** uses the recur-with-carry KISS+Monster algorithm described in the  
NOTES     **rndKMi** Technical Notes.

## rndKMgam

PURPOSE     Computes Gamma pseudo-random numbers.

FORMAT     {  $x, newstate$  } = **rndKMgam**( $r, c, alpha, state$ );

INPUT      $r$             scalar, number of rows of resulting matrix.  
             $c$             scalar, number of columns of resulting matrix.  
             $alpha$          $r \times c$  matrix, or  $r \times 1$  vector, or  $1 \times c$  vector, or scalar, shape argument  
                            for gamma distribution.  
             $state$         scalar or  $500 \times 1$  vector.  
                            **Scalar case:**  
                             $state$  = starting seed value only. If -1, **GAUSS** computes the starting  
                            seed based on the system clock.  
                            **500×1 vector case:**  
                             $state$  = the state vector returned from a previous call to one of the  
                            **rndKM** random number functions.

OUTPUT     $x$              $r \times c$  matrix, gamma distributed random numbers.  
               $newstate$      $500 \times 1$  vector, the updated state.

REMARKS    The properties of the pseudo-random numbers in  $x$  are:

$$E(x) = \alpha, Var(x) = \alpha$$

$$x > 0, \alpha > 0$$

To generate **gamma**( $\alpha, \theta$ ) pseudo-random numbers where  $\theta$  is a scale parameter, multiply the result of **rndKMgam** by  $\theta$ .

Thus

$$z = \theta * \text{rndgam}(1, 1, \alpha);$$

has the properties

$$E(z) = \alpha * \theta, Var(z) = \alpha * \theta^2$$

$$z > 0, \alpha > 0, \theta > 0$$

$r$  and  $c$  will be truncated to integers if necessary.

SOURCE    `randkm.src`

TECHNICAL    **rndKMgam** uses the recur-with-carry KISS+Monster algorithm described in the  
 NOTES        **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** *r* scalar, row dimension.

*c* scalar, column dimension.

*state* scalar or 500×1 vector.

**Scalar case:**

*state* = starting seed value. If -1, **GAUSS** computes the starting seed based on the system clock.

**500×1 vector case:**

*state* = the state vector returned from a previous call to one of the **rndKM** random number generators.

**OUTPUT** *y* *r*×*c* matrix of random integers between 0 and  $2^{32} - 1$ , inclusive.

*newstate* 500×1 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 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>.

**r**

**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 *r* scalar, row dimension.  
*c* scalar, column dimension.  
*state* scalar or 500×1 vector.

**Scalar case:**

*state* = starting seed value. If -1, **GAUSS** computes the starting seed based on the system clock.

**500×1 vector case:**

*state* = the state vector returned from a previous call to one of the **rndKM** random number generators.

OUTPUT    *y*            *r*×*c* matrix of standard normal random numbers.  
          *newstate*    500×1 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.

```
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;  
end;  
  
mean = meanc(submean);  
print mean;
```

SEE ALSO    **rndKMu**, **rndKMi**

TECHNICAL    **rndKMn** calls the uniform random number generator that is the basis for **rndKMu**  
NOTES        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.

**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> × <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>p</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>	scalar or 500×1 vector.

**Scalar case:**

*state* = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

**500×1 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> × <i>c</i> matrix, negative binomial distributed random numbers.
<i>newstate</i>	500×1 vector, the updated state.

**REMARKS**     The properties of the pseudo-random numbers in *x* are:

$$E(x) = \frac{k * p}{(1 - p)}, Var(x) = \frac{k * p}{(1 - p)^2}$$

## rndKMp

---

$$x = 0, 1, \dots, k > 0, 0 < p < 1$$

$r$  and  $c$  will be truncated to integers if necessary.

SOURCE    `randkm.src`

TECHNICAL    **rndKMnb** uses the recur-with-carry KISS+Monster algorithm described in the  
NOTES    **rndKM**i Technical Notes.

### rndKMp

PURPOSE    Computes Poisson pseudo-random numbers.

FORMAT    {  $x, newstate$  } = **rndKMp**( $r, c, lambda, state$ );

INPUT     $r$             scalar, number of rows of resulting matrix.

$c$             scalar, number of columns of resulting matrix.

$lambda$      $r \times c$  matrix, or  $r \times 1$  vector, or  $1 \times c$  vector, or scalar, shape argument  
                          for Poisson distribution.

$state$       scalar or  $500 \times 1$  vector.

**Scalar case:**

$state$  = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.

**500×1 vector case:**

$state$  = the state vector returned from a previous call to one of the **rndKM** random number functions.

OUTPUT     $x$              $r \times c$  matrix, Poisson distributed random numbers.

$newstate$      $500 \times 1$  vector, the updated state.

REMARKS     The properties of the pseudo-random numbers in  $x$  are:

$$E(x) = \lambda, Var(x) = \lambda$$

$$x = 0, 1, \dots, \lambda > 0$$

$r$  and  $c$  will be truncated to integers if necessary.

SOURCE     `randkm.src`

TECHNICAL     **rndKMp** uses the recur-with-carry KISS+Monster algorithm described in the  
NOTES     **rndKMi** Technical Notes.

**r**

**rndKMu**

PURPOSE     Returns a matrix of uniform (pseudo) random variables and the state of the random number generator.

FORMAT     {  $y, newstate$  } = **rndKMu**( $r, c, state$ );

INPUT      $r$             scalar, row dimension.  
             $c$             scalar, column dimension.  
             $state$        scalar, 2×1 vector, or 500×1 vector.  
                 **Scalar case:**  
                  $state$  = starting seed value. If -1, **GAUSS** computes the starting seed based on the system clock.  
                 **2×1 vector case:**  
                 [ 1 ] the starting seed, uses the system clock if -1  
                 [ 2 ] 0 for 0 ≤  $y$  < 1  
                        1 for 0 ≤  $y$  ≤ 1

### **500×1 vector case:**

*state* = the state vector returned from a previous call to one of the **rndKM** random number generators.

OUTPUT    *y*            *r*×*c* matrix of uniform random numbers,  $0 \leq y < 1$ .  
          *newstate*    500×1 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;  
end;  
  
mean = meanc(submean);  
print 0.5-mean;
```

SEE ALSO    **rndKMn**, **rndKMi**

TECHNICAL    **rndKMu** uses the recur-with-carry KISS-Monster algorithm described in the  
NOTES        **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.

**FORMAT**     { *x*, *newstate* } = **rndKMvm**(*r*, *c*, *m*, *k*, *state*);

**INPUT**     *r*            scalar, number of rows of resulting matrix.  
              *c*            scalar, number of columns of resulting matrix.  
              *m*            *r*×*c* matrix, or *r*×1 vector, or 1×*c* vector, or scalar, means for vm distribution.  
              *k*            *r*×*c* matrix, or *r*×1 vector, or 1×*c* vector, or scalar, shape argument for vm distribution.  
              *state*        scalar or 500×1 vector.

**Scalar case:**  
*state* = starting seed value only. If -1, **GAUSS** computes the starting seed based on the system clock.  
**500×1 vector case:**  
*state* = the state vector returned from a previous call to one of the **rndKM** random number functions.

**OUTPUT**     *x*            *r*×*c* matrix, von Mises distributed random numbers.  
              *newstate*    500×1 vector, the updated state.

**REMARKS**     *r* and *c* will be truncated to integers if necessary.

**SOURCE**     randkm.src

**TECHNICAL**     **rndKMvm** uses the recur-with-carry KISS+Monster algorithm described in the  
**NOTES**           **rndKM**i Technical Notes.

## rndLCbeta

**PURPOSE**    Computes beta pseudo-random numbers.

**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>	$r \times c$ matrix, or $r \times 1$ vector, or $1 \times c$ vector, or scalar, first shape argument for beta distribution.
<i>b</i>	$r \times c$ matrix, or $r \times 1$ vector, or $1 \times c$ vector, or scalar, second shape argument for beta distribution.
<i>state</i>	scalar, or $3 \times 1$ vector, or $4 \times 1$ 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.

### $3 \times 1$ vector case:

[ 1 ] the starting seed, uses the system clock if -1

[ 2 ] the multiplicative constant

[ 3 ] the additive constant

### $4 \times 1$ vector case:

*state* = the state vector returned from a previous call to one of the **rndLC** random number generators.

**OUTPUT**

<i>x</i>	$r \times c$ matrix, beta distributed random numbers.
<i>newstate</i>	$4 \times 1$ vector:
	[ 1 ] the updated seed

- [ 2 ] the multiplicative constant
- [ 3 ] the additive constant
- [ 4 ] the original initialization seed

REMARKS    The properties of the pseudo-random numbers in  $x$  are:

$$E(x) = \frac{a}{a+b}, Var(x) = \frac{(a*b)}{(a+b+1)*(a+b)^2}$$

$$0 < x < 1, a > 0, b > 0$$

$r$  and  $c$  will be truncated to integers if necessary.

SOURCE    `randlc.src`

TECHNICAL    This function uses a linear congruential method, discussed in Kennedy, W.J. Jr.,  
NOTES        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.

FORMAT    {  $x, newstate$  } = **rndLCgam**( $r, c, alpha, state$ );

INPUT     $r$             scalar, number of rows of resulting matrix.

## rndLCgam

---

*c* scalar, number of columns of resulting matrix.

*alpha*  $r \times c$  matrix, or  $r \times 1$  vector, or  $1 \times c$  vector, or scalar, shape argument for gamma distribution.

*state* scalar, or  $3 \times 1$  vector, or  $4 \times 1$  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.

### $3 \times 1$ vector case:

[ 1 ] the starting seed, uses the system clock if -1

[ 2 ] the multiplicative constant

[ 3 ] the additive constant

### $4 \times 1$ 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, gamma distributed random numbers.

*newstate*  $4 \times 1$  vector:

[ 1 ] the updated seed

[ 2 ] the multiplicative constant

[ 3 ] the additive constant

[ 4 ] the original initialization seed

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(alpha,theta)** pseudo-random numbers where *theta* is a scale parameter, multiply the result of **rndLCgam** by *theta*.

Thus

$$z = theta * rndgam(1,1,alpha);$$

has the properties

$$E(z) = alpha * theta, Var(z) = alpha * theta^2$$

$$z > 0, alpha > 0, theta > 0$$

*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.

**rndLCi**

**PURPOSE**    Returns a matrix of random integers,  $0 \leq y < 2^{32}$ , and the state of the random number generator.

## **rndLCi**

---

FORMAT    { *y,newstate* } = **rndLCi**(*r,c,state*);

INPUT    *r*            scalar, row dimension.  
          *c*            scalar, column dimension.  
          *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*<0, **GAUSS** computes the starting seed based on the system clock.

**3×1 vector case:**

[ 1 ] the starting seed, uses the system clock if < 0

[ 2 ] the multiplicative constant

[ 3 ] the additive constant

**4×1 vector case:**

*state* = the state vector returned from a previous call to one of the **rndLC** random number generators.

OUTPUT    *y*            *r*×*c* matrix of random integers between 0 and  $2^{32} - 1$ , inclusive.  
          *newstate*    4×1 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 = 20000000000;
k = 10000000;
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**

**r**

**rndLCn**

**PURPOSE** Returns a matrix of standard normal (pseudo) random variables and the state of the random number generator.

**FORMAT** { *y,newstate* } = **rndLCn**(*r,c,state*);

**INPUT** *r* scalar, row dimension.  
*c* scalar, column dimension.  
*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* < 0, **GAUSS** computes the starting seed based on the system clock.

### **3×1 vector case:**

[ 1 ] the starting seed, uses the system clock if < 0

[ 2 ] the multiplicative constant

[ 3 ] the additive constant

### **4×1 vector case:**

*state* = the state vector returned from a previous call to one of the **rndLC** random number generators.

OUTPUT    *y*            *r*×*c* matrix of standard normal random numbers.

*newstate*    4×1 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 = 20000000000;  
k = 1000000;  
c = 0;  
submean = {};  
  
do while c < n;  
    { y,state } = rndLCn(k,1,state);  
    submean = submean | meanc(y);  
    c = c + k;  
end;`

```
mean = meanc(submean);  
print mean;
```

SEE ALSO    **rndLCu, rndLCi, rndcon, rndmult**

TECHNICAL    The normal random number generator is based on the uniform random number  
NOTES       generator, 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. 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.

**r**

**rndLCnb**

PURPOSE    Computes negative binomial pseudo-random numbers.

FORMAT    { *x, newstate* } = **rndLCnb**(*r, c, k, p, state*);

- INPUT    *r*            scalar, number of rows of resulting matrix.
- c*            scalar, number of columns of resulting matrix.
- k*            *r*×*c* matrix, or *r*×1 vector, or 1×*c* vector, or scalar, “event” argument  
                         for negative binomial distribution.
- p*            *r*×*c* matrix, or *r*×1 vector, or 1×*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.

**3×1 vector case:**

[ 1 ] the starting seed, uses the system clock if -1

[ 2 ] the multiplicative constant

[ 3 ] the additive constant

**4×1 vector case:**

*state* = the state vector returned from a previous call to one of the **rndLC** random number generators.

OUTPUT    *x*            *r*×*c* matrix, negative binomial distributed random numbers.  
              *newstate*    4×1 vector:  
                          [ 1 ] the updated seed  
                          [ 2 ] the multiplicative constant  
                          [ 3 ] the additive constant  
                          [ 4 ] the original initialization seed

REMARKS    The properties of the pseudo-random numbers in *x* are:

$$E(x) = \frac{k * p}{(1 - p)}, Var(x) = \frac{k * p}{(1 - p)^2}$$

$$x = 0, 1, \dots, k > 0, 0 < p < 1$$

*r* and *c* will be truncated to integers if necessary.

SOURCE    `randlc.src`

TECHNICAL    This function uses a linear congruential method, discussed in Kennedy, W.J. Jr.,  
 NOTES        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.

**FORMAT**     { *x*,*newstate* } = **rndLCp**(*r*,*c*,*lambda*,*state*);

**INPUT**

*r*               scalar, number of rows of resulting matrix.

*c*               scalar, number of columns of resulting matrix.

*lambda*        *r*×*c* matrix, or *r*×1 vector, or 1×*c* vector, or scalar, shape argument for Poisson 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.

**3×1 vector case:**

[ 1 ] the starting seed, uses the system clock if -1

[ 2 ] the multiplicative constant

[ 3 ] the additive constant

**4×1 vector case:**

*state* = the state vector returned from a previous call to one of the **rndLC** random number generators.

**OUTPUT**     *x*                *r*×*c* matrix, Poisson distributed random numbers.

**r**

## **rndLCu**

---

*newstate* 4×1 vector:  
[ 1 ] the updated seed  
[ 2 ] the multiplicative constant  
[ 3 ] the additive constant  
[ 4 ] the original initialization seed

REMARKS The properties of the pseudo-random numbers in  $x$  are:

$$E(x) = \textit{lambda}, \textit{Var}(x) = \textit{lambda}$$

$$x = 0, 1, \dots, \textit{lambda} > 0$$

$r$  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

$$\textit{new\_seed} = (((a * \textit{seed}) \% 2^{32}) + c) \% 2^{32}$$

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.

FORMAT {  $y, \textit{newstate}$  } = **rndLCu**( $r, c, \textit{state}$ );





INPUT	<i>r</i>	scalar, row dimension.
	<i>c</i>	scalar, column dimension.
	<i>state</i>	scalar, or 3×1 vector, or 4×1 vector. <b>Scalar case:</b> <i>state</i> = 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 <b>rndcon</b> and <b>rndmult</b> . If <i>state</i> <0, <b>GAUSS</b> computes the starting seed based on the system clock. <b>3×1 vector case:</b> [ 1 ] the starting seed, uses the system clock if < 0 [ 2 ] the multiplicative constant [ 3 ] the additive constant <b>4×1 vector case:</b> <i>state</i> = the state vector returned from a previous call to one of the <b>rndLC</b> random number generators.
OUTPUT	<i>y</i>	<i>r</i> × <i>c</i> matrix of uniform random numbers, 0<= <i>y</i> <1.
	<i>newstate</i>	4×1 vector: [ 1 ] the updated seed [ 2 ] the multiplicative constant [ 3 ] the additive constant [ 4 ] the original initialization seed
REMARKS	<i>r</i> and <i>c</i> 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<sup>32</sup>.

## **rndLCvm**

---

EXAMPLE     `state = 13;`  
              `n = 20000000000;`  
              `k = 10000000;`  
              `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     This function uses a linear congruential method, discussed in Kennedy, W. J. Jr.,  
NOTES           and J. E. Gentle, *Statistical Computing*, Marcel Dekker, Inc., 1980, pp. 136-147.

## **rndLCvm**

PURPOSE     Computes von Mises pseudo-random numbers.

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>	$r \times c$ matrix, or $r \times 1$ vector, or $1 \times c$ vector, or scalar, means for vm distribution.
	<i>k</i>	$r \times c$ matrix, or $r \times 1$ vector, or $1 \times c$ vector, or scalar, shape argument for vm distribution.

	<i>state</i>	<p>scalar, or 3×1 vector, or 4×1 vector.</p> <p><b>Scalar case:</b>  <i>state</i> = 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 <b>rndcon</b> and <b>rndmult</b>.  If <i>state</i> = -1, <b>GAUSS</b> computes the starting seed based on the system clock.</p> <p><b>3×1 vector case:</b>  [ 1 ] the starting seed, uses the system clock if -1  [ 2 ] the multiplicative constant  [ 3 ] the additive constant</p> <p><b>4×1 vector case:</b>  <i>state</i> = the state vector returned from a previous call to one of the <b>rndLC</b> random number generators.</p>
OUTPUT	<i>x</i>  <i>newstate</i>	<p><i>r</i>×<i>c</i> matrix, von Mises distributed random numbers.</p> <p>4×1 vector:  [ 1 ] the updated seed  [ 2 ] the multiplicative constant  [ 3 ] the additive constant  [ 4 ] the original initialization seed</p>
REMARKS		<i>r</i> and <i>c</i> will be truncated to integers if necessary.
SOURCE		randlc.src
TECHNICAL NOTES		<p>This function uses a linear congruential method, discussed in Kennedy, W.J. Jr., and J.E. Gentle, <i>Statistical Computing</i>, Marcel Dekker, Inc. 1980, pp. 136-147. Each seed is generated from the preceding seed using the formula</p> $new\_seed = (((a * seed) \% 2^{32}) + c) \% 2^{32}$ <p>where <b>%</b> is the mod operator and where <i>a</i> is the multiplicative constant and <i>c</i> is the additive constant.</p>

### **rndn**

**PURPOSE**     Creates a matrix of standard Normal (pseudo) random numbers.

**FORMAT**      $y = \text{rndn}(r, c);$

**INPUT**        $r$             scalar, row dimension.  
                  $c$             scalar, column dimension.

**OUTPUT**      $y$              $r \times c$  matrix of Normal random numbers having a mean of 0 and standard deviation of 1.

**REMARKS**      $r$  and  $c$  will be truncated to integers if necessary.

The Normal random number generator is based upon the uniform random number generator. To reseed them both, use the **rndseed** statement. The other parameters of the uniform generator can be changed using **rndcon**, **rndmod**, and **rndmult**.

**EXAMPLE**      $x = \text{rndn}(8100, 1);$   
                  $m = \text{meanc}(x);$   
                  $s = \text{stdc}(x);$

$m = 0.002810$

$s = 0.997087$

In this example, a sample of 8100 Normal random numbers is drawn, and the mean and standard deviation are computed for the sample.

**SEE ALSO**     **rndu**, **rndcon**

**TECHNICAL NOTES**     This function uses 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*. Vol. 71 No. 356, Dec. 1976, 893-96.

**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.  
               $c$            scalar, number of columns of resulting matrix.  
               $k$            M×N matrix, E×E conformable with  $r \times c$  resulting matrix, “event” parameters for negative binomial distribution.  
               $p$            K×L matrix, E×E 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

## rndp

---

### rndp

**PURPOSE**    Computes pseudo-random numbers with Poisson distribution.

**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.
$\text{lambda}$	M×N matrix, E×E 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:

$$\begin{aligned} E(x) &= \text{lambda} \\ \text{Var}(x) &= \text{lambda} \\ x &= 0, 1, 2, \dots \\ \text{lambda} &> 0 \end{aligned}$$

**SOURCE**    random.src

### rndu

**PURPOSE**    Creates a matrix of uniform (pseudo) random variables.

**FORMAT**     $y = \text{rndu}(r, c);$

INPUT	<i>r</i>	scalar, row dimension.
	<i>c</i>	scalar, column dimension.
OUTPUT	<i>y</i>	<i>r</i> × <i>c</i> matrix of uniform random variables between 0 and 1.
REMARKS	<i>r</i> and <i>c</i> will be truncated to integers if necessary.	

This generator is automatically seeded using the clock when **GAUSS** is first started. However, that can be overridden using the **rndseed** statement.

The seed is automatically updated as a random number is generated (see above under **rndcon**). Thus, 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. This is a 32-bit generator, though, so the range is sufficient for most applications.

EXAMPLE    *x* = rndu(8100,1);  
              *y* = meanc(*x*);  
              *z* = stdc(*x*);

*y* = 0.500205

*z* = 0.289197

In this example, a sample of 8100 uniform random numbers is generated, and the mean and standard deviation are computed for the sample.

SEE ALSO    **rndn, rndcon, rndmod, rndmult, rndseed**

TECHNICAL    This function uses a multiplicative-congruential method. This method is  
NOTES        discussed in Kennedy, W.J., Jr., and J.E. Gentle. *Statistical Computing*. Marcel Dekker, Inc., NY, 1980, 136-147.

r

## rotater

---

### 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, $E \times E$ conformable with $r \times c$ , means for von Mises distribution.
	$k$	$L \times M$ matrix, $E \times E$ 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	

### rotater

PURPOSE	Rotates the rows of a matrix.	
FORMAT	$y = \text{rotater}(x, r);$	
INPUT	$x$	$N \times K$ matrix to be rotated.
	$r$	$N \times 1$ or $1 \times 1$ 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* % **cols**(*x*)).

EXAMPLE     *y* = **rotater**(*x*,*r*);

If  $x = \begin{matrix} 1 & 2 & 3 \\ 4 & 5 & 6 \end{matrix}$  and  $r = \begin{matrix} 1 \\ -1 \end{matrix}$  Then  $y = \begin{matrix} 3 & 1 & 2 \\ 5 & 6 & 4 \end{matrix}$

If  $x = \begin{matrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \\ 10 & 11 & 12 \end{matrix}$  and  $r = \begin{matrix} 0 \\ 1 \\ 2 \\ 3 \end{matrix}$  Then  $y = \begin{matrix} 1 & 2 & 3 \\ 6 & 4 & 5 \\ 8 & 9 & 7 \\ 10 & 11 & 12 \end{matrix}$

r

SEE ALSO     **shiftr**

round

PURPOSE     Round to the nearest integer.

FORMAT     *y* = **round**(*x*);

INPUT       *x*             N×K matrix or N-dimensional array.

OUTPUT     *y*             N×K matrix or N-dimensional array containing the rounded elements of *x*.

EXAMPLE     let *x* = { 77.68     -14.10,

## rows

---

