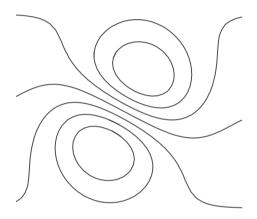
# Calerga



# Sysquake for LaTeX User Manual

Embedding numerical results and graphics in LaTeX

January 2017

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Sysquake for LaTeX User Manual, January 2017. Calerga Sàrl, La Tour-de-Peilz, Switzerland.

Most of the material in Sysquake for LaTeX User Manual has first been written as a set of XHTML files, with lots of cross-reference links. Since (X)HTML is not very well suited for printing, it has been converted to LaTeX with the help of a home-made conversion utility. Additional XML tags have been used to benefit from LaTeX features: e.g. raster images have been replaced with EPS images, equations have been converted from text to real mathematic notation, and a table of contents and an index have been added. The same method has been used to create the material for the help command. Thanks to the make utility, the whole process is completely automatic. This system has proved to be very flexible to maintain three useful formats in parallel: two for on-line help, and one for high-quality printing.

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## **Chapter 1**

# Introduction

LaTeX is a typesetting environment for generating high-quality documents, for reading on-screen (typically as PDF) or for printing. Developed by Leslie Lamport with the support of many other contributors and the American Mathematics Society (AMS), LaTeX is extremely popular for scientific publications. LaTeX is a set of macros for TeX. From TeX, it inherits the high quality of text layout and equation formatting, and implementations on most computer platforms; it adds facilities for sections, page layout, table of contents, index, etc.

Like TeX, LaTeX takes as input a text file which contains the text to typeset and commands introduced with an escape character. To create a document, the author typically writes this file in a text editor, processes it with latex (a command-line tool), displays the result in a viewer, and continues with more iterations until a satisfactory result is obtained. Numerical results and graphics come usually from other applications such as Sysquake.

Sysquake for LaTeX is a LaTeX package and a command tool which let you embed Sysquake commands directly in LaTeX source files. With it, no need to copy and format numerical results or generate EPS file by hand; the package takes care of cumbersome tasks, letting you focus on the creative work. Sysquake for LaTeX uses the same language and libraries as Sysquake and other products of Calerga.

## **Chapter 2**

# Installing Sysquake for LaTeX

Installing Sysquake for LaTeX consists in three steps:

- Copy all Sysquake for LaTeX files to your hard disk. The default location is /opt/SysquakeLx on unix systems, and C:\Program Files\SysquakeForLaTeX on Windows.
- Update environment variables so that sysquakelatextool is found by the operating system and finds library files. This consists in adding the path of the directory containing sysquakelatextool to PATH, and setting a new environment variable SYSQUAKELXPATH to the semicolon-separated list of paths to library directories.
- Make your TeX distribution able to load the sysquake.sty package. In TDS-based distributions (TeX Live, teTex, MiKTeX, etc.), this means adding the path of TexFiles to the list of TDS roots.

LaTeX must also be able to execute external programs with \write18. Depending on the platform, this may require to add an option argument to pdflatex, such as -shell-escape or -enable-write18.

**Caution:** enabling external programs can be dangerous if you typeset tex files from dubious origin. Depending on your needs, you may prefer a separate configuration to typeset tex files which use Sysquake for LaTeX, keeping the standard, more secure configuration for other files. Sysquake for LaTeX should be used only with tex files you trust.

#### 2.1 Windows

#### **MiKTeX**

Install the latest release of MiKTeX (e.g. MiKTeX 2.9.6161). Then run the installer of Sysquake for LaTeX and follow the instructions.

The last steps, as explained in the installer, are manual:

- Make your TeX distribution aware of Sysquake for LaTeX. You should add a new root directory to the list of TDS directories. In the Start menu, submenu MiKTeX 2.9 (or equivalent), MiKTeX Settings (or MiKTeX Settings (Admin)) and select the Roots tab. Click the Add button, select directory TexFiles of Sysquake for LaTeX, and confirm, clicking the main OK button of the MiKTeX Options window.
- Configure your LaTeX editor to run pdflatex.exe with the argument -shell-escape. In TeXworks, for instance, in menu Edit>Preferences, Typesetting tab, double-click pdfLaTeX+MakeIndex+BibTeX in the Processing tool list, click the "+" button, type -tex-option=-shell-escape, then click the up arrow button to move the new option before \$fullname\$.

#### TeX Live

Install the latest release of TeX Live (e.g. TeX Live 2016), then run Sysquake for LaTeX installer and follow the instructions.

The last steps are manual:

- Make sure sysquakelatextool.exe is in the search path. Open the Command Prompt application (e.g. by entering cmd in menu Start>Run), and type sysquakelatextool. You should read a few lines beginning with a percent. If you get an error message instead (e.g. 'sysquakelatextool' is not recognized...), you should add its path to the PATH environment variable. Select menu Start>Control Panel, then System, Advanced tab, Environment Variables button, double-click the User variable PATH and add sysquakelatextool.exe's path followed by a semicolon at the beginning of what already exists. E.g.

C:\Program Files\SysquakeForLaTeX\bin\Win32;...

- Configure your LaTeX editor to run pdflatex.exe with the argument -shell-escape. In TeXworks, for instance, in menu Edit>Preferences, Typesetting tab, double-click pdfLaTeX+MakeIndex+BibTeX in the Processing tool list, click the "+" button, type -tex-option=-shell-escape, then click the up arrow button to move the new option before \$fullname\$. Install 11

Edit the local texmf.cnf (typically in C:\texlive\2011; you can find it by running kpsewhich.exe texmf.conf in the Command Prompt). If there is already a line starting with TEXMFLOCAL, add a comma followed by the path of TexFiles in SysquakeForLaTeX, without any quote even if there are spaces; otherwise, add the following line (specify the correct path if you have chosen a different install directory):

TEXMFLOCAL = \$SELFAUTOPARENT/../texmf-local,C:\Program Files\SysguakeFo

#### Other distributions

For other TeX distributions, please read their documentation. Sysquake for LaTeX installer is totally independent from any other software; you should just configure TeX so that it finds the file sysquake.sty and can run sysquakelatextool.exe.

#### 2.2 macOS

On macOS, Sysquake for LaTeX is distributed as a package. To install it, just double-click its icon and accept everything. You will need an administrator password. An alias of the documentation entry file is put on the Desktop.

The macOS package stores the distribution in /opt/SysquakeLx and creates the following symbolic links, also creating directories if they do not already exist:

You should make sure to enable the execution of external programs with \write18. In TeXShop, for instance, open TeXShop preferences (menu TeXShop>Preferences), display the Engine tab, then in pdfTeX/Latex field, replace pdflatex -file-