```
          4.73   -158.88 };  
y = round(x);
```

```
      78.00   -14.00  
y =    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`  $N \times K$  matrix or sparse matrix.

**OUTPUT** `y` scalar, number of rows in the specified matrix.

**REMARKS** If `x` is an empty matrix, **rows(x)** and **cols(x)** return 0.

**EXAMPLE** `x = ones(3,5);`  
`y = rows(x);`

```
      1  1  1  1  1  
x =   1  1  1  1  1  
      1  1  1  1  1
```

```
y = 3
```

SEE ALSO **cols, show**

rowsf

PURPOSE

Returns the number of rows in a **GAUSS** data set (.dat) file or **GAUSS** matrix (.fmt) file.

FORMAT

$y = \text{rowsf}(f);$

INPUT

$f$  file handle of an open file.

OUTPUT

$y$  scalar, number of rows in the specified file.

EXAMPLE

```
open fp = myfile;
r = rowsf(fp);
c = colsf(fp);
```

SEE ALSO

colsf, open, typef

r

rref

PURPOSE

Computes the reduced row echelon form of a matrix.

FORMAT

$y = \text{rref}(x);$

INPUT

$x$  M×N matrix.

OUTPUT

$y$  M×N matrix containing reduced row echelon form of  $x$ .

REMARKS

The tolerance used for zeroing elements is computed inside the procedure using:

$$tol = \maxc(m|n) * eps * \maxc(abs(\text{sumc}(x')));$$

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    `let x[3,3] = 1 2 3  
                  4 5 6  
                  7 8 9;  
y = rref(x);`

$$y = \begin{bmatrix} 1 & 0 & -1 \\ 0 & 1 & 2 \\ 0 & 0 & 0 \end{bmatrix}$$

SOURCE    `rref.src`

---

**run**

PURPOSE    Runs a source code or compiled code program.

FORMAT **run** *filename*;

INPUT *filename* literal or ^string, name of file to run.

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.

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.

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**.

<b>run /gauss/myprog.prg;</b>	No additional search will be made if the file is not found.
<b>run myprog.prg;</b>	The directories listed in <b>src_path</b> will be searched for <b>myprog.prg</b> 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

r

## satostrC

---

starting **GAUSS**.

### EXAMPLE    **Example 1**

```
run myprog.prg;
```

### **Example 2**

```
name = "myprog.prg";  
run ^name;
```

SEE ALSO    **#include**

## satostrC

**PURPOSE**    Copies from one string array to another using a C language format specifier string for each element.

**FORMAT**     $y = \text{satostrC}(sa,fmt);$

**INPUT**    *sa*             $N \times M$  string array.  
          *fmt*             $1 \times 1$ ,  $1 \times M$ , or  $M \times 1$  format specifier for each element copy.

**OUTPUT**    *y*             $N \times M$  formatted string array.

**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**;

INPUT     *vflag*            version flag.

- v89            not supported
- v92            supported on UNIX, Windows
- v96            supported on all platforms

See also FILE I/O, Chapter 20, 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**.

*path*            literal or ^string, a default path to use for this and subsequent **save**'s.

*x*                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.

*lpath*           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.

*y*                the symbol to be saved to *lpath*.

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:

S

## save

---

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;
```



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** *fname* 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,

S

## saved

---

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** *x*             $N \times K$  matrix to save in .dat file.  
*dataset*        string, name of data set.  
*vnames*        string or  $K \times 1$  character vector, names for the columns 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    **loadadd, writer, create**

## savestruct

**S**

PURPOSE    Saves a matrix of structures to a file on the disk.

FORMAT    *retcode* = **saveStruct**(*instance*,*file\_name*);

INPUT    *instance*    M×N matrix, instances of a structure.  
           *file\_name*   string, name of file on disk to contain matrix of structures.

OUTPUT    *retcode*    scalar, 0 if successful, otherwise 1.

REMARKS    The file on the disk will be given a .fsr extension

EXAMPLE    #include ds.sdf

```

struct DS p0;
p0 = reshape(dsCreate,2,3);
retc = saveStruct(p2,"p2");

```

## scale

---

### savewind

PURPOSE	Save the current graphic panel configuration to a file.	
LIBRARY	pgraph	
FORMAT	<i>err</i> = <b>savewind</b> ( <i>filename</i> );	
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 GRAPHIC PANELS, Section <a href="#">24.3</a> .	
SOURCE	pwindow.src	
SEE ALSO	<b>loadwind</b>	

### 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.	
LIBRARY	pgraph	
FORMAT	<b>scale</b> ( <i>x</i> , <i>y</i> );	
INPUT	<i>x</i>	matrix, the X axis data.
	<i>y</i>	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 **ytics** 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.

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**

S

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.

LIBRARY `pgraph`

FORMAT **scale3d**( $x,y,z$ );

INPUT  $x$  matrix, the X axis data.  
 $y$  matrix, the Y axis data.

## scalerr

---

$z$  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**  $y = \text{scalerr}(c);$

**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**.

S

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  $5 \times 5$  array, in which each element corresponds to a  $10 \times 10$  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:

## scalinfnanmiss

---

	trap 1	trap 0
function	error code	error message
<b>chol</b>	10	<b>Matrix not positive definite</b>
<b>invpd</b>	20	<b>Matrix not positive definite</b>
<b>solpd</b>	30	<b>Matrix not positive definite</b>
<b>/</b>	40	<b>Matrix not positive definite</b> <i>(second argument not square)</i>
	41	<b>Matrix singular</b> <i>(second argument is square)</i>
<b>inv</b>	50	<b>Matrix singular</b>

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.

SEE ALSO    **error, trap, trapchk**

## scalinfnanmiss

PURPOSE    Returns true if the argument is a scalar infinity, NaN, or missing value.

FORMAT     $y = \text{scalinfnanmiss}(x);$

INPUT     $x$             N×K matrix.



OUTPUT  $y$  scalar, 1 if  $x$  is a scalar, infinity, NaN, or missing value, else 0.

SEE ALSO **isinfnanmiss**, **issmiss**, **scalmiss**

## scalmiss

PURPOSE Tests to see if its argument is a scalar missing value.

FORMAT  $y = \text{scalmiss}(x);$

INPUT  $x$   $N \times K$  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.

The **issmiss** function will test each element of the matrix and return 1 if it encounters any missing values. **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 **issmiss** would return a 1 for complex  $x = . + 1i$ , and a 0 for  $x = 1 + .i$ .

EXAMPLE

```
clear s;
do until eof(fp);
    y = readr(fp,nr);
    y = packr(y);
    if scalmiss(y);
        continue;
    endif;
```

S

```

        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 2×2 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** *sch* real N×N matrix in Real Schur form, i.e., upper triangular except for possibly 2×2 blocks on the diagonal.  
*trans* real N×N matrix, the associated transformation matrix.

**OUTPUT** *schc* N×N matrix, possibly complex, strictly upper triangular. The diagonal entries are the eigenvalues.  
*transc* N×N 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

PURPOSE    Computes the Schur form of a square matrix.

FORMAT     $\{ s, z \} = \mathbf{schur}(x)$

INPUT     $x$              $K \times K$  matrix.

OUTPUT     $s$              $K \times K$  matrix, Schur form.

$z$              $K \times K$  matrix, transformation matrix.

REMARKS    **schur** computes the real Schur form of a square matrix. The real Schur form is an upper quasi-triangular matrix, that is, it is block triangular where the blocks are  $2 \times 2$  submatrices which correspond to complex eigenvalues of  $x$ . If  $x$  has no complex eigenvalues,  $s$  will be strictly upper triangular. To convert  $s$  to the complex Schur form, use the **Run-Time Library** function **schtoc**.

$x$  is first reduced to upper Hessenberg form using orthogonal similarity transformations, then reduced to Schur form through a sequence of QR decompositions.

**schur** uses the ORTRAN, ORTHES and HQR2 functions from EISPACK.

$z$  is an orthogonal matrix that transforms  $x$  into  $s$  and vice versa. Thus

$$s = z' x z$$

S

## screen

---

and since  $z$  is orthogonal,

$$x = zsz'$$

EXAMPLE    `let x[3,3] = 1 2 3  
                  4 5 6  
                  7 8 9;  
          { s, z } = schur(x);`

```
          16.11684397    4.89897949    0.00000000  
s =    -0.00000000    -1.11684397    -0.00000000  
          0.00000000    0.00000000    -0.00000000
```

```
          0.23197069    0.88290596    0.40824829  
z =    0.52532209    0.23952042    -0.81649658  
          0.81867350    -0.40386512    0.40824829
```

SEE ALSO    **hess**

## 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.

EXAMPLE     `output file = mydata.asc reset;  
screen off;  
format /m1/rz 1,8;  
open fp = mydata;  
do until eof(fp);  
    print readr(fp,200);;  
end;  
fp = close(fp);  
end;`

**S**

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**

PURPOSE     Searches the source path and (if specified) the `src` subdirectory of the **GAUSS** installation directory for a specified file.

## seekr

---

FORMAT *fpath* = **searchsourcepath**(*fname*, *srcdir*);

INPUT *fname* string, name of file to search for.

*srcdir* scalar, one of the following:

**0** do not search in the **src** subdirectory of the **GAUSS** installation directory.

**1** search the **src** subdirectory first.

**2** search the **src** subdirectory last.

OUTPUT *fpath* string, the path of *fname*, or null string if *fname* 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 *fh* scalar, file handle of an open file.

*r* scalar, the row number to which the pointer is to be moved.

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)**

S

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*            N×K matrix or string array.  
          *e*            N×1 vector of 1's and 0's.

## seqa, seqm

---

**OUTPUT**     *y*             M×K matrix or string array consisting of the rows of *x* for which there is a 1 in the corresponding row of *e*.

**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**  $y = \text{seqa}(\text{start}, \text{inc}, n);$

$y = \text{seqm}(\text{start}, \text{inc}, n);$

**INPUT** *start* scalar specifying the first element.

*inc* scalar specifying increment.

*n* scalar specifying the number of elements in the sequence.

**OUTPUT** *y*  $n \times 1$  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,

$\text{seqa}(1, 1, 10);$

will create a column vector containing the numbers 1, 2, ... 10.

For **seqm**, *y* will contain a first element equal to *start*, the second equal to *start\*inc*, and the last equal to *start\*inc<sup>n-1</sup>*.

For instance,

$\text{seqm}(10, 10, 10);$

will create a column vector containing the numbers 10, 100, ... 10<sup>10</sup>.

**EXAMPLE**  $a = \text{seqa}(2, 2, 10)';$

$m = \text{seqm}(2, 2, 10)';$

```

a  =  2 4 6 8 10 12 14 16 18 20
m  =  2 4 8 16 32 64 128 256 512 1024

```

Note that the results have been transposed in this example. Both functions return  $N \times 1$  (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>	$M \times 1$ 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  $N \times 1$  vector, then *src* must be a scalar. If *loc* is an  $[N-1] \times 1$  vector, then *src* must be a 1-dimensional array or a  $1 \times L$  vector, where L is the size of the fastest moving dimension of the array. If *loc* is an  $[N-2] \times 1$  vector, then *src* must be a  $K \times L$  matrix, or a  $K \times L$  2-dimensional array, where K is the size of the second fastest moving dimension.

Otherwise, if *loc* is an  $M \times 1$  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  $4 \times 5 \times 6$  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    **putarray**

S

setdif

PURPOSE	Returns the unique elements in one vector that are not present in a second vector.		
FORMAT	$y = \text{setdif}(v1, v2, typ);$		
INPUT	$v1$	$N \times 1$ vector.	
	$v2$	$M \times 1$ vector.	
	$typ$	scalar, type of data.	
	<b>0</b>	character, case sensitive.	
	<b>1</b>	numeric.	
	<b>2</b>	character, case insensitive.	
OUTPUT	$y$	$L \times 1$ vector containing all unique values that are in $v1$ and are not in $v2$ , sorted in ascending order.	
REMARKS	Place smaller vector first for fastest operation.		

## setdifsa

---

When there are a lot of duplicates, it is faster to remove them first with **unique** before calling this function.

EXAMPLE    `let v1 = mary jane linda john;`  
             `let v2 = mary sally;`  
             `typ = 0;`  
             `y = setdif(v1,v2,typ);`

```
           JANE
y =      JOHN
           LINDA
```

SOURCE    `setdif.src`

SEE ALSO    **setdifsa**

## 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      *sv1*          N×1 or 1×N string vector.  
             *sv2*          M×1 or 1×M string vector.

OUTPUT    *sy*          L×1 vector containing all unique values that are in *sv1* and are not in *sv2*, 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" };`

`sy = setdifsa(sv1,sv2);  
sy =jane  
     john  
     linda`

SOURCE     `setdif.src`

SEE ALSO    **setdif**

## setvars

S

PURPOSE    Reads the variable names from a data set header and creates global matrices with the same names.

FORMAT     `nvec = setvars(dataset);`

INPUT       *dataset*     string, the name of the **GAUSS** data set. Do not use a file extension.

OUTPUT      *nvec*        N×1 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**

## setwind

---

### setvwrmode

PURPOSE	Sets the graphics viewer mode.		
LIBRARY	pgraph		
FORMAT	<i>oldmode</i> = <b>setvwrmode</b> ( <i>mode</i> );		
INPUT	<i>mode</i>	string, new mode or null string.	
	<b>one</b>	Use only one viewer.	
	<b>many</b>	Use a new viewer for each graph.	
OUTPUT	<i>oldmode</i>	string, previous <i>mode</i> .	
REMARKS	If <i>mode</i> is a null string, the current <i>mode</i> will be returned with no changes made.		
	If “ <b>one</b> ” is set, the viewer executable will be <code>vwr.exe</code> .		
EXAMPLE	<pre>oldmode = setvwrmode("one"); call setvwrmode(oldmode);</pre>		
SOURCE	pgraph.src		
SEE ALSO	<b>pqgwin</b>		

### setwind

PURPOSE	Sets the current graphic panel to a previously created graphic panel number.		
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 GRAPHICS PANELS, Section 24.3.

SOURCE    `pwindow.src`

SEE ALSO    **begwind**, **endwind**, **getwind**, **nextwind**, **makewind**, **window**

S

shell

PURPOSE   Executes an operating system command.

FORMAT    **shell** [*s*];

INPUT     *s*            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).

EXAMPLE   `comstr = "ls ./src";`  
              `shell ^comstr;`

This lists the contents of the `./src` subdirectory, then returns to **GAUSS**.

## shiftr

---

```
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 = shiftr(x,s,f);`

<b>INPUT</b>	<i>x</i>	N×K matrix to be shifted.
	<i>s</i>	scalar or N×1 vector specifying the amount of shift.
	<i>f</i>	scalar or N×1 vector specifying the value to fill in.

**OUTPUT** *y* N×K 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.

**EXAMPLE** `y = shiftr(x,s,f);`



If  $x = \begin{smallmatrix} 1 & 2 \\ 3 & 4 \end{smallmatrix}$  and  $s = \begin{smallmatrix} 1 \\ -1 \end{smallmatrix}$  and  $f = \begin{smallmatrix} 99 \\ 999 \end{smallmatrix}$

Then  $y = \begin{smallmatrix} 99 & 1 \\ 4 & 999 \end{smallmatrix}$

If  $x = \begin{smallmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{smallmatrix}$  and  $s = \begin{smallmatrix} 0 \\ 1 \\ 2 \end{smallmatrix}$  and  $f = 0$

Then  $y = \begin{smallmatrix} 1 & 2 & 3 \\ 0 & 4 & 5 \\ 0 & 0 & 7 \end{smallmatrix}$

S

SEE ALSO **rotater**

PURPOSE Displays the global symbol table.

FORMAT **show** *[-flags]* *[[symbol]]*;

INPUT *flags* flags to specify the symbol type that is shown.

- k** keywords
- p** procedures
- f** **fn** functions
- m** matrices
- s** strings
- g** show only symbols with global references
- l** show only symbols with all local references

## show

---

*symbol* 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.

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

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 1×1 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*;`

## sin

---

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  $\frac{\pi}{180}$ .

**EXAMPLE**

```
let x = { 0, .5, 1, 1.5 };
y = sin(x);
```

```

          0.00000000
          0.47942554
y =       0.84147098
          0.99749499
```

SEE ALSO `atan`, `cos`, `sinh`, `pi`

## singleindex

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

**FORMAT** `si = singleindex(i,o);`

**INPUT** *i*            N×1 vector of indices into an N-dimensional array.  
*o*            N×1 vector of orders of an N-dimensional array.

**OUTPUT** *si*            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:

**S**

## sinh

---

```
ai =
      2.00000000
      1.00000000
      3.00000000
vi =      15.0000000
getarray(a,ai) =      49.0000000
v[vi] =      49.0000000
```

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     $x$              $N \times K$  matrix.

OUTPUT    $y$              $N \times K$  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);`

```
      -1.570796
      -0.785398
      0.000000
x =    0.785398
      1.570796
      3.141593
```

```
      -2.301299
      -0.868671
      0.000000
y =    0.868671
      2.301299
      11.548739
```

S

SOURCE    trig.src

sleep

PURPOSE    Sleeps for a specified number of seconds.

FORMAT    *unslept* = **sleep**(*secs*);

INPUT      *secs*           scalar, number of seconds to sleep.

OUTPUT    *unslept*       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.

## solpd

---

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**  $b$   $N \times K$  matrix or M-dimensional array where the last two dimensions are  $N \times K$ .

$A$   $N \times N$  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  $\text{inv}(A) * b$ .

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  $4 \times 2$  arrays contained in  $b$  and  $4 \times 4$  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  $\mathbf{X}$  is a matrix of independent variables, and  $\mathbf{Y}$  is a vector containing the dependent variable, then the following code will compute the least squares coefficients of the regression of  $\mathbf{Y}$  on  $\mathbf{X}$ :

```
b = solpd(X'Y,X'X);
```

EXAMPLE

```
n = 5; format /lo 16,8;
A = rndn(n,n);
A = A'A;
x = rndn(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 **scalerr, chol, invpd, trap**

**sortc, sortcc**

**PURPOSE**     Sorts a matrix of numeric or character data.

**FORMAT**      $y = \text{sortc}(x, c);$   
 $y = \text{sortcc}(x, c);$

**INPUT**        $x$              $N \times K$  matrix.  
 $c$             scalar specifying one column of  $x$  to sort on.

**OUTPUT**       $y$              $N \times K$  matrix equal to  $x$  and sorted on the column  $c$ .

**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  $-\infty$ .

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:

$\text{rev}(\text{sortc}(x, c))$

**EXAMPLE**      $\text{let } x[3,3] = \begin{matrix} 4 & 7 & 3 \\ 1 & 3 & 2 \\ 3 & 4 & 8; \end{matrix}$   
 $y = \text{sortc}(x, 1);$

```

      4  7  3
x =   1  3  2
      3  4  8

```

```

      1  3  2
y =   3  4  8
      4  7  3

```

SEE ALSO **rev**

## sortd

S

**PURPOSE** Sorts a data file on disk with respect to a specified variable.

**FORMAT** **sortd**(*infile*,*outfile*,*keyvar*,*keytyp*);

**INPUT** *infile* string, name of input file.  
*outfile* string, name of output file, must be different.  
*keyvar* string, name of key variable.  
*keytyp* scalar, 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 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

## sorthc, sorthcc

---

SEE ALSO `sortmc`, `sortc`, `sortcc`, `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** `x` `N`×`K` matrix or string array.  
`c` scalar specifying one column of `x` to sort on.

**OUTPUT** `y` `N`×`K` matrix or string array equal to `x` and sorted on the column `c`.

**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;
y = sorthc(x,1);

```

```

      4 7 3
x =  1 3 2
      3 4 8

```

```

      1 3 2
y =  3 4 8
      4 7 3

```

SEE ALSO **sortc, rev**

S

## sortind, sortindc

**PURPOSE** Returns the sorted index of  $x$ .

**FORMAT**  $ind = \text{sortind}(x);$   
 $ind = \text{sortindc}(x);$

**INPUT**  $x$   $N \times 1$  column vector.

**OUTPUT**  $ind$   $N \times 1$  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.

## sortmc

---

EXAMPLE    `let x = 3 8 2 5 1 6 9`  
             `ind = sortind(x);`  
             `y = x[ind];`

             3  
             8  
             2  
`x =` 5  
             1  
             6  
             9

             5  
             3  
             1  
`ind =` 4  
             6  
             2  
             7

             1  
             2  
             3  
`y =` 5  
             6  
             8  
             9

## sortmc

PURPOSE    Sorts a matrix on multiple columns.

FORMAT     $y = \text{sortmc}(x, v);$

INPUT     $x$              $N \times K$  matrix to be sorted.  
            $v$              $L \times 1$  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    $y$              $N \times K$  sorted matrix.

SOURCE   `sortmc.src`

SEE ALSO   **sortd, sortc, sortcc, sorthc, sorthcc**

**S**

**sortr, sortrc**

PURPOSE   Sorts rows of a matrix of numeric or character data.

FORMAT     $y = \text{sortr}(x, r);$   
               $y = \text{sortrc}(x, r);$

INPUT     $x$              $N \times K$  matrix.  
            $r$             scalar, row of  $x$  on which to sort.

OUTPUT    $y$              $N \times K$  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    `let x = { 4 7 3,`  
                  `1 3 2,`  
                  `3 4 8 };`

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

```
      3 4 7  
y = 2 1 1  
      8 3 3
```

## 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);`

<b>INPUT</b>	<i>r</i>	scalar, rows of output matrix.
	<i>c</i>	scalar, columns of output matrix.
	<i>vals</i>	N×1 vector, non-zero values.
	<i>rinds</i>	N×1 vector, row indices of corresponding non-zero values.
	<i>cinds</i>	N×1 vector, column indices of corresponding non-zero values.



OUTPUT

$y$

$r \times c$  sparse matrix.

REMARKS

Since sparse matrices are strongly typed in **GAUSS**,  $y$  must be defined as a sparse matrix before the call to **spCreate**.

EXAMPLE

```

sparse matrix y;
vals = { 1,2,3,4 };
rinds = { 2,5,8,13 };
cinds = { 4,1,9,5 };

y = spCreate(15,10,vals,rinds,cinds);

```

This example creates a  $15 \times 10$  sparse matrix **y**, containing the following non-zero values:

Non-zero value	Index
1	(2,4)
2	(5,1)
3	(8,9)
4	(13,5)

SEE ALSO

**packedToSp**, **denseToSp**, **spEye**

spDenseSubmat

PURPOSE

Returns a dense submatrix of a sparse matrix.

FORMAT

$y = \text{spDenseSubmat}(x, rinds, cinds);$

INPUT

$x$

$M \times N$  sparse matrix.

$rinds$

$K \times 1$  vector, row indices.

$cinds$

$L \times 1$  vector, column indices.

OUTPUT

$y$

$K \times L$  dense matrix, the intersection of  $rinds$  and  $cinds$ .

## spDiagRvMat

---

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 };`  
  
             `y = denseToSp(x,0);`  
             `d = spDenseSubmat(y,0,1|3|4);`

```
      0 0 10
      0 0 0
d =   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	<i>x</i>	M×N sparse matrix.
	<i>inds</i>	K×2 vector or scalar 0, row and column indices into <i>x</i> at which to place the corresponding submatrices in <i>a</i> .
	<i>size</i>	K×2 vector or scalar 0, sizes of the corresponding submatrices in <i>a</i> .
	<i>a</i>	K×L×P array, containing the submatrices to insert into <i>x</i> .

OUTPUT      $y$               $M \times N$  sparse matrix, a copy of  $x$  containing the specified insertions.

REMARKS     Each row of  $ind$  must contain the row and column indices, respectively, that form the starting point for the insertion of the corresponding submatrix in  $a$ . If  $ind$  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;  
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 };  
a = arrayinit(4|3|4,0);  
a[1,1:2,1:2] = sx1;  
a[2,..] = sx2;  
a[3,1:2,1:3] = sx3;  
a[4,1:2,1] = sx4;  
inds = 0;  
siz = { 2 2, 3 4, 2 3, 2 1 };  
y = spDiagRvMat(x,inds,siz,a);  
  
dx = spToDense(x);  
dy = spToDense(y);`

```

      1  0  0  0  0  0  0  0  0  0
      0  1  0  0  0  0  0  0  0  0
      0  0  1  0  0  0  0  0  0  0
      0  0  0  1  0  0  0  0  0  0
dx =  0  0  0  0  1  0  0  0  0  0
      0  0  0  0  0  1  0  0  0  0
      0  0  0  0  0  0  1  0  0  0
      0  0  0  0  0  0  0  1  0  0
      0  0  0  0  0  0  0  0  1  0
      0  0  0  0  0  0  0  0  0  1

```

```

      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
dy =  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

```

## spEye

**PURPOSE**    Creates a sparse identity matrix.

**FORMAT**     $y = \text{spEye}(n);$

**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** `sparse matrix y;  
y = spEye(3);  
d = spDenseSubmat(y,0,0);`

```

          1.0000000  0.0000000  0.0000000
d = 0.0000000  1.0000000  0.0000000
      0.0000000  0.0000000  1.0000000

```

**SEE ALSO** `spCreate`, `spOnes`, `denseToSp`

S

## 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** *x* M×N sparse matrix.

**OUTPUT** *vals* N×1 vector, non-zero values in *x*.  
*rinds* N×1 vector, row indices of corresponding non-zero values.  
*cinds* N×1 vector, column indices of corresponding non-zero values.

**EXAMPLE** `sparse matrix y;  
x = { 0 0 0 10,  
 0 2 0 0,  
 0 0 0 0,  
 5 0 0 0,`

## spline

---

```
0 0 0 3 };  
  
y = denseToSp(x,0);  
{ v,r,c } = spGetNZE(y);
```

```
10  
2  
v = 5  
3
```

```
1  
2  
r = 4  
5
```

```
4  
2  
c = 1  
4
```

SEE ALSO **spNumNZE**

## spline

**PURPOSE** Computes a two-dimensional interpolatory spline.

**FORMAT** { *u,v,w* } = **spline**(*x,y,z,sigma,g*);

**INPUT** *x*            1×K vector, x-abcissae (x-axis values).  
         *y*            N×1 vector, y-abcissae (y-axis values).

	<i>z</i>	K×N matrix, ordinates (z-axis values).
	<i>sigma</i>	scalar, tension factor.
	<i>g</i>	scalar, grid size factor.
OUTPUT	<i>u</i>	1×(K*g) vector, x-abcissae, regularly spaced.
	<i>v</i>	(N*g)×1 vector, y-abcissae, regularly spaced.
	<i>w</i>	(K*g)×(N*g) matrix, interpolated ordinates.
REMARKS	<p><i>sigma</i> contains the tension factor. This value indicates the curviness desired. If <i>sigma</i> is nearly zero (e.g., .001), the resulting surface is approximately the tensor product of cubic splines. If <i>sigma</i> is large (e.g., 50.0), the resulting surface is approximately bi-linear. If <i>sigma</i> equals zero, tensor products of cubic splines result. A standard value for <i>sigma</i> is approximately 1.</p> <p><i>g</i> is the grid size factor. It determines the fineness of the output grid. For <i>g</i> = 1, the output matrices are identical to the input matrices. For <i>g</i> = 2, the output grid is twice as fine as the input grid, i.e., <i>u</i> will have twice as many columns as <i>x</i>, <i>v</i> will have twice as many rows as <i>y</i>, and <i>w</i> will have twice as many rows and columns as <i>z</i>.</p>	
SOURCE	spline.src	

S

spNumNZE

PURPOSE	Returns the number of non-zero elements in a sparse matrix.	
FORMAT	<i>n</i> = <b>spNumNZE</b> ( <i>x</i> ) ;	
INPUT	<i>x</i>	M×N sparse matrix.
OUTPUT	<i>n</i>	scalar, the number of non-zero elements in <i>x</i> .

## spOnes

---

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);

            n = 4

SEE ALSO    **spGetNZE**

## spOnes

PURPOSE    Generates a sparse matrix containing only ones and zeros

FORMAT    **y = spOnes(r,c,rinds,cinds);**

INPUT      *r*            scalar, rows of output matrix.  
            *c*            scalar, columns of output matrix.  
            *rinds*        N×1 vector, row indices of ones.  
            *cinds*        N×1 vector, column indices of ones.

OUTPUT    *y*            *r*×*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    sparse matrix y;



```
rinds = { 1, 3, 5 };
cinds = { 2, 1, 3 };
y = spOnes(5,4,rinds,cinds);
d = spDenseSubmat(y,0,0);
```

```

          0.0000000  1.0000000  0.0000000  0.0000000
          0.0000000  0.0000000  0.0000000  0.0000000
d = 1.0000000  0.0000000  0.0000000  0.0000000
          0.0000000  0.0000000  0.0000000  0.0000000
          0.0000000  0.0000000  1.0000000  0.0000000
```

SEE ALSO    **spCreate, spEye, spZeros, denseToSp**

S

SpreadsheetReadM

PURPOSE    Reads and writes Excel files.

FORMAT    *xlsmat* = **SpreadsheetReadM**(*file*,*range*,*sheet*);

INPUT    *file*            string, name of .xls file.  
          *range*        string, *range* to read or write; e.g., “a1:b20”.  
          *sheet*        scalar, *sheet* number.

OUTPUT    *xlsmat*        matrix of numbers read from Excel.

PORTABILITY    **Windows** only

REMARKS    If the read functions fail, they will return a scalar error code which can be decoded with **scalerr**. If the write function fails, it returns a non-zero error number.

SEE ALSO    **scalerr, error**

## SpreadsheetReadSA

---

### SpreadsheetReadSA

PURPOSE Reads and writes Excel files.

FORMAT *xlssa* = **SpreadsheetReadSA**(*file*,*range*,*sheet*);

INPUT *file* string, name of .xls *file*.  
*range* string, *range* to read or write; e.g., “a1:b20”.  
*sheet* scalar, *sheet* number.

OUTPUT *xlssa* string array read from Excel.

PORTABILITY **Windows** only

REMARKS If the read functions fail, they will return a scalar error code which can be decoded with **scalerr**. If the write function fails, it returns a non-zero error number.

SEE ALSO **scalerr**, **error**

### SpreadsheetWrite

PURPOSE Reads and writes Excel files.

FORMAT *xlsret* = **SpreadsheetWrite**(*data*,*file*,*range*,*sheet*);

INPUT *data* matrix, string or string array, *data* to write.  
*file* string, name of .xls file.  
*range* string, *range* to read or write; e.g., “a1:b20”.

	<i>sheet</i>	scalar, <i>sheet</i> number.
OUTPUT	<i>xlsret</i>	success code, 0 if successful, else error code.
PORTABILITY	<b>Windows</b> only	
REMARKS	If the read functions fail, they will return a scalar error code which can be decoded with <b>scalerr</b> . If the write function fails, it returns a non-zero error number.	
SEE ALSO	<b>scalerr</b> , <b>error</b>	

spScale

S

PURPOSE	Scales a sparse matrix.	
FORMAT	{ <i>a r s</i> } = <b>spScale</b> ( <i>x</i> );	
INPUT	<i>x</i>	M×N sparse matrix.
OUTPUT	<i>a</i>	M×N scaled sparse matrix.
	<i>r</i>	M×1 vector, row scale factors.
	<i>s</i>	N×1 vector, column scale factors.
REMARKS	<b>spScale</b> scales the elements of the matrix by powers of 10 so that they are all within (-10,10).	
EXAMPLE	<pre>x = { 25  -12   0,       3    0 -11,       8 -100   0 };</pre> <pre>declare sparse matrix sm, smsc;</pre>	

## spSubmat

---

```
sm = denseToSp(x,0);  
  
{ smsc, r, c } = spScale(sm);
```

```
      25  -12   0  
sm =   3   0 -11  
      8 -100   0
```

```
      2.5 -1.2  0.0  
smsc =   0.3  0.0 -1.1  
      0.08 -1.0  0.0
```

```
      0.1  
r =   0.1  
      0.01
```

```
      1.0  
c =   1.0  
      1.0
```

## spSubmat

**PURPOSE** Returns a sparse submatrix of a sparse matrix.

**FORMAT** `y = spSubmat(x,rinds,cinds);`

<b>INPUT</b>	<i>x</i>	M×N sparse matrix.
	<i>rinds</i>	K×1 vector, row indices.
	<i>cinds</i>	L×1 vector, column indices.

OUTPUT  $s$   $K \times L$  sparse matrix, the intersection of *rinds* and *cinds*.

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);
z = spSubmat(y,1|3|4,0);
d = spDenseSubmat(z,0,0);

```

```

      0  0  0 10
d =  0  0  0  0
      5  0  0  0

```

SEE ALSO **spDenseSubmat**

## spToDense

PURPOSE Converts a sparse matrix to a dense matrix.

FORMAT  $y = \text{spToDense}(x);$

INPUT  $x$   $M \times N$  sparse matrix.

## spTrTDense

---

OUTPUT     $y$              $M \times N$  dense matrix.

REMARKS    A dense matrix is just a normal format matrix.

EXAMPLE    `sparse matrix y;`  
              `y = spEye(4);`  
              `d = spToDense(y);`

$$d = \begin{pmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix}$$

SEE ALSO    `spDenseSubmat`, `denseToSp`

## spTrTDense

PURPOSE    Multiplies a sparse matrix transposed by a dense matrix.

FORMAT    `y = spTrTDense(s,d);`

INPUT       $s$              $N \times M$  sparse matrix.

$d$              $N \times L$  dense matrix.

OUTPUT     $y$              $M \times L$  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**  $y = \text{spTScalar}(s, \text{scal}, \text{rinds}, \text{cinds});$

**INPUT**

$s$	$N \times M$ sparse matrix.
$\text{scal}$	scalar.
$\text{rinds}$	$K \times 1$ vector of row indices.
$\text{cinds}$	$L \times 1$ vector of column indices.

**OUTPUT**  $y$   $K \times L$  sparse matrix.

**REMARKS** Only the elements of  $s$  specified by  $\text{rinds}$  and  $\text{cinds}$  will be multiplied by  $\text{scal}$ . All other elements will be unchanged in the result.

To select all rows or all columns, input a scalar 0 for  $\text{rinds}$  or  $\text{cinds}$ .

Since sparse matrices are strongly typed in **GAUSS**,  $y$  must be defined as a sparse matrix before the call to **spTScalar**.

**EXAMPLE**

```

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 };
y = spTScalar(x,10,rinds,cinds);
d = spDenseSubmat(y,0,0);

```

**S**

## spZeros

---

```
      3   0   2  10
d =   0  40   0   0
      5   0   0  30
      0  10   2   0
```

SEE ALSO **spTrTDense**

## spZeros

**PURPOSE** Creates a sparse matrix containing no non-zero values.

**FORMAT** `y = spZeros(r,c);`

**INPUT** *r* scalar, rows of output matrix.  
*c* scalar, columns of output matrix.

**OUTPUT** *y*  $r \times c$  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*;  
`y = spZeros(4,3);`  
`d = spDenseSubmat(y,0,0);`

```
      0.0000000  0.0000000  0.0000000
d =   0.0000000  0.0000000  0.0000000
      0.0000000  0.0000000  0.0000000
      0.0000000  0.0000000  0.0000000
```

SEE ALSO **spOnes, spEye, createSp**



**PURPOSE**     Solves the nonlinear programming problem using a sequential quadratic programming method.

**FORMAT**     { *x,f,lagr,retcode* } = **sqpSolve**(&*fct*,*start*);

**INPUT**     *&fct*     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.

*start*     K×1 vector of start values.

**GLOBAL INPUT**     **\_sqp\_A**     M×K matrix, linear equality constraint coefficients.

**\_sqp\_B**     M×1 vector, linear equality constraint constants.

                     These globals are used to specify linear equality constraints of the following type:

$$\_sqp\_A * X = \_sqp\_B$$

where *X* is the K×1 unknown parameter vector.

**\_sqp\_EqProc**     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×1 vector of parameters, and one output argument, the R×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:

$$\text{\_sqp\_C} * X \geq \text{\_sqp\_D}$$

where  $X$  is the  $K \times 1$  unknown parameter vector.

**\_sqp\_IneqProc** scalar, pointer to a procedure that computes the 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  $K \times 1$  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:

$$p[1] * p[2] \geq p[3]$$

The procedure for this is:

```
proc ineqproc(p);
    retp(p[1]*[2]-p[3]);
endp;
```

**\_sqp\_Bounds**  $K \times 2$  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  $1 \times 2$  matrix may be used. Default is:

$$\begin{bmatrix} [1] & -1e256 & [2] & 1e256 \end{bmatrix}$$

**\_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;
```

tells **sqpSolve** that a gradient procedure exists and where to find it. The user-provided procedure has two input arguments, a  $K \times 1$  vector of parameter values and an  $N \times P$  matrix of data. The procedure returns a single output argument, an  $N \times K$  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.

**\_sqp\_HessProc** 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  $P \times 1$  vector of parameter values and an  $N \times K$  data matrix. The procedure returns a single output argument, the  $P \times P$  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\_ParNames**  $K \times 1$  character vector, parameter names.

**\_sqp\_PrintIters** scalar, if nonzero, prints iteration information. Default = 0. Can be toggled during iterations by pressing P on the keyboard.

**\_sqp\_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.

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**\_sqp\_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.

**\_\_output** scalar, if nonzero, results are printed. Default = 0.

OUTPUT *x*  $K \times 1$  vector of parameters at minimum.

*f* scalar, function evaluated at *x*.

*lagr* vector, created using **vput**. Contains the Lagrangean for the constraints. They may be extracted with the **vread** command using the following strings:

‘‘lineq’’ Lagrangeans of linear equality constraints,  
 ‘‘nlineq’’ 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.

*retcode* 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

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    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.

INCLUDE sqpsolvemt.sdf

FORMAT *out1* = **sqpSolveMT**(&*fct*,*par1*,*data1*,*c1*);

INPUT &*fct* pointer to a procedure that computes the function to be minimized.

This procedure must have two input arguments, an instance of structure of type **PV** and an instance of a structure of type **DS**, and one output argument, either a 1×1 scalar or an N×1 vector of function values evaluated at the parameters stored in the **PV** instance using data stored in the **DS** instance.

*par1* an instance of structure of type **PV**. The *par1* instance is passed to the user-provided procedure pointed to by *&fct*. *par1* is constructed using the “**pack**” functions.

*data1* an array of instances of a **DS** structure. This array is passed to the user-provided pointed by *&fct* to be used in the objective function. **sqpSolveMT** does not look at this structure. Each instance contains the the following members which can be set in whatever way that is convenient for computing the objective function:

*data1*[*i*].*dataMatrix* N×K matrix, data matrix.

*data1*[*i*].*dataArray* N×K×L.. array, data array.

*data1*[*i*].*vnames* string array, variable names (optional).

*data1*[*i*].*dsname* string, data name (optional).

*data1*[*i*].*type* scalar, type of data (optional).

*c1* 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 *c1*, the members are:

*c1.A* M×K matrix, linear equality constraint coefficients:  $c1.A * p = c1.B$  where *p* is a vector of the parameters.

*c1.B* M×1 vector, linear equality constraint constants:  $c1.A * p = c1.B$  where *p* is a vector of the parameters.

*c1.C* M×K matrix, linear inequality constraint coefficients:  $c1.C * p \geq c1.D$  where *p* is a vector of the parameters.

*c1.D* M×1 vector, linear inequality constraint constants:  $c1.C * p \geq c1.D$  where *p* is a vector of the parameters.

<i>cl.eqProc</i>	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 <b>SQPdata</b> , and one output argument, a vector of computed equality constraints. For more details see Remarks below. Default = ., i.e., no equality procedure.
<i>cl.weights</i>	vector, weights for objective function returning a vector. Default = 1.
<i>cl.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 <b>SQPdata</b> , and one output argument, a vector of computed inequality constraints. For more details see Remarks below. Default = ., i.e., no inequality procedure.
<i>cl.bounds</i>	1×2 or K×2 matrix, bounds on parameters. If 1×2 all parameters have same bounds. Default = -1e256 1e256 .
<i>cl.covType</i>	scalar, if 2, QML covariance matrix, else if 0, no covariance matrix is computed, else ML covariance matrix is computed.
<i>cl.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>cl.hessProc</i>	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. Default = ., i.e., no Hessian procedure has been provided.
<i>cl.maxIters</i>	scalar, maximum number of iterations. Default = 1e+5.
<i>cl.dirTol</i>	scalar, convergence tolerance for gradient of

estimated coefficients. Default = 1e-5. When this criterion has been satisfied **SQPSolve** exits the iterations.

- cl.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.
- cl.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.
- cl.output* scalar, if nonzero, results are printed. Default = 0.
- cl.printIters* scalar, if nonzero, prints iteration information. Default = 0.

OUTPUT     *out1*     an instance of an **sqpSolveMTout** structure. For an instance named *out1*, the members are:

- out1.par*     an instance of structure of type **PV** containing the parameter estimates will be placed in the member matrix *out1.par*.
- out1.fct*     scalar, function evaluated at *x*.
- out1.lagr*     an instance of a **SQPLagrange** structure containing the Lagrangeans for the constraints. The members are:
- out1.lagr.lineq*     M×1 vector,  
Lagrangeans of linear equality constraints.
- out1.lagr.nlineq*     N×1 vector,  
Lagrangeans of nonlinear equality constraints.
- out1.lagr.linineq*     P×1 vector,  
Lagrangeans of linear inequality constraints.



*out1.lagr.nlineq*  $Q \times 1$  vector,  
Lagrangeans of  
nonlinear inequality  
constraints.

*out1.lagr.bounds*  $K \times 2$  matrix,  
Lagrangeans of  
bounds.

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.

*out1.retcode*

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.

**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 have one input argument that is an instance of a structure of type struct **PV** and a second argument that is an instance of a structure of type struct **DS**. On input to the call to **sqpSolveMT**, the first argument contains starting values for the parameters and the second argument any required data. The data are passed in a separate argument because the structure in the first argument will be copied as it is passed through procedure calls which would be very costly if it contained large data matrices. Since **sqpSolveMT** makes no

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changes to the second argument it will be passed by pointer thus saving time because its contents aren't copied.

Both of the structures of type **PV** are set up using the **PV** “**pack**” procedures, **pvPack**, **pvpPackm**, **pvpPacks**, and **pvpPacksm**. 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 Micherlitz(struct PV par1, struct DS data1);
    local p0,e,s2,x,y;
    p0 = pvUnpack(par1,"parameters");
    y = data1.dataMatrix[.,1];
    x = data1.dataMatrix[.,2];
    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 first and second columns of a data matrix stored in a single **DS** structure. Alternatively these two columns of data can be entered into a vector of **DS** structures, one for each column of data:

```
proc Micherlitz(struct PV par1, struct DS data1);
    local p0,e,s2,x,y;
    p0 = pvUnpack(par1,"parameters");
    y = data1[1].dataMatrix;
    x = data1[2].dataMatrix;
    e = y - p0[1] - p0[2]*exp(-p0[3]*x);
    retp(e'*e);
endp;
```

The syntax is similar for the optional user-provided procedures. 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 ineqConst(struct PV par1, struct DS data1);
    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:

```
#include sqpSolveMT.sdf

struct DS d0;
d0 = dsCreate;

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);
d0.dataMatrix = y~x;
struct sqpSolveMTControl c0;
c0 = sqpSolveMTControlCreate;
```

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```
c0.bounds = 0~100; /* constrains parameters */
                /* to be positive */

struct PV par1;
par1 = pvCreate;
pvPack(par1,.92|2.62|.114,"parameters");
struct sqpSolveMTout out1;
out1 = sqpSolveMT(&Micherlitz,par1,d0,c0);

print " parameter estimates ";
print pvUnPack(out1.par,"parameters");

proc Micherlitz(struct PV par1, struct DS data1);
    local p0,e,s2,x,y;
    p0 = pvUnpack(par1,"parameters");
    y = data1.dataMatrix[.,1];
    x = data1.dataMatrix[.,2];
    e = y - p0[1] - p0[2]*exp(-p0[3] * x);
    retp(e'*e);
endp;

proc grad(struct PV par1, struct DS data1);
    local p0,e,w,g,r,x,y;
    p0 = pvUnpack(par1,"parameters");
    y = data1.dataMatrix[.,1];
    x = data1.dataMatrix[.,2];
    g = zeros(3,1);
    w = exp(-p0[3] * x);
    e = y - p0[1] - p0[2]*w;
    r = e'*w;
    g[1] = -2*sumc(e);
    g[2] = -2*r;
    g[3] = 2*p0[1]*p0[2]*r;
    retp(g);
endp;
```

SOURCE    sqpsolvemt.src

SEE ALSO    **sqpSolveMTControlCreate**, **sqpSolveMTlagrangeCreate**,  
**sqpSolveOutCreate**

sqpSolveMTControlCreate

PURPOSE    Creates an instance of a structure of type **sqpSolveMTcontrol** set to default values.

INCLUDE    sqpsolvemt.sdf

FORMAT    *s* = **sqpSolveMTControlCreate**;

OUTPUT    *s*            instance of structure of type **sqpSolveMTControl**.

SOURCE    sqpsolvemt.src

SEE ALSO    **sqpSolve**

S

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**.

## **sqpSolveMToutCreate**

---

SOURCE    `sqpsolvemt.src`

SEE ALSO    **sqpSolve**

### **sqpSolveMToutCreate**

PURPOSE    Creates an instance of a structure of type **sqpSolveMTout** set to default values.

INCLUDE    `sqpsolvemt.sdf`

FORMAT    `s = sqpSolveMToutCreate;`

OUTPUT    `s`            instance of structure of type **sqpSolveMTout**.

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        $x$                  $N \times K$  matrix or  $N$ -dimensional array.

OUTPUT       $y$                  $N \times K$  matrix or  $N$ -dimensional array, the square roots of each element of  $x$ .

S

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.

EXAMPLE      $\text{let } x[2,2] = 1 \ 2 \ 3 \ 4;$   
                  $y = \text{sqrt}(x);$

$x =$         1.00000000    2.00000000  
              3.00000000    4.00000000

$y =$         1.00000000    1.41421356  
              1.73205081    2.00000000

**stdc**

**PURPOSE**     Computes the standard deviation of the elements in each column of a matrix.

**FORMAT**      $y = \text{stdc}(x);$

**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:

$$\text{sqrt}(1/(N-1) * \text{sumc}((x - \text{meanc}(x))' * (x - \text{meanc}(x))))$$

Thus, the divisor is  $N-1$  rather than  $N$ , where  $N$  is the number of elements being summed. To convert to the alternate definition, multiply by

$$\text{sqrt}((N-1)/N)$$

**EXAMPLE**      $y = \text{rndn}(8100, 1);$   
                   $\text{std} = \text{stdc}(y);$

$$\text{std} = 1.008377$$

In this example, 8100 standard Normal random variables are generated, and their standard deviation is computed.

**SEE ALSO**     **meanc**



## stdsc

**PURPOSE** Computes the standard deviation of the elements in each column of a matrix.

**FORMAT**  $y = \text{stdsc}(x);$

**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:

$$\text{sqrt}(1/(N) * \text{sumc}((x - \text{meanc}(x))' )^2))$$

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**  $y = \text{rndn}(8100, 1);$   
 $\text{std} = \text{stdsc}(y);$

$\text{std} = 1.0151475$

In this example, 8100 standard Normal random variables are generated, and their standard deviation is computed.

**SEE ALSO** **stdc**, **astds**, **meanc**

S

## stof

---

### stocv

**PURPOSE**     Converts a string to a character vector.

**FORMAT**      $v = \text{stocv}(s);$

**INPUT**         $s$                 string, to be converted to character vector.

**OUTPUT**       $v$                 N×1 character vector, contains the contents of  $s$ .

**REMARKS**     **stocv** breaks  $s$  up into a vector of 8-character length matrix elements. Note that the character information in the vector is not guaranteed to be null-terminated.

**EXAMPLE**      $s = \text{"Now is the time for all good men"};$   
                   $v = \text{stocv}(s);$

```
          "Now is t"  
v =      "he time "  
          "for all "  
          "good men"
```

**SEE ALSO**     **cvtos, vget, vlist, vput, vread**

### stof

**PURPOSE**     Converts a string to floating point.

**FORMAT**      $y = \text{stof}(x);$

**INPUT**         $x$                 string or N×K 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 3×1 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

s

**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.

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 N×M string array to an N×1 string vector by combining each element in a column separated by a user-defined delimiter string.

## strindx

---

FORMAT     $y = \text{strcombine}(sa, delim, qchar);$

INPUT    *sa*             $N \times M$  string array.  
          *delim*         $1 \times 1$ ,  $1 \times M$ , or  $M \times 1$  delimiter string.  
          *qchar*       scalar,  $2 \times 1$ , or  $1 \times 2$  string vector containing quote characters as required:  
                          scalar:                            Use this character as quote character.  
  If this is 0, no quotes are added.  
                           $2 \times 1$  or  $1 \times 2$  string vector:    Contains left and right quote characters.

OUTPUT     $y$              $N \times 1$  string vector result.

SOURCE    `strfns.src`

SEE ALSO    **satostrC**

## strindx

PURPOSE    Finds the index of one string within another string.

FORMAT     $y = \text{strindx}(where, what, start);$

INPUT    *where*        string or scalar, the data to be searched.  
          *what*        string or scalar, the substring to be searched for in *where*.  
          *start*       scalar, the starting point of the search in *where* for an occurrence of *what*. The index of the first character in a string is 1.

OUTPUT     $y$             scalar containing the index of the first occurrence of *what*, within *where*, which is greater than or equal to *start*. 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);

y = 8
```

This function is used with **strsect** for extracting substrings.

SEE ALSO    **strrindx, strlen, strsect, strput**

strlen

s

PURPOSE	Returns the length of a string.		
FORMAT	$y = \text{strlen}(x);$		
INPUT	$x$	string, N×K matrix of character data, or N×K string array.	
OUTPUT	$y$	scalar containing the exact length of the string $x$ , or N×K matrix or string array containing the lengths of the elements in $x$ .	
REMARKS	<p>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.</p> <p>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.</p>		
EXAMPLE	$x = \text{"How long?"};$ $y = \text{strlen}(x);$		

## strput

---

$y = 9$

SEE ALSO **strsect, strindx, strrindx**

## strput

PURPOSE Lays a substring over a string.

FORMAT  $y = \mathbf{strput}(\mathit{substr}, \mathit{str}, \mathit{off})$ ;

INPUT *substr* string, the substring to be laid over the other string.  
*str* string, the string to receive the substring.  
*off* scalar, the offset in *str* to place *substr*. The offset of the first byte is 1.

OUTPUT  $y$  string, the new string.

EXAMPLE 

```
str = "max";  
sub = "imum";  
f = 4;  
y = strput(sub, str, f);  
print y;
```

produces:

`maximum`

SOURCE `strput.src`

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** `y` scalar containing the index of the last occurrence of *what*, within *where*, which is less than or equal to *start*. 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;

pos = 11

name = "ols.src"
```

S

## strsect

---

**strrindx** can be used with **strsect** for extracting substrings.

SEE ALSO **strindx, strlen, strsect, strput**

### strsect

**PURPOSE** Extracts a substring of a string.

**FORMAT** `y = strsect(str,start,len);`

<b>INPUT</b>	<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.

<b>OUTPUT</b>	<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);`

`y = "example"`

SEE ALSO **strlen, strindx, strrindx**



strsplit

PURPOSE     Splits an N×1 string vector into an N×K string array of the individual tokens.

FORMAT     *sa* = **strsplit**(*sv*);

INPUT       *sv*            N×1 string array.

OUTPUT      *sa*            N×K string array.

REMARKS     Each row of *sv* must contain the same number of tokens. The following characters are considered delimiters between tokens:

- space            ASCII 32
- tab              ASCII 9
- comma           ASCII 44
- newline          ASCII 10
- carriage return ASCII 13

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     

```
let string sv = {
    "dog 'cat fish' moose",
    "lion, zebra, elk",
    "seal owl whale"
};

sa = strsplit(sv);
```

S

## strsplitPad

---

```
          'dog'  'cat fish' 'moose'  
sa =      'lion' 'zebra'   'elk'  
          'seal' 'owl'    'whale'
```

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*,*cols*);

**INPUT**    *sv*            N×1 string array.  
          *cols*          scalar, number of columns of output string array.

**OUTPUT**   *sa*            N×*cols* string array.

**REMARKS** Rows containing more than *cols* tokens are truncated and rows containing fewer than *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

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  let string sv = {
           "dog 'cat fish' moose",
           "lion, zebra, elk, bird",
           "seal owl whale"
         };

           sa = strsplitPad(sv, 4);

           'dog'  'cat fish'  'moose'  ""
sa =      'lion'  'zebra'    'elk'     'bird'
           'seal' 'owl'     'whale'   ""
```

SEE ALSO    **strsplit**

S

strtodt

**PURPOSE**    Converts a string array of dates to a matrix in DT scalar format.

**FORMAT**     $x = \text{strtodt}(sa,fmt);$

**INPUT**      *sa*            N×K string array containing dates.  
              *fmt*           string containing date/time format characters.

**OUTPUT**     *x*            N×K 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  
  
              20050910223505  
  
              represents 22:35:05 or 10:35:05 PM on September 10, 2005.  
  
              The following formats are supported:

## strtodt

---

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("2005-07-12 10:18:32",  
              "YYYY-MO-DD HH:MI:SS");  
              print x;`

produces:

`20050712101832.0`

`x = strtodt("2005-07-12 10:18:32", "YYYY-MO-DD");  
print x;`

produces:

`200507120000000.0`

`x = strtodt("10:18:32", "HH:MI:SS");  
print x;`

produces:

`101832.0`

`x = strtodt("05-28-05", "MO-DD-YR");  
print x;`

produces:

20050528000000.0

SEE ALSO **dttostr**, **dttoutc**, **utctodt**

strtof

S

- PURPOSE    Converts a string array to a numeric matrix.
- FORMAT     $x = \text{strtof}(sa);$
- INPUT       $sa$             N×K string array containing numeric data.
- OUTPUT     $x$              N×K matrix.
- REMARKS    This function supports real matrices only. Use **strtofcplx** for complex data.
- SEE ALSO    **strtofcplx**, **ftostrC**

strtofcplx

- PURPOSE    Converts a string array to a complex numeric matrix.
- FORMAT     $x = \text{strtofcplx}(sa);$
- INPUT       $sa$             N×K string array containing numeric data.
- OUTPUT     $x$              N×K complex matrix.

## strtriml

---

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**, **ftostrC**

## strtriml

PURPOSE     Strips all whitespace characters from the left side of each element in a string array.

FORMAT      $y = \text{strtriml}(sa);$

INPUT        $sa$              $N \times M$  string array.

OUTPUT      $y$              $N \times M$  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      $y = \text{strtrimr}(sa);$

INPUT        $sa$              $N \times M$  string array.

OUTPUT     *y*             N×M 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       *sa*             N×K string array.  
              *maxlen*       1×K or 1×1 matrix, maximum length.

OUTPUT     *y*             N×K string array result.

SEE ALSO     `strtriml`, `strtrimr`, `strtrunc1`, `strtruncpad`, `strtruncr`

S

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       *sa*             N×M, N×1, 1×M, or 1×1 string array.  
              *ntrunc*       N×M, N×1, 1×M, or 1×1 matrix containing the number of characters to strip.

## strtruncpad

---

OUTPUT     *y*                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* = **strtruncpad**(*sa*,*maxlen*);

INPUT        *sa*                N×K string array.  
              *maxlen*        1×K or 1×1 matrix, maximum length.

OUTPUT     *y*                N×K string array result.

SEE ALSO     **strtriml, strtrimr, strtrunc, strtruncr, 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        *sa*                N×M, N×1, 1×M, or 1×1 string array.  
              *ntrunc*        N×M, N×1, 1×M, or 1×1 matrix containing the number of characters to strip.



OUTPUT     *y*             String array result.

SOURCE     `strfns.src`

SEE ALSO     `strtriml`, `strtrimr`, `strtrunc`, `strtrunc1`, `strtruncpad`

submat

S

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*             N×K matrix.  
              *r*             L×M matrix of row indices.  
              *c*             P×Q matrix of column indices.

OUTPUT     *y*             (L\*M)×(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     `let x[3,4] = 1 2 3 4 5 6 7 8 9 10 11 12;`  
              `let v1 = 1 3;`  
              `let v2 = 2 4;`  
              `y = submat(x,v1,v2);`  
              `z = submat(x,0,v2);`

$$\mathbf{x} = \begin{matrix} & \begin{matrix} 1 & 2 & 3 & 4 \end{matrix} \\ \begin{matrix} 5 & 6 & 7 & 8 \\ 9 & 10 & 11 & 12 \end{matrix} & \end{matrix}$$

## subscat

---

$$y = \begin{array}{cc} 2 & 4 \\ 10 & 12 \end{array}$$
$$z = \begin{array}{cc} 2 & 4 \\ 6 & 8 \\ 10 & 12 \end{array}$$

SEE ALSO **diag, vec, reshape**

## subscat

**PURPOSE** Changes the values in a vector depending on the category a particular element falls in.

**FORMAT**  $y = \text{subscat}(x, v, s);$

**INPUT**  $x$   $N \times 1$  vector.

$v$   $P \times 1$  numeric vector, containing breakpoints specifying the ranges within which substitution is to be made. This **MUST** be sorted in ascending order.

$v$  can contain a missing value as a separate category if the missing value is the first element in  $v$ .

If  $v$  is a scalar, all matches must be exact for a substitution to be made.

$s$   $P \times 1$  vector, containing values to be substituted.

**OUTPUT**  $y$   $N \times 1$  vector, with the elements in  $s$  substituted for the original elements of  $x$  according to which of the regions the elements of  $x$  fall

into:

$$\begin{array}{rcll} & x \leq v[1] & \rightarrow & s[1] \\ v[1] < x \leq v[2] & \rightarrow & & s[2] \\ & \dots & & \\ v[p-1] < x \leq v[p] & \rightarrow & & s[p] \\ & x > v[p] & \rightarrow & \text{the original value of } x \end{array}$$

If missing is not a category specified in *v*, missings in *x* are passed through without change.

```
EXAMPLE  let x = 1 2 3 4 5 6 7 8 9 10;
          let v = 4 5 8;
          let s = 10 5 0;
          y = subscat(x,v,s);
```

S

```

          10
          10
          10
          10
          5
y =       0
          0
          0
          9
          10
```

PURPOSE Substitutes new values for old values in a matrix, depending on the outcome of a logical expression.

FORMAT `y = substute(x,e,v);`

## substute

---

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.

                         Elements of  $x$  will be changed if the corresponding element of  $e$  is 1.

$v$              $P \times Q$  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     $x = \{ \begin{array}{l} Y \ 55 \ 30, \\ N \ 57 \ 18, \\ Y \ 24 \ 3, \\ N \ 63 \ 38, \\ Y \ 55 \ 32, \\ N \ 37 \ 11 \end{array} \};$

$e = x[.,1] \ .\$=\backslash, = "Y" \ .\text{and} \ x[.,2] \ .>= 55 \ .\text{and} \ x[.,3] \ .>= 30;$   
 $x[.,1] = \text{substute}(x[.,1],e,"R");$

$e = \begin{array}{c} 1 \\ 0 \\ 0 \\ 0 \\ 1 \\ 0 \end{array}$

Here is what **x** looks like after substitution:

```

      R  55  30
      N  57  18
      Y  24   3
x =    N  63  38
      R  55  32
      N  37  11
    
```

SOURCE    `datatran.src`

SEE ALSO    `code`, `recode`

S

subvec

PURPOSE    Extracts an N×1 vector of elements from an N×K matrix.

FORMAT    `y = subvec(x,ci);`

INPUT    *x*            N×K matrix.  
          *ci*          N×1 vector of column indices.

OUTPUT    *y*            N×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    `x = reshape(seqa(1,1,12),4,3);`  
             `ci = { 2,3,1,3 };`  
             `y = subvec(x,ci);`

## sumc

---

```
      1  2  3
x =   4  5  6
      7  8  9
     10 11 12
```

```
      2
ci =   3
      1
      3
```

```
      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×K matrix or L-dimensional array where the last two dimensions are N×K.

**OUTPUT**    *y*            K×1 vector or L-dimensional array where the last two dimensions are K×1.

**EXAMPLE**    `x = reshape(seqa(1,1,12),3,4);`  
              `y = sumc(x);`

---

```

      1  2  3  4
x =  5  6  7  8
      9 10 11 12

```

```

      15
y =   18
      21
      24

```

```

a = areshape(seqa(1,1,24),2|3|4);
z = sumc(a);

```

S

**a** is a 2×3×4 array such that:

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

```

      1  2  3  4
      5  6  7  8
      9 10 11 12

```

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

```

      13 14 15 16
      17 18 19 20
      21 22 23 24

```

**z** is a 2×4×1 array such that:

## sumr

---

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

15  
18  
21  
24

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

51  
54  
57  
60

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.

**FORMAT**  $y = \text{sumr}(x);$

**INPUT**  $x$   $N \times K$  matrix or L-dimensional array where the last two dimensions are  $N \times K$ .

**OUTPUT**  $y$   $N \times 1$  vector or L-dimensional array where the last two dimensions are  $N \times 1$ .

**EXAMPLE**  $x = \text{reshape}(\text{seqa}(1, 1, 12), 3, 4);$   
 $y = \text{sumr}(x);$



---

```
      1  2  3  4
x =   5  6  7  8
      9 10 11 12
```

```
      10
y =   26
      42
```

```
a = areshape(seqa(1,1,24),2|3|4);
z = sumr(a);
```

**a** is a 2×3×4 array such that:

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

```
      1  2  3  4
      5  6  7  8
      9 10 11 12
```

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

```
      13 14 15 16
      17 18 19 20
      21 22 23 24
```

**z** is a 2×3×1 array such that:

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

```
      10
      26
      42
```

## surface

---

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

58  
74  
90

SEE ALSO **sumc**

## surface

**PURPOSE**    Graphs a 3-D surface.

**LIBRARY**    pgraph

**FORMAT**    **surface**(*x,y,z*);

<b>INPUT</b>	<i>x</i>	1×K vector, the X axis data.
	<i>y</i>	N×1 vector, the Y axis data.
	<i>z</i>	N×K matrix, the matrix of height data to be plotted.

<b>GLOBAL INPUT</b>	<b>_psurf</b>	2×1 vector, controls 3-D surface characteristics. <b>[1]</b> if 1, show hidden lines. Default 0. <b>[2]</b> 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.
	<b>_pticout</b>	scalar, if 0 (default), tick marks point inward, if 1, tick marks point outward.
	<b>_pzclr</b>	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:

```
_pzclr = { 0.0  1,
          0.0 10,
          0.2 15 };
_pzclr[1,1] = minc(minc(z));
```

See COLORS, Section 24.5, for the list of available colors.

## svd

---

REMARKS	<b>surface</b> uses only the minimum and maximum of the <i>X</i> axis data in generating the graph and tick marks.
SOURCE	psurface.src
SEE ALSO	<b>volume</b> , <b>view</b>

## svd

PURPOSE	Computes the singular values of a matrix.	
FORMAT	$s = \text{svd}(x);$	
INPUT	$x$	$N \times P$ matrix whose singular values are to be computed.
OUTPUT	$s$	$M \times 1$ vector, where $M = \min(N, P)$ , containing the singular values of $x$ arranged in descending order.
GLOBAL INPUT	<b>_svderr</b>	scalar, if not all of the singular values can be computed, <b>_svderr</b> will be nonzero. The singular values in $s[\text{\_svderr}+1], \dots, s[M]$ will be correct.
REMARKS	Error handling is controlled with the low bit of the trap flag.  <b>trap 0</b> set <b>_svderr</b> and terminate with message <b>trap 1</b> set <b>_svderr</b> and continue execution	
EXAMPLE	$x = \begin{Bmatrix} 4 & 3 & 6 & 7, \\ & 8 & 2 & 9 & 5 \end{Bmatrix};$ $y = \text{svd}(x);$ $y = \begin{matrix} 16.521787 \\ 3.3212254 \end{matrix}$	

SOURCE    `svd.src`

SEE ALSO    `svd1`, `svd2`, `svds`

## svd1

PURPOSE    Computes the singular value decomposition of a matrix so that:  $x = u * s * v'$ .

FORMAT    `{ u,s,v } = svd1(x);`

INPUT     $x$              $N \times P$  matrix whose singular values are to be computed.

OUTPUT     $u$              $N \times N$  matrix, the left singular vectors of  $x$ .

$s$              $N \times P$  diagonal matrix, containing the singular values of  $x$  arranged in descending order on the principal diagonal.

$v$              $P \times P$  matrix, the right singular vectors of  $x$ .

GLOBAL    `_svderr`            scalar, if all of the singular values are correct, `_svderr` is 0.  
OUTPUT    If not all of the singular values can be computed, `_svderr` is set and the diagonal elements of  $s$  with indices greater than `_svderr` are correct.

REMARKS    Error handling is controlled with the low bit of the trap flag.

**trap 0**    set `_svderr` and terminate with message

**trap 1**    set `_svderr` and continue execution

EXAMPLE    `x = rndu(3,3);`  
              `{ u, s, v } = svd1(x);`

```

                0.97847012  0.20538614  0.59906497
x = 0.85474208  0.79673540  0.22482095
        0.33340653  0.74443792  0.75698778
```

S

## svd2

---

$$u = \begin{pmatrix} -0.57955818 & 0.65204491 & 0.48882486 \\ -0.61005618 & 0.05056673 & -0.79074298 \\ -0.54031821 & -0.75649219 & 0.36847767 \end{pmatrix}$$
$$s = \begin{pmatrix} 1.84994646 & 0.00000000 & 0.00000000 \\ 0.00000000 & 0.60370542 & 0.00000000 \\ 0.00000000 & 0.00000000 & 0.47539239 \end{pmatrix}$$
$$v = \begin{pmatrix} -0.68578561 & 0.71062560 & -0.15719208 \\ -0.54451302 & -0.64427479 & -0.53704336 \\ -0.48291165 & -0.28270348 & 0.82877927 \end{pmatrix}$$

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	<b>_svderr</b>	scalar, if all of the singular values are correct, <b>_svderr</b> is 0. If not all of the singular values can be computed, <b>_svderr</b> is set and the diagonal elements of <i>s</i> with indices greater than <b>_svderr</b> are correct.
REMARKS	Error handling is controlled with the low bit of the trap flag.  <b>trap 0</b> set <b>_svderr</b> and terminate with message <b>trap 1</b> set <b>_svderr</b> and continue execution	
SOURCE	svd.src	
SEE ALSO	<b>svd, svd1, svdcusv</b>	

S

svdcusv

PURPOSE	Computes the singular value decomposition of <i>x</i> so that: $x = u * s * v'$ (compact <i>u</i> ).	
FORMAT	{ <i>u,s,v</i> } = <b>svdcusv(x)</b> ;	
INPUT	<i>x</i>	N×P matrix or K-dimensional array where the last two dimensions are N×P, whose singular values are to be computed.
OUTPUT	<i>u</i>	N×N or N×P matrix or K-dimensional array where the last two dimensions are N×N or N×P, the left singular vectors of <i>x</i> . If N > P, <i>u</i> is N×P, containing only the P left singular vectors of <i>x</i> .
	<i>s</i>	N×P or P×P diagonal matrix or K-dimensional array where the last two dimensions describe N×P or P×P diagonal arrays, the singular values of <i>x</i> arranged in descending order on the principal diagonal. If N > P, <i>s</i> is P×P.
	<i>v</i>	P×P matrix or K-dimensional array where the last two dimensions are P×P, the right singular vectors of <i>x</i> .

## svds

---

**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  $10 \times 4 \times 5$  array  $x$ ,  $u$  will be a  $10 \times 4 \times 4$  array containing the left singular vectors of each of the 10 corresponding  $4 \times 5$  arrays contained in  $x$ .  $s$  will be a  $10 \times 4 \times 5$  array and  $v$  will be a  $10 \times 5 \times 5$  array both containing their respective results for each of the 10 corresponding  $4 \times 5$  arrays contained in  $x$ .

If not all of the singular values can be computed,  $s[1,1]$  is set to a scalar error code. Use **scalerr** to convert this to an integer. The diagonal elements of  $s$  with indices greater than **scalerr**( $s[1,1]$ ) are correct. If **scalerr**( $s[1,1]$ ) returns a 0, all of the singular values have been computed.

**SEE ALSO** **svd2**, **svds**, **svdusv**

## svds

**PURPOSE** Computes the singular values of a  $x$ .

**FORMAT**  $s = \text{svds}(x);$

**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.

**REMARKS** 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  $10 \times 4 \times 5$  array  $x$ ,  $s$  will be a  $10 \times 4 \times 1$  array containing the singular values of each of the 10  $4 \times 5$  arrays contained in  $x$ .

If not all of the singular values can be computed,  $s[1]$  is set to a scalar error



code. Use **scalerr** to convert this to an integer. The elements of  $s$  with indices greater than **scalerr**( $s[1]$ ) are correct. If **scalerr**( $s[1]$ ) returns a 0, all of the singular values have been computed.

SEE ALSO **svd**, **svdcusv**, **svdusv**

## svdusv

**PURPOSE** Computes the singular value decomposition of  $x$  so that:  $x = u * s * v'$ .

**FORMAT** {  $u, s, v$  } = **svdusv**( $x$ );

**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**  $u$   $N \times N$  matrix or K-dimensional array where the last two dimensions are  $N \times N$ , the left singular vectors of  $x$ .

$s$   $N \times P$  diagonal matrix or K-dimensional array where the last two dimensions describe  $N \times P$  diagonal arrays, the singular values of  $x$  arranged in descending order on the principal diagonal.

$v$   $P \times P$  matrix or K-dimensional array where the last two dimensions are  $P \times P$ , the right singular vectors of  $x$ .

**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  $10 \times 4 \times 5$  array  $x$ ,  $u$  will be a  $10 \times 4 \times 4$  array containing the left singular vectors of each of the 10 corresponding  $4 \times 5$  arrays contained in  $x$ .  $s$  will be a  $10 \times 4 \times 5$  array and  $v$  will be a  $10 \times 5 \times 5$  array both containing their respective results for each of the 10 corresponding  $4 \times 5$  arrays contained in  $x$ .

If not all of the singular values can be computed,  $s[1,1]$  is set to a scalar error code. Use **scalerr** to convert this to an integer. The diagonal elements of  $s$

S

## sysstate

---

with indices greater than **scalerr**(*s*[1,1]) are correct. If **scalerr**(*s*[1,1]) returns a 0, all of the singular values have been computed.

SEE ALSO    **svd1**, **svdcusv**, **svds**

## sysstate

PURPOSE    Gets or sets general system parameters.

FORMAT    { *rets...* } = **sysstate**(*case*,*y*);

REMARKS    The available cases are as follows:

<b>Case 1</b>	<b>Version Information</b> Returns the current <b>GAUSS</b> version information in an 8-element numeric vector.
<b>Cases 2-7</b>	<b>GAUSS System Paths</b> Gets or sets <b>GAUSS</b> system path.
<b>Case 8</b>	<b>Complex Number Toggle</b> Controls automatic generation of complex numbers in <b>sqrt</b> , <b>ln</b> , and <b>log</b> for negative arguments.
<b>Case 9</b>	<b>Complex Trailing Character</b> Gets or sets trailing character for the imaginary part of a complex number.
<b>Case 10</b>	<b>Printer Width</b> Gets or sets <b>lprint</b> width.
<b>Case 11</b>	<b>Auxiliary Output Width</b> Gets or sets the auxiliary output width.
<b>Case 13</b>	<b>LU Tolerance</b> Gets or sets singularity tolerance for LU decomposition in current thread.

---

<b>Case 14</b>	<b>Cholesky Tolerance</b> Gets or sets singularity tolerance for Cholesky decomposition in current thread.
<b>Case 15</b>	<b>Screen State</b> Gets or sets window state as controlled by <b>screen</b> command.
<b>Case 18</b>	<b>Auxiliary Output</b> Gets auxiliary output parameters.
<b>Case 19</b>	<b>Get/Set Format</b> Gets or sets format parameters.
<b>Case 21</b>	<b>Imaginary Tolerance</b> Gets or sets imaginary tolerance in current thread.
<b>Case 22</b>	<b>Source Path</b> Gets or sets the path the compiler will search for source files.
<b>Case 24</b>	<b>Dynamic Library Directory</b> Gets or sets the path for the default dynamic library directory.
<b>Case 25</b>	<b>Temporary File Path</b> Gets or sets the path <b>GAUSS</b> will use for temporary files.
<b>Case 26</b>	<b>Interface Mode</b> Returns the current interface mode.
<b>Case 28</b>	<b>Random Number Generator Parameters</b> Gets or sets parameters used by the random number generation commands.
<b>Case 30</b>	<b>Base Year Toggle</b> Specifies whether year value returned by <b>date</b> is to include base year (1900) or not.
<b>Case 32</b>	<b>Global LU Tolerance</b> Gets or sets global singularity tolerance for LU decomposition.
<b>Case 33</b>	<b>Global Cholesky Tolerance</b> 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**  $vi = \text{sysstate}(1,0);$

**OUTPUT**  $vi$  8×1 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
- 3 IBM RS/6000
- 4 HP 9000
- 5 SGI MIPS
- 6 DEC Alpha
- 7 Mac 32-bit PowerPC

$vi[5]$  indicates the operating system on which **GAUSS** is running:

- 1 DOS
- 2 SunOS 4.1.x
- 3 Solaris 2.x

- 4 AIX
- 5 HP-UX
- 6 IRIX
- 7 OSF/1
- 8 OS/2
- 9 Windows

Cases 2-7: GAUSS System Paths

PURPOSE	Gets or sets <b>GAUSS</b> system path.	
FORMAT	<i>oldpath</i> = <b>sysstate</b> ( <i>case</i> , <i>path</i> );	
INPUT	<i>case</i>	scalar 2-7, path to set. <ul style="list-style-type: none"><li>2 .exe file location.</li><li>3 <b>loadexe</b> path.</li><li>4 <b>save</b> path.</li><li>5 <b>load</b>, <b>loadm</b> path.</li><li>6 <b>loadf</b>, <b>loadp</b> path.</li><li>7 <b>loads</b> path.</li></ul>
	<i>path</i>	scalar 0 to get path, or string containing the new path.
OUTPUT	<i>oldpath</i>	string, original path.
REMARKS	If <i>path</i> 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 <b>sqrt</b> , <b>ln</b> and <b>log</b> for negative arguments.	
FORMAT	<i>oldstate</i> = <b>sysstate</b> (8, <i>state</i> );	

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## sysstate

---

INPUT	<i>state</i>	scalar, 1, 0, or -1
OUTPUT	<i>oldstate</i>	scalar, the original state.
REMARKS	If <i>state</i> = 1, <b>log</b> , <b>ln</b> , and <b>sqrt</b> will return complex numbers for negative arguments. If <i>state</i> = 0, the program will terminate with an error message when negative numbers are passed to <b>log</b> , <b>ln</b> , and <b>sqrt</b> . If <i>state</i> = -1, the current state is returned and left unchanged. The default state is 1.	

### Case 9: Complex Trailing Character

PURPOSE	Gets or sets trailing character for the imaginary part of a complex number.	
FORMAT	<i>oldtrail</i> = <b>sysstate(9, trail)</b> ;	
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 “ <b>i</b> ”.	

### Case 10: Printer Width

PURPOSE	Gets or sets <b>lprint</b> width.	
FORMAT	<i>oldwidth</i> = <b>sysstate(10, width)</b> ;	
INPUT	<i>width</i>	scalar, new printer width.
OUTPUT	<i>oldwidth</i>	scalar, the current original width.

---

REMARKS    If *width* is 0, the printer width will not be changed.

             This may also be set with the **lpwidth** command.

SEE ALSO    **lpwidth**

### 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**

### 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.

## sysstate

---

REMARKS     The tolerance must be  $\geq 0$ . 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       *tol*            scalar, new tolerance.

OUTPUT      *oldtol*       scalar, the original tolerance.

REMARKS     The tolerance must be  $\geq 0$ . 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.

SEE ALSO     **chol, invpd, solpd**

### Case 15: Screen State

PURPOSE     Gets or sets window state as controlled by **screen** command.

FORMAT     *oldstate* = **sysstate(15, *state*)**;



---

INPUT	<i>state</i>	scalar, new window state.
OUTPUT	<i>oldstate</i>	scalar, the original window state.
REMARKS	If <i>state</i> = 1, window output is turned on. If <i>state</i> = 0, window output is turned off. If <i>state</i> = -1, the state is returned unchanged.	
SEE ALSO	<b>screen</b>	

**Case 18: Auxiliary Output**

PURPOSE	Gets auxiliary output parameters.	
FORMAT	{ <i>state</i> , <i>name</i> } = <b>sysstate(18,0)</b> ;	
OUTPUT	<i>state</i>	scalar, auxiliary output state, 1 - on, 0 - off.
	<i>name</i>	string, auxiliary output filename.
SEE ALSO	<b>output</b>	

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**Case 19: Get/Set Format**

PURPOSE	Gets or sets format parameters.	
FORMAT	<i>oldfmt</i> = <b>sysstate(19,fmt)</b> ;	
INPUT	<i>fmt</i>	scalar or 11×1 column vector containing the new format parameters. Usually this will have come from a previous <b>sysstate(19,0)</b> call. See Output for description of matrix.
OUTPUT	<i>oldfmt</i>	11×1 vector containing the current format parameters. The characters in quotes are components of the format string that gets passed through to the C library <b>sprintf</b> function:

- [1] format conversion type:
  - 0 string format (“**s**”).
  - 1 compact format (“**g**”).
  - 2 auto format (“**#g**”).
  - 3 scientific format (“**e**”).
  - 4 decimal format (“**f**”).
  - 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 *fnt* is scalar 0, then the format parameters will be left unchanged.

See the **format** and **print** commands for more information on the formatting parameters.

S

**SEE ALSO** **format, print**

### **Case 21: Imaginary Tolerance**

**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.

## sysstate

---

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 *path* scalar 0 to get path, or string containing the new path.

OUTPUT *oldpath* 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)**;

INPUT *path* scalar 0 to get path, or string containing the new path.

OUTPUT *oldpath* 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.

## sysstate

---

FORMAT    *mode* = **sysstate(26,0)**;

OUTPUT    *mode*        scalar, interface mode flag

**0**    non-X mode

**1**    terminal (-v) mode

**2**    X Windows mode

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      *prms*        scalar 0 to get parameters, or 3×1 matrix of new parameters.

**[1]** seed,             $0 < \text{seed} < 2^{32}$

**[2]** multiplier,     $0 < \text{mult} < 2^{32}$

**[3]** constant,      $0 \leq \text{const} < 2^{32}$

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 <b>date</b> is to include base year (1900) or not.
FORMAT	<i>oldstate</i> = <b>sysstate(30, state)</b> ;
INPUT	<i>state</i> scalar, 1, 0, or missing value.
OUTPUT	<i>oldstate</i> scalar, the original state.
REMARKS	<p>Internally, <b>date</b> acquires the number of years since 1900. <b>sysstate</b> case 30 specifies whether <b>date</b> should add the base year to that value or not. If <i>state</i> = 1, <b>date</b> adds 1900, returning a fully-qualified 4-digit year.</p> <p>If <i>state</i> = 0, <b>date</b> returns the number of years since 1900. If <i>state</i> is a missing value, the current state is returned. The default state is 1.</p>

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**Case 32: Global LU Tolerance**

PURPOSE	Gets or sets global singularity tolerance for LU decomposition.
FORMAT	<i>oldtol</i> = <b>sysstate(32, tol)</b> ;
INPUT	<i>tol</i> scalar, new tolerance.
OUTPUT	<i>oldtol</i> scalar, the original tolerance.
REMARKS	<p>The tolerance must be <math>\geq 0</math>. If <i>tol</i> is negative, the tolerance is returned and left unchanged.</p> <p>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.</p>
SEE ALSO	<b>croutp, inv</b>

## sysstate

---

### Case 33: Global Cholesky Tolerance

PURPOSE	Gets or sets global singularity tolerance for Cholesky decomposition.	
FORMAT	<i>oldtol</i> = <b>sysstate(33,<i>tol</i>)</b> ;	
INPUT	<i>tol</i>	scalar, new tolerance.
OUTPUT	<i>oldtol</i>	scalar, the original tolerance.
REMARKS	<p>The tolerance must be <math>\geq 0</math>. If <i>tol</i> is negative, the tolerance is returned and left unchanged.</p> <p>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.</p>	
SEE ALSO	<b>chol</b> , <b>invpd</b> , <b>solpd</b>	

### Case 34: Global Imaginary Tolerance

PURPOSE	Gets or sets the global imaginary tolerance.	
FORMAT	<i>oldtol</i> = <b>sysstate(34,<i>tol</i>)</b> ;	
INPUT	<i>tol</i>	scalar, the new tolerance.
OUTPUT	<i>oldtol</i>	scalar, the original tolerance.
REMARKS	<p>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.</p> <p>If <i>tol</i>&lt;0, the current tolerance is returned.</p>	



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 *c* 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**

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## tab

PURPOSE Tabs the cursor to a specified text column.

FORMAT **tab**(*col*);  
**print** **expr1** **expr2** **tab**(*col1*) **expr3** **tab**(*col2*) **expr4** ...;

INPUT *col* scalar, the column position to tab to.

## **tan**

---

**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**     $y = \tan(x);$

**INPUT**       $x$             N×K matrix or N-dimensional array.

**OUTPUT**     $y$             N×K 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  $\frac{\pi}{180}$ .

**EXAMPLE**    `let x = 0 .5 1 1.5;`  
              `y = tan(x);`

```
0.00000000
0.54630249
y = 1.55740772
14.10141995
```

SEE ALSO    **atan, pi**

tanh

PURPOSE    Computes the hyperbolic tangent.

FORMAT     $y = \tanh(x);$

INPUT     $x$              $N \times K$  matrix or  $N$ -dimensional array.

OUTPUT     $y$              $N \times K$  matrix or  $N$ -dimensional array containing the hyperbolic tangents of the elements of  $x$ .

EXAMPLE    `let x = -0.5   -0.25   0   0.25   0.5   1;`  
             `x = x * pi;`  
             `y = tanh(x);`

```
-1.570796
-0.785398
x = 0.000000
0.785398
1.570796
3.141593
```

## tempname

---

```
          -0.917152
          -0.655794
          0.000000
y =       0.655794
          0.917152
          0.996272
```

SOURCE    trig.src

## tempname

PURPOSE    Creates a temporary file with a unique name.

FORMAT    *tname* = **tempname**(*path*,*pre*,*suf*);

INPUT    *path*        string, path where the file will reside.  
         *pre*        string, a prefix to begin the file name with.  
         *suf*        string, a suffix to end the file name with.

OUTPUT    *tname*        string, unique temporary file name of the form  
                 *path/preXXXXnnnnnsuf*, 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.

time

PURPOSE

Returns the current system time.

FORMAT

$y = \text{time};$

OUTPUT

$y$             4×1 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

t

timedt

PURPOSE

Returns system date and time in DT scalar format.

FORMAT

$dt = \text{timedt};$

OUTPUT

$dt$             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

20050306071511

## **timestr**

---

represents 07:15:11 or 7:15:11 AM on March 6, 2005.

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* 4×1 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.

**FORMAT** *tc* = **timeutc**;

**OUTPUT** *tc* scalar, number of seconds since January 1, 1970 Greenwich Mean Time.

**EXAMPLE** `tc = timeutc;  
utv = utctodtv(tc);`

`tc = 1125511090`

`utv = 2005 8 31 10 58 10 3 242`

**SEE ALSO** **dtvnormal**, **utctodtv**

t

## title

**PURPOSE** Sets the title for the graph.

**LIBRARY** **pgraph**

**FORMAT** **title**(*str*);

**INPUT** *str* string, the title to display above the graph.

## tkf2eps

---

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 [24.4.1](#).

SOURCE `pgraph.src`

SEE ALSO `xlabel`, `ylabel`, `fonts`

## tkf2eps

PURPOSE Converts a .tkf file to an Encapsulated PostScript file.

LIBRARY `pgraph`

FORMAT `ret = tkf2eps(tekfile,epsfile);`

INPUT *tekfile* string, name of .tkf file.  
*epsfile* string, name of Encapsulated PostScript file.

OUTPUT *ret* 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.

LIBRARY     `pgraph`

FORMAT     `ret = tkf2ps(tekfile,psfile);`

INPUT       *tekfile*       string, name of `.tkf` file.  
              *epsfile*       string, name of Encapsulated PostScript file.

OUTPUT      *ret*           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 Encapsulated PostScript file and a PostScript manual if you want to modify these parameters.

t

tocart

PURPOSE     Converts from polar to cartesian coordinates.

FORMAT     `xy = tocart(r,theta);`

INPUT       *r*                 $N \times K$  real matrix, radius.  
              *theta*         $L \times M$  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.

## todaydt

---

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    *dt*            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

20050306130525

represents 13:05:25 or 1:05:25 PM on March 6, 2005.

SOURCE    `time.src`

SEE ALSO    **timedt**, **timeutc**, **dtdate**

### toeplitz

PURPOSE    Creates a Toeplitz matrix from a column vector.

FORMAT    `t = toeplitz(x);`

INPUT    *x*            K×1 vector.

OUTPUT    *t*            K×K Toeplitz matrix.

```
EXAMPLE  x = seqa(1,1,5);
          y = toeplitz(x);
```

```
          1
          2
x =       3
          4
          5

          1 2 3 4 5
          2 1 2 3 4
y =       3 2 1 2 3
          4 3 2 1 2
          5 4 3 2 1
```

```
SOURCE  toeplitz.src
```

t

PURPOSE Extracts the leading token from a string.

```
FORMAT  { token, str_left } = token(str);
```

INPUT str string, the string to parse.

OUTPUT token string, the first token in str.  
 str\_left string, str minus token.

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.

## topolar

---

EXAMPLE Here is a keyword that uses **token** to parse its string parameter:

```
keyword add(s);
  local tok,sum;
  sum = 0;
  do until s $=\,,= "";
    { tok, s } = token(s);
    sum = sum + stof(tok);
  endo;
  format /rd 1,2;
  print "Sum is: " sum;
endp;
```

If you type:

```
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>	N×K 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>	N×K real matrix, radius.
	<i>theta</i>	N×K real matrix, angle in radians.
SOURCE	coord.src	

trace

PURPOSE Allows the user to trace program execution for debugging purposes.

FORMAT **trace** *new*;  
**trace** *new*, *mask*;

INPUT *new* scalar, new value for trace flag.  
*mask* 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:

```
trace 16,16;  turn on output to printer
trace 0,16;   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**

**PURPOSE** Sets the trap flag to enable or disable trapping of numerical errors.

**FORMAT** **trap** *new*;  
**trap** *new*, *mask*;

**INPUT** *new* scalar, new trap value.  
*mask* scalar, optional mask to allow leaving some bits of the trap flag unchanged.

**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

**EXAMPLE**

```
proc(0) = printinv(x);
    local oldval,y;
    oldval = trapchk(1);
    trap 1,1;
    y = inv(x);
    trap oldval,1;
    if scalerr(y);
        errorlog "WARNING: x is singular";
    else;
```

```
        print "y" y;  
    endif;  
endp;
```

In this example the result of **inv** is trapped in case **x** is singular. The trap state is reset to the original value after the call to **inv**.

Calling **printinv** as follows:

```
x = eye(3);  
printinv(x);
```

produces:

```
y =  
    1.00000000    0.00000000    0.00000000  
    0.00000000    1.00000000    0.00000000  
    0.00000000    0.00000000    1.00000000
```

while

```
x = ones(3,3);  
printinv(x);
```

produces:

```
WARNING: x is singular
```

SEE ALSO **scalerr**, **trapchk**, **error**



trapchk

PURPOSE Tests the value of the trap flag.

FORMAT  $y = \text{trapchk}(m);$

INPUT  $m$  scalar mask value.

OUTPUT  $y$  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:

t

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

# trigamma

---

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

xM×N matrix or N-dimensional array.
- OUTPUT

yM×N 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      $x$               $N \times K$  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      $y$               $R \times K$  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      $x = \text{rndu}(5,3);$   
                 $y = \text{trimr}(x,2,1);$

$x =$              0.76042751   0.33841579   0.01844780  
                 0.05334503   0.38939785   0.65029973  
                 0.93077511   0.06961078   0.04207563  
                 0.53640701   0.06640062   0.07222560  
                 0.14084669   0.06033813   0.69449247

$y =$              0.93077511   0.06961078   0.04207563  
                 0.53640701   0.06640062   0.07222560

SEE ALSO     **submat, rotater, shiftr**

## type

---

### trunc

**PURPOSE**    Converts numbers to integers by truncating the fractional portion.

**FORMAT**     $y = \text{trunc}(x);$

**INPUT**       $x$              $N \times K$  matrix or  $N$ -dimensional array.

**OUTPUT**     $y$              $N \times K$  matrix or  $N$ -dimensional array containing the truncated elements of  $x$ .

**EXAMPLE**     $x = 100 * \text{rndn}(2, 2);$   
                   $y = \text{trunc}(x);$

$$x = \begin{bmatrix} 77.68 & -14.10 \\ 4.73 & -158.88 \end{bmatrix}$$
$$y = \begin{bmatrix} 77.00 & -14.00 \\ 4.00 & -158.00 \end{bmatrix}$$

**SEE ALSO**    **ceil, floor, round**

### type

**PURPOSE**    Returns the symbol table type of its 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

REMARKS    **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.

EXAMPLE    `k = { "CHARS" };`  
             `print k;`  
             `if type(k) =\,= 6;`  
                 `k = "" $+ k; /* force matrix to string */`  
             `endif;`  
             `print k;`

produces:

                                 +DEN

CHARS

SEE ALSO    **typecv**, **typedef**

t

## 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** `x` string, or N×1 character vector or string array which contains the names of variables whose type is to be determined.

**OUTPUT** `y` scalar or N×1 vector containing the types of the respective symbols in `x`.

**REMARKS** The values returned by **typecv** for the various variable types are as follows:

- 5 keyword (**keyword**)
- 6 matrix (numeric, character, or mixed)
- 8 procedure (**proc**)
- 9 function (**fn**)
- 13 string
- 15 string array
- 17 structure
- 21 array
- 23 structure pointer

**typecv** 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 = "Montana";
fn area(r) = pi*r*r;
let names = xvar yvar area;
y = typecv(names);
```

```
          XVAR
names =   YVAR
          AREA
```

6  
y = 13  
9

SEE ALSO **type, typedef, 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 *fp* scalar, file handle of an open file.

OUTPUT *y* 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 

```
infile = "dat1";
outfile = "dat2";
open fin = ^infile;
names = getname(infile);
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**

t

## union

---

### union

**PURPOSE** Returns the union of two vectors with duplicates removed.

**FORMAT** `y = union(v1,v2,flag);`

<b>INPUT</b>	<i>v1</i>	N×1 vector.
	<i>v2</i>	M×1 vector.
	<i>flag</i>	scalar, 1 if numeric data, 0 if character.

**OUTPUT** *y* L×1 vector containing all unique values that are in *v1* and *v2*, sorted in ascending order.

**REMARKS** The combined elements of *v1* and *v2* must fit into a single vector.

**EXAMPLE** `let v1 = mary jane linda john;  
let v2 = mary sally;  
x = union(v1,v2,0);`

```
      JANE  
      JOHN  
x =  LINDA  
      MARY  
      SALLY
```

### unionsa

**PURPOSE** Returns the union of two string vectors with duplicates removed.



FORMAT    `y = unionsa(sv1,sv2);`

INPUT     *sv1*        N×1 or 1×N string vector.  
            *sv2*        M×1 or 1×M string vector.

OUTPUT    *y*         L×1 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);`

                 jane  
                 john  
*y* = linda  
                 mary  
                 sally

SOURCE    `unionsa.src`

SEE ALSO   **union**

u

uniqindx

PURPOSE   Computes the sorted index of *x*, leaving out duplicate elements.

FORMAT    `index = uniqindx(x,flag);`

INPUT     *x*         N×1 or 1×N vector.  
            *flag*       scalar, 1 if numeric data, 0 if character.

## uniqindxsa

---

OUTPUT    *index*        M×1 vector, indices corresponding to the elements of *x* 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;`  
             `ind = uniqindx(x,1);`  
             `y = x[ind];`

```

              7
              6
ind = 5
      2
      1
```

```

      1
      2
y = 3
     4
     5
```

SEE ALSO    **unique**, **uniqindxsa**

## uniqindxsa

PURPOSE    Computes the sorted index of a string vector, omitting duplicate elements.

FORMAT    *ind* = **uniqindxsa**(*sv*);

INPUT      *sv*            N×1 or 1×N string vector.

OUTPUT     *ind*            M×1 vector, indices corresponding to the elements of *sv* 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",
             "jane","cindy","betty"};
ind = uniqindxsa(sv);
y = sv[ind];
```

```

              7
              6
ind = 5
      2
      1
```

```

      betty
      cindy
y = jane
    linda
    mary
```

SOURCE     `uniquesa.src`

SEE ALSO    **unique**, **uniquesa**, **uniqindx**

u

unique

PURPOSE    Sorts and removes duplicate elements from a vector.

FORMAT     `y = unique(x,flag);`

## uniquesa

---

INPUT	$x$	$N \times 1$ or $1 \times N$ vector.
	$flag$	scalar, 1 if numeric data, 0 if character.
OUTPUT	$y$	$M \times 1$ vector, sorted $x$ with the duplicates removed.
EXAMPLE	<pre>let x = 5 4 4 3 3 2 1; y = unique(x,1);</pre>	

```
      1  
      2  
y =   3  
      4  
      5
```

SEE ALSO    **uniquesa**, **uniqindx**

## uniquesa

PURPOSE    Removes duplicate elements from a string vector.

FORMAT     $y = \text{uniquesa}(sv);$

INPUT	$sv$	$N \times 1$ or $1 \times N$ string vector.
-------	------	---

OUTPUT	$y$	sorted $M \times 1$ string vector containing all unique elements found in $sv$ .
--------	-----	--

EXAMPLE	<pre>string sv1 = { "mary", "jane", "mary", "linda", "john", "jane" }; y = uniquesa(sv);</pre>	
---------	--	--

```
      jane
y =    john
      linda
      mary
```

SOURCE    `uniquesa.src`

SEE ALSO    `unique`, `uniqindxsa`, `uniqindx`

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);`

INPUT      `x`            N×K matrix.

OUTPUT    `u`            N×K 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    `x = { 1  2 -1,`  
              `2  3 -2,`  
              `1 -2  1 };`  
  
            `u = upmat(x);`  
            `u1 = upmat1(x);`

u

## upper

---

The resulting matrices are

$$\mathbf{u} = \begin{bmatrix} 1 & 2 & -1 \\ 0 & 3 & -2 \\ 0 & 0 & 1 \end{bmatrix}$$

$$\mathbf{u1} = \begin{bmatrix} 1 & 2 & -1 \\ 0 & 1 & -2 \\ 0 & 0 & 1 \end{bmatrix}$$

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 N×K matrix, or string array containing the character data to be converted to uppercase.

OUTPUT `y` string, or N×K 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 `x = "uppercase";`

---

```
y = upper(x);
```

```
y = "UPPERCASE"
```

SEE ALSO **lower**

use

**PURPOSE** Loads a compiled file at the beginning of the compilation of a source program.

**FORMAT** **use** *fname*;

**INPUT** *fname* literal or ^string, the name of a compiled file created using the **compile** or the **saveall** 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:

u

```
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.

**use** can appear only ONCE at the TOP of a program.

SEE ALSO **compile, run, saveall**



---

PURPOSE	Converts UTC scalar format to DT scalar format.	
FORMAT	<i>dt</i> = <b>utctodt</b> ( <i>utc</i> );	
INPUT	<i>utc</i>	N×1 vector, UTC scalar format.
OUTPUT	<i>dt</i>	N×1 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	<pre>tc = 1126290409; print "tc = " tc; dt = utctodt(tc); print "dt = " dt;</pre> <p>produces:</p> <pre>tc = 1126290409 dt = 20050909112649</pre>	
SOURCE	time.src	
SEE ALSO	<b>dtvnormal</b> , <b>timeutc</b> , <b>utctodtv</b> , <b>dtctodtv</b> , <b>dtvtodt</b> , <b>dtctoutc</b> , <b>dtvtodt</b> , <b>strtodt</b> , <b>dttostr</b>	

u

---

**utctodtv**

PURPOSE	Converts UTC scalar format to DTV vector format.	
FORMAT	<i>dtv</i> = <b>utctodtv</b> ( <i>utc</i> );	

## utrisol

---

INPUT	<i>utc</i>	N×1 vector, UTC scalar format.																
OUTPUT	<i>dtv</i>	N×8 matrix, DTV vector format.																
REMARKS	<p>A UTC scalar gives the number of seconds since or before January 1, 1970 Greenwich Mean Time.</p> <p>Each row of <i>dtv</i>, in DTV vector format, contains:</p> <table><tr><td>[N,1]</td><td>Year, four digit integer.</td></tr><tr><td>[N,2]</td><td>Month in Year, 1-12.</td></tr><tr><td>[N,3]</td><td>Day of month, 1-31.</td></tr><tr><td>[N,4]</td><td>Hours since midnight, 0-23.</td></tr><tr><td>[N,5]</td><td>Minutes, 0-59.</td></tr><tr><td>[N,6]</td><td>Seconds, 0-59.</td></tr><tr><td>[N,7]</td><td>Day of week, 0-6, 0=Sunday.</td></tr><tr><td>[N,8]</td><td>Days since Jan 1 of current year, 0-365.</td></tr></table>		[N,1]	Year, four digit integer.	[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.
[N,1]	Year, four digit integer.																	
[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	<pre>tc = timeutc; print "tc = " tc; dtv = utctodtv(tc); print "dtv = " dtv;</pre> <p>produces:</p> <pre>tc = 1126290724 dtv = 2005  9    9    11   32   4    5    251</pre>																	
SEE ALSO	<b>dtvnormal</b> , <b>timeutc</b> , <b>utctodt</b> , <b>dtctodtv</b> , <b>dtctoutc</b> , <b>dtvtodt</b> , <b>dtvtoutc</b> , <b>strtodt</b> , <b>dttostr</b>																	

## utrisol

**PURPOSE**    Computes the solution of  $Ux = b$  where  $U$  is an upper triangular matrix.

FORMAT	$x = \text{utrisol}(b,U);$	
INPUT	$b$	P×K matrix.
	$U$	P×P upper triangular matrix.
OUTPUT	$x$	P×K matrix, solution of $Ux = b$ .
REMARKS	<b>utrisol</b> 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., <b>utrisol</b> applies a back solve to $Ux[:,i] = b[:,i]$ .	

PURPOSE	Converts a string into a matrix of its ASCII values.	
FORMAT	$y = \text{vals}(s);$	
INPUT	$s$	string of length N where $N > 0$ .
OUTPUT	$y$	N×1 matrix containing the ASCII values of the characters in the string $s$ .
REMARKS	If the string is null, the function will fail and an error message will be given.	
EXAMPLE	<pre>k0:     k = key;     if not k;         goto k0;     endif;     if k = \, = vals("Y") or k = \, = vals("y");         goto doit;     else;         end;</pre>	

## varget

---

```
endif;
```

```
doit:
```

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 jump to the label **doit**, otherwise the program will end.

SEE ALSO **chrs, ftos, stof**

## varget

**PURPOSE**    Accesses a global variable whose name is given as a string argument.

**FORMAT**    `y = varget(s);`

**INPUT**      `s`            string containing the name of the global symbol you wish to access.

**OUTPUT**    `y`            contents of the variable whose name is in `s`.

**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**    `dog = rndn(2,2);`  
              `y = varget("dog");`

```
dog =  -0.83429985  0.34782433
       0.91032546  1.75446391
```

---

```

y =  -0.83429985  0.34782433
      0.91032546  1.75446391

```

SEE ALSO    **typecv, varput**

## vargetl

**PURPOSE**    Accesses a local variable whose name is given as a string argument.

**FORMAT**    `y = vargetl(s);`

**INPUT**      `s`            string containing the name of the local symbol you wish to access.

**OUTPUT**    `y`            contents of the variable whose name is in `s`.

**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 dog;
  local x,y;
  x = rndn(2,2);
  y = vargetl("x");
  print "x" x;
  print "y" y;
  retp(y);
endp;
z = dog;
print "z" z;

```

```

x
-0.543851    -0.181701
-0.108873    0.0648738

```

**v**

## varmall

---

y	-0.543851	-0.181701
	-0.108873	0.0648738
z	-0.543851	-0.181701
	-0.108873	0.0648738

SEE ALSO **varput1**

## varmall

**PURPOSE** Computes log-likelihood of a Vector ARMA model.

**FORMAT**  $ll = \text{varmall}(w, \phi, \theta, vc);$

**INPUT**  $w$   $N \times K$  matrix, time series.

$\phi$   $(K \times P) \times K$  matrix, AR coefficient matrices.

$\theta$   $(K \times Q) \times K$  matrix, MA coefficient matrices.

$vc$   $K \times K$  matrix, covariance matrix.

**OUTPUT**  $ll$  scalar, log-likelihood. If the calculation fails  $ll$  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.

v

varmares

PURPOSE     Computes residuals of a Vector ARMA model.

FORMAT     *res* = **varmares**(*w,phi,theta*);

INPUT     *w*             $N \times K$  matrix, time series.  
             *phi*           $(K * P) \times K$  matrix, AR coefficient matrices.  
             *theta*         $(K * Q) \times K$  matrix, MA coefficient matrices.

OUTPUT     *res*             $N \times K$  matrix, residuals. If the calculation fails *res* is set to missing value with error code:

## varput

---

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.

EXAMPLE     `source = rndn(2,2);`  
               `targname = "target";`  
               `if not varput(source,targname);`  
                   `print "Symbol table full";`  
                   `end;`  
               `endif;`

```
source =  -0.93519984  0.40642598
          -0.36867581  2.57623519
```

```
target =  -0.93519984  0.40642598
          -0.36867581  2.57623519
```

SEE ALSO     **varget**, **typecv**

v

## varputl

**PURPOSE**     Allows a matrix, array, string, or string array to be assigned to a local symbol whose name is given as a string argument.

**FORMAT**     `y = varputl(x,n);`

<b>INPUT</b>	<i>x</i>	matrix, array, string, or string array which is to be assigned to the target variable.
	<i>n</i>	string containing the name of the local symbol which will be the target variable.

## varputl

---

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 dog(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;
    do while putvar $/= vars;
        print "Assign x (" $vars ")": ";
        putvar = upper(cons);
        print;
    endo;
    call varputl(x,putvar);
    retp(a+b*c-d/e);
endp;

format /rds 2,1;
i = 0;
do until i >= 5;
    z = dog(17);
    print "    z is " z;
    i = i + 1;
endo;
```

produces:

```
Assign x ( A B C D E ): a
      z is 22.3
Assign x ( A B C D E ): b
      z is 51.3
Assign x ( A B C D E ): c
      z is 34.3
Assign x ( A B C D E ): d
```

```

      z is 4.6
Assign x ( A  B  C  D  E ): e
      z is 6.7

```

SEE ALSO    **varget1**

vartypef

PURPOSE	Returns a vector of ones and zeros that indicate whether variables in a data set are character or numeric.		
FORMAT	$y = \mathbf{vartypef}(f);$		
INPUT	$f$	file handle of an open file.	
OUTPUT	$y$	$N \times 1$ 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 <b>v96</b> data set format.		

v

vcm, vcx

PURPOSE	Computes a variance-covariance matrix.		
FORMAT	$vc = \textbf{vcm}(m);$  $vc = \textbf{vcx}(x);$		
INPUT	$m$	$K \times K$ moment ( $x'x$ ) matrix. A constant term <b>MUST</b> have been the first variable when the moment matrix was computed.	

## vcms, vcxs

---

	$x$	N×K matrix of data.
OUTPUT	$vc$	K×K variance-covariance matrix.
SOURCE	corr.src	
SEE ALSO	momentd	

## vcms, vcxs

PURPOSE	Computes a sample variance-covariance matrix.	
FORMAT	$vc = \text{vcms}(m);$ $vc = \text{vcxs}(x);$	
INPUT	$m$	K×K moment ( $x'x$ ) matrix. A constant term <b>MUST</b> have been the first variable when the moment matrix was computed.
	$x$	N×K matrix of data.
OUTPUT	$vc$	K×K variance-covariance matrix.
REMARKS	Computes sample covariance matrix, that is, it divides the sample size, <b>N</b> , rather than <b>N</b> - 1. For population covariance matrix which uses <b>N</b> - 1 rather than <b>N</b> see <b>vcm</b> or <b>vcx</b> .	
SOURCE	corrs.src	
SEE ALSO	momentd, corrms, corrvcs, corrxs	

**PURPOSE** Creates a column vector by appending the columns/rows of a matrix to each other.

**FORMAT**  $yc = \mathbf{vec}(x);$   
 $yr = \mathbf{vecr}(x);$

**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**  $x = \begin{Bmatrix} 1 & 2, \\ 3 & 4 \end{Bmatrix};$   
 $yc = \mathbf{vec}(x);$   
 $yr = \mathbf{vecr}(x);$

$$x = \begin{pmatrix} 1.000000 & 2.000000 \\ 3.000000 & 4.000000 \end{pmatrix}$$
$$yc = \begin{pmatrix} 1.000000 \\ 3.000000 \\ 2.000000 \\ 4.000000 \end{pmatrix}$$

## vech

---

```
yr = 1.000000
      2.000000
      3.000000
      4.000000
```

### 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**  $x = \text{seqa}(10,10,3) + \text{seqa}(1,1,3)';$   
 $v = \mathbf{vech}(x);$   
 $sx = \mathbf{xpnd}(v);$

```
x = 11 12 13
     21 22 23
     31 32 33
```

```

      11
      21
      22
v =    31
      32
      33

      11  21  31
sx =    21  22  32
      31  32  33
```

SEE ALSO    **xpnd**

vector (dataloop)

**v**

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.

## vget

---

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    *dbuf*        N×1 vector, a data buffer containing various strings and matrices.  
          *name*       string, the name of the string or matrix to extract from *dbuf*.

OUTPUT    *x*            L×M matrix or string, the item extracted from *dbuf*.  
          *dbufnew*   K×1 vector, the remainder of *dbuf* 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.

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

v

PURPOSE To set the position of the observer in plot coordinates for 3-D plots.

LIBRARY pgraph

FORMAT **viewxyz**( $x,y,z$ );

INPUT  $x$  scalar, the X position in plot coordinates.  
 $y$  scalar, the Y position in plot coordinates.  
 $z$  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.

## vlist

---

SOURCE    `pgraph.src`

SEE ALSO    **volume, view**

## vlist

PURPOSE    Lists the contents of a data buffer constructed with **vput**.

FORMAT    **vlist**(*dbuf*);

INPUT    *dbuf*         $N \times 1$  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    *dbuf*         $N \times 1$  vector, a data buffer containing various strings and matrices.

OUTPUT    *cv*          $K \times 1$  character vector containing the names of the elements of *dbuf*.

SEE ALSO    **vget, vput, vread, vtypecv**

## volume

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

**LIBRARY** pgraph

**FORMAT** **volume**(*x,y,z*);

**INPUT** *x* scalar, the X length of the 3-D workbox.  
*y* scalar, the Y length of the 3-D workbox.  
*z* 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**

v

## vput

**PURPOSE** Inserts a matrix or string into a data buffer.

**FORMAT** *dbufnew* = **vput**(*dbuf,x,xname*);

**INPUT** *dbuf* N×1 vector, a data buffer containing various strings and matrices. If *dbuf* is a scalar 0, a new data buffer will be created.  
*x* L×M matrix or string, item to be inserted into *dbuf*.  
*xname* string, the name of *x*, will be inserted with *x* into *dbuf*.

## vread

---

OUTPUT	<i>dbufnew</i>	$K \times 1$ vector, the data buffer after <i>x</i> and <i>xname</i> have been inserted.
REMARKS	If <i>dbuf</i> already contains <i>x</i> , the new value of <i>x</i> will replace the old one.	
SOURCE	<code>vpack.src</code>	
SEE ALSO	<b>vget</b> , <b>vlist</b> , <b>vread</b>	

## vread

PURPOSE	Reads a string or matrix from a data buffer constructed with <b>vput</b> .	
FORMAT	$x = \mathbf{vread}(dbuf, xname);$	
INPUT	<i>dbuf</i>	$N \times 1$ 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>	$L \times M$ matrix or string, the item read from <i>dbuf</i> .
REMARKS	<b>vread</b> , unlike <b>vget</b> , does not change the contents of <i>dbuf</i> . Reading <i>x</i> from <i>dbuf</i> does not remove it from <i>dbuf</i> .	
SOURCE	<code>vpack.src</code>	
SEE ALSO	<b>vget</b> , <b>vlist</b> , <b>vput</b>	

## vtypecv

PURPOSE	Returns the types of the elements of a data buffer constructed with <b>vput</b> .
---------	---

---

**FORMAT**    *cv* = **vtypecv**(*dbuf*);  
**INPUT**     *dbuf*        N×1 vector, a data buffer containing various strings and matrices.  
**OUTPUT**    *cv*                K×1 character vector containing the types of the elements of *dbuf*.  
**SEE ALSO**    **vget**, **vput**, **vread**, **vnamecv**

## 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**

v

## walkindex

**PURPOSE**    Walks the index of an array forward or backward through a specified dimension.  
**FORMAT**     *ni* = **walkindex**(*i,o,dim*);

## window

---

INPUT	<i>i</i>	M×1 vector of indices into an array, where M≤N.
	<i>o</i>	N×1 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>	M×1 vector of indices, the new index.
REMARKS	<b>walkindex</b> will return a scalar error code if the index cannot walk further in the specified dimension and direction.	
EXAMPLE	<pre>orders = (3,4,5,6,7); a = arrayinit(orders,1); ind = { 2,3,3 }; ind = walkindex(ind,orders,-2);</pre>	

```
      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.	
LIBRARY	pgraph	
FORMAT	<b>window</b> ( <i>row,col,typ</i> );	
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	<p>The graphic panels will be numbered from 1 to (<i>row</i>)×(<i>col</i>) starting from the left topmost graphic panel and moving right.</p> <p>See <b>makewind</b> for creating graphic panels of a specific size and position. (For more information, see GRAPHIC PANELS, Section 24.3.</p>	
SOURCE	pwindow.src	
SEE ALSO	<b>endwind, begwind, setwind, nextwind, getwind, makewind</b>	

W

writer

PURPOSE	Writes a matrix to a <b>GAUSS</b> data set.	
FORMAT	<i>y</i> = <b>writer</b> ( <i>fh,x</i> );	
INPUT	<i>fh</i>	handle of the file that data is to be written to.

## writer

---

	$x$	$N \times K$ matrix.
OUTPUT	$y$	scalar specifying the number of rows of data actually written to the data set.
REMARKS	<p>The file must have been opened with <b>create</b>, <b>open for append</b>, or <b>open for update</b>.</p> <p>The data in <math>x</math> will be written to the data set whose handle is <math>fh</math> starting at the current pointer position in the file. The pointer position in the file will be updated, so the next call to <b>writer</b> will put the next block of data after the first block. (See <b>open</b> and <b>create</b> for the initial pointer positions in the file for reading and writing.)</p> <p><math>x</math> must have the same number of columns as the data set. <b>colsf</b> returns the number of columns in a data set.</p> <p><b>writer</b> returns the number of rows actually written to the data set. If <math>y</math> does not equal <b>rows</b>(<math>x</math>), the disk is probably full.</p> <p>If the data set is not double precision, the data will be rounded as it is written out.</p> <p>If the data contain character elements, the file must be double precision or the character information will be lost.</p> <p>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 <b>miss</b> function:</p> $x = \text{miss}(x, -32768);$ <p>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 <b>create</b> for details on creating a complex data set.)</p>	
EXAMPLE	<pre>create fp = data with x,10,8;</pre>	



---

```

if fp =\,= -1;
    errorlog "Can't create output file";
end;
endif;
c = 0;
do until c >= 10000;
    y = rndn(100,10);
    k = writer(fp,y);
    if k /= rows(y);
        errorlog "Disk Full";
        fp = close(fp);
        end;
    endif;
    c = c+k;
end;
fp = close(fp);

```

In this example, a 10000×10 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**

w

xlabel

PURPOSE Sets a label for the X axis.

LIBRARY `pgraph`

FORMAT **xlabel**(*str*);

INPUT *str* string, the label for the X axis.

SOURCE `pgraph.src`

## xlsGetSheetCount

---

SEE ALSO    **title, ylabel, xlabel**

### xlsGetSheetCount

PURPOSE    Gets the number of sheets in an Excel<sup>®</sup> spreadsheet.

FORMAT    *nsheets* = **xlsGetSheetCount**(*file*);

INPUT    *file*            string, name of .xls file.

OUTPUT    *nsheets*       scalar, sheet count or an error code.

PORTABILITY    **Windows** only

REMARKS    If **xlsGetSheetCount** fails, it will return a scalar error code, which can be decoded with **scalerr**.

SEE ALSO    **xlsGetSheetSize, xlsGetSheetTypes, xlsMakeRange**

### xlsGetSheetSize

PURPOSE    Gets the size (rows and columns) of a specified sheet in an Excel<sup>®</sup> spreadsheet.

FORMAT    { *rows, cols* } = **xlsGetSheetSize**(*file, sheet*);

INPUT    *file*            string, name of .xls file.  
          *sheet*       scalar, sheet index (1-based).

OUTPUT    *rows*           scalar, number of rows.

*cols*      scalar, number of columns.

PORTABILITY    **Windows** only

REMARKS      If **xlsGetSheetSize** fails, it will return a scalar error code, which can be decoded with **scalerr**.

SEE ALSO      **xlsGetSheetCount**, **xlsGetSheetTypes**, **xlsMakeRange**

## xlsGetSheetTypes

PURPOSE      Gets the cell format types of a row in an Excel<sup>®</sup> spreadsheet.

FORMAT      *nsheets* = **xlsGetSheetTypes**(*file*,*sheet*,*row*);

INPUT      *file*          string, name of .xls file.  
              *sheet*       scalar, sheet index (1-based).  
              *row*          scalar, the row of cells to be scanned.

OUTPUT      *types*          1×K 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** only

REMARKS      K is the number of columns found in the spreadsheet.

If **xlsGetSheetTypes** fails, it will return a scalar error code, which can be decoded with **scalerr**.

**X**

## xlsMakeRange

---

SEE ALSO    **xlsGetSheetCount**, **xlsGetSheetSize**, **xlsMakeRange**

### xlsMakeRange

PURPOSE    Builds an Excel<sup>®</sup> range string from a row/column pair.

FORMAT    *range* = **xlsMakeRange**(*row*, *col*);

INPUT      *row*            scalar or 2×1 vector.

*col*            scalar or 2×1 vector.

OUTPUT    *range*            string, an Excel<sup>®</sup>-formatted range specifier.

PORTABILITY    **Windows** only

REMARKS    If *row* is a 2×1 vector, it is interpreted as follows

*row*[1]    starting row

*row*[2]    ending row

            If *col* is a 2×1 vector, it is interpreted as follows

*col*[1]    starting column

*col*[2]    ending column

SEE ALSO    **xlsGetSheetCount**, **xlsGetSheetSize**, **xlsGetSheetTypes**

PURPOSE Reads from an Excel<sup>®</sup> spreadsheet into a **GAUSS** matrix.

FORMAT *mat* = **xlsreadm**(*file*,*range*,*sheet*,*vls*);

INPUT *file* string, name of .xls file.

*range* string, range to read, e.g. **a2:b20**, or the starting point of the read, e.g. **a2**.

*sheet* scalar, sheet number.

*vls* null string or 9×1 matrix, specifies the conversion of Excel<sup>®</sup> 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 *mat* matrix or a Microsoft error code.

PORTABILITY **Windows** only

REMARKS If range is a null string, then by default the read will begin at cell **a1**.

The *vls* argument lets users control the import of Excel<sup>®</sup> empty cells and special types, according to the following table:

## xlsreadsa

---

Row Number	Excel <sup>®</sup> 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 9999.99, and all other empty cells and special types to **GAUSS** missing values:

```
vls = reshape(error(0),9,1);  
  
vls[4] = 9999.99;
```

SEE ALSO **xlsReadSA**, **xlsWrite**, **xlsWriteM**, **xlsWriteSA**, **xlsGetSheetCount**, **xlsGetSheetSize**, **xlsGetSheetTypes**, **xlsMakeRange**

## xlsreadsa

**PURPOSE** Reads from an Excel<sup>®</sup> spreadsheet into a **GAUSS** string array or string.

**FORMAT** *s* = **xlsreadsa**(*file*,*range*,*sheet*,*vls*);

<b>INPUT</b>	<i>file</i>	string, name of .xls file.
	<i>range</i>	string, range to read, e.g. <b>a2:b20</b> or the starting point of the read, e.g. <b>a2</b> .
	<i>sheet</i>	scalar, sheet number.

*vls* null string or 9×1 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.

OUTPUT *s* string array or string or a Microsoft error code.

PORTABILITY **Windows** only

REMARKS If *range* is a null string, then by default the read will begin at cell **a1**.

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 “Division by Zero”, and all other empty cells and special types to null strings:

```
vls = reshape("",9,1);  
vls[4] = "Division by Zero";
```

SEE ALSO **xlsReadM, xlsWrite, xlsWriteM, xlsWriteSA, xlsGetSheetCount, xlsGetSheetSize, xlsGetSheetTypes, xlsMakeRange**

### xlsWrite

**PURPOSE** Writes a **GAUSS** matrix, string, or string array to an Excel<sup>®</sup> spreadsheet.

**FORMAT** *ret* = **xlsWrite**(*data*,*file*,*range*,*sheet*,*vls*);

**INPUT**

<i>data</i>	matrix, string, or string array.
<i>file</i>	string, name of .xls file.
<i>range</i>	string, the starting point of the write, e.g. <b>a2</b> .
<i>sheet</i>	scalar, sheet number.
<i>vls</i>	null string or 9×1 matrix or string array, specifies the conversion of <b>GAUSS</b> values or characters into Excel <sup>®</sup> empty cells and special types (see Remarks). A null string results in all <b>GAUSS</b> missing values and null strings being converted to empty cells.

**OUTPUT** *ret* scalar, 0 if success or a Microsoft error code.

**PORTABILITY** **Windows** only

**REMARKS** The *vls* argument lets users control the export to Excel<sup>®</sup> empty cells and special types, according to the following table:

Row Number	Excel <sup>®</sup> 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<sup>®</sup> and convert all **GAUSS** missing values to empty cells in Excel<sup>®</sup>:

```
vls = reshape(error(0),9,1);

vls[4] = 9999.99;
```

SEE ALSO **xlsReadSA, xlsReadm, xlsWriteM, xlsWriteSA, xlsGetSheetCount, xlsGetSheetSize, xlsGetSheetTypes, xlsMakeRange**

xlsWritem

**PURPOSE**     Writes a **GAUSS** matrix to an Excel<sup>®</sup> spreadsheet.

**FORMAT**     *ret* = **xlsWritem**(*data,file,range,sheet,vls*);

**INPUT**

<i>data</i>	matrix.
<i>file</i>	string, name of .xls file.
<i>range</i>	string, the starting point of the write, e.g. <b>a2</b> .
<i>sheet</i>	scalar, sheet number.
<i>vls</i>	null string or 9×1 matrix, specifies the conversion of <b>GAUSS</b> values into Excel <sup>®</sup> empty cells and special types (see Remarks). A null string results in all <b>GAUSS</b> missing values being converted to empty cells.

**OUTPUT**     *ret*            scalar, 0 if success or a Microsoft error code.

**PORTABILITY**     **Windows** only

**REMARKS**     The *vls* argument lets users control the export to Excel<sup>®</sup> empty cells and special types, according to the following table:

## xlswritesa

---

Row Number	Excel <sup>®</sup> 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<sup>®</sup> and convert all **GAUSS** missing values to empty cells in Excel<sup>®</sup>:

```
vls = reshape(error(0),9,1);
```

```
vls[4] = 9999.99;
```

SEE ALSO **xlsReadSA**, **xlsReadm**, **xlsWrite**, **xlsWriteSA**, **xlsGetSheetCount**, **xlsGetSheetSize**, **xlsGetSheetTypes**, **xlsMakeRange**

## xlswritesa

**PURPOSE** Writes a **GAUSS** string or string array to an Excel<sup>®</sup> spreadsheet.

**FORMAT** *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. <b>a2</b> .
<i>sheet</i>	scalar, sheet number.

*vls* null string or 9×1 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.

OUTPUT *ret* scalar, 0 if success or a Microsoft error code.

PORTABILITY **Windows** only

REMARKS 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";
```

SEE ALSO **xlsReadM, xlsWrite, xlsWriteM, xlsReadSA, xlsGetSheetCount, xlsGetSheetSize, xlsGetSheetTypes, xlsMakeRange**



## xpnd

---

**PURPOSE**     Expands a column vector into a symmetric matrix.

**FORMAT**      $x = \mathbf{xpnd}(v);$

**INPUT**        $v$               $K \times 1$  vector, to be expanded into a symmetric matrix.

**OUTPUT**       $x$               $M \times M$  matrix, the results of taking  $v$  and filling in a symmetric matrix with its elements.

$$M = ((-1 + \sqrt{1+8*K}))/2)$$

**REMARKS**     If  $v$  does not contain the right number of elements, (that is, if  $\sqrt{1 + 8*K}$  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**      $x = \{ \begin{array}{l} 1, \\ 2, 3, \\ 4, 5, 6, \\ 7, 8, 9, 10 \end{array} \};$   
 $y = \mathbf{xpnd}(x);$

$x = \begin{array}{c} 1 \\ 2 \\ 3 \\ 4 \\ 5 \\ 6 \\ 7 \\ 8 \\ 9 \\ 10 \end{array}$

$$y = \begin{matrix} & 1 & 2 & 4 & 7 \\ & 2 & 3 & 5 & 8 \\ 4 & 5 & 6 & 9 \\ & 7 & 8 & 9 & 10 \end{matrix}$$

SEE ALSO    **vech**

## xtics

**PURPOSE**    Sets and fixes scaling, axes numbering and tick marks for the X axis.

**LIBRARY**    **pgraph**

**FORMAT**    **xtics**(*min,max,step,minordiv*);

**INPUT**    *min*            scalar, the minimum value.  
               *max*            scalar, the maximum value.  
               *step*           scalar, the value between major tick marks.  
               *minordiv*   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 **xtics** 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**

SEE ALSO    **scale**, **ytics**, **ztics**

**X**

## xyz

---

### xy

PURPOSE	Graphs X vs. Y using Cartesian coordinates.	
LIBRARY	pgraph	
FORMAT	<b>xy</b> ( <i>x</i> , <i>y</i> );	
INPUT	<i>x</i>	N×1 or N×M matrix. Each column contains the X values for a particular line.
	<i>y</i>	N×1 or N×M 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	
SEE ALSO	<b>xyz</b> , <b>logx</b> , <b>logy</b> , <b>loglog</b>	

### xyz

PURPOSE	Graphs X vs. Y vs. Z using Cartesian coordinates.	
LIBRARY	pgraph	
FORMAT	<b>xyz</b> ( <i>x</i> , <i>y</i> , <i>z</i> );	
INPUT	<i>x</i>	N×1 or N×K matrix. Each column contains the X values for a particular line.

	<i>y</i>	N×1 or N×K matrix. Each column contains the Y values for a particular line.
	<i>z</i>	N×1 or N×K 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.	
SOURCE	pxyz.src	

ylabel

PURPOSE	Sets a label for the Y axis.	
LIBRARY	pgraph	
FORMAT	<b>ylabel</b> ( <i>str</i> );	
INPUT	<i>str</i>	string, the label for the Y axis.
SOURCE	pgraph.src	
SEE ALSO	<b>title</b> , <b>xlabel</b> , <b>ylabel</b>	

x

ytics

PURPOSE	Sets and fixes scaling, axes numbering and tick marks for the Y axis.	
LIBRARY	pgraph	

## zeros

---

FORMAT    **yticks**(*min*,*max*,*step*,*minordiv*);

INPUT    *min*        scalar, the minimum value.  
          *max*        scalar, the maximum value.  
          *step*       scalar, the value between major tick marks.  
          *minordiv*   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 **yticks** 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 **xticks** and **yticks** functions.

SOURCE   *pscale.src*

SEE ALSO   **scale**, **xticks**, **zticks**

## zeros

PURPOSE   Creates a matrix of zeros.

FORMAT    *y* = **zeros**(*r*,*c*);

INPUT    *r*            scalar, the number of rows.  
          *c*            scalar, the number of columns.

OUTPUT    *y*            *r*×*c* matrix of zeros.

REMARKS   This is faster than **ones**.



Noninteger arguments will be truncated to an integer.

EXAMPLE    `y = zeros(3,2);`

```
          0.000000  0.000000
y = 0.000000  0.000000
      0.000000  0.000000
```

SEE ALSO    `ones`, `eye`

zlabel

PURPOSE    Sets a label for the *Z* axis.

LIBRARY    `pgraph`

FORMAT    `zlabel(str);`

INPUT    *str*            string, the label for the *Z* axis.

SOURCE    `pgraph.src`

SEE ALSO    `title`, `xlabel`, `ylabel`

**Z**

ztics

## ztics

---

PURPOSE	Sets and fixes scaling, axes numbering and tick marks for the Z axis.	
LIBRARY	pgraph	
FORMAT	<b>ztics</b> ( <i>min</i> , <i>max</i> , <i>step</i> , <i>minordiv</i> );	
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 <b>contour</b> , 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 <b>graphset</b> is called.  This gives you direct control over the axes endpoints and tick marks. If <b>ztics</b> is called after a call to <b>scale3d</b> , it will override <b>scale3d</b> .	
SOURCE	pscale.src	
SEE ALSO	<b>scale3d</b> , <b>xtics</b> , <b>ytics</b> , <b>contour</b>	

# Obsolete Commands D

The following commands will no longer be supported and therefore should not be used when creating new programs.

<b>color</b>	<b>eigch</b>
<b>coreleft</b>	<b>eigch2</b>
<b>csrtype</b>	<b>eigr</b>
<b>denseSubmat</b>	<b>eigr2</b>
<b>dfree</b>	<b>eigrs</b>
<b>disable</b>	<b>eigrs2</b>
<b>editm</b>	<b>enable</b>
<b>eigcg</b>	<b>export</b>
<b>eigcg2</b>	

<b>exportf</b>	<b>ndpclex</b>
<b>files</b>	<b>ndpcntrl</b>
<b>font</b>	<b>plot</b>
<b>FontLoad</b>	<b>plotsym</b>
<b>FontUnload</b>	<b>prcsn</b>
<b>FontUnloadAll</b>	<b>print on/off</b>
<b>graph</b>	<b>rndns</b>
<b>import</b>	<b>rndus</b>
<b>importf</b>	<b>scroll</b>
<b>isSparse</b>	<b>setvmode</b>
<b>line</b>	<b>sparseCols</b>
<b>lpos</b>	<b>sparseEye</b>
<b>lprint</b>	<b>sparseFD</b>
<b>lprint on/off</b>	<b>sparseFP</b>
<b>lpwidth</b>	<b>sparseHConcat</b>
<b>lshow</b>	<b>sparseNZE</b>
<b>medit</b>	<b>sparseOnes</b>
<b>nametype</b>	<b>sparseRows</b>
<b>ndpchck</b>	

---

<code>sparseScale</code>	<code>WinGetColorCells</code>
<code>sparseSet</code>	<code>WinGetCursor</code>
<code>sparseSolve</code>	<code>WinMove</code>
<code>sparseSubmat</code>	<code>WinOpenPQG</code>
<code>sparseTD</code>	<code>WinOpenText</code>
<code>sparseTranspose</code>	<code>WinOpenTTY</code>
<code>sparseTrTD</code>	<code>WinPan</code>
<code>sparseTScalar</code>	<code>WinPrint</code>
<code>sparseVConcat</code>	<code>WinPrintPQG</code>
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<code>spline2d</code>	<code>WinRefreshArea</code>
<code>vartype</code>	<code>WinResize</code>
<code>WinClear</code>	<code>WinSetActive</code>
<code>WinClearArea</code>	<code>WinSetBackground</code>
<code>WinClearTTYlog</code>	<code>WinSetColorCells</code>
<code>WinClose</code>	<code>WinSetColormap</code>
<code>WinCloseAll</code>	<code>WinSetCursor</code>
<code>WinGetActive</code>	<code>WinSetForeground</code>
<code>WinGetAttributes</code>	

**WinSetRefresh**

**WinZoomPQG**

**WinSetTextWrap**

# Colors E

Colors

0	Black	8	Dark Grey
1	Blue	9	Light Blue
2	Green	10	Light Green
3	Cyan	11	Light Cyan
4	Red	12	Light Red
5	Magenta	13	Light Magenta
6	Brown	14	Yellow
7	Grey	15	White





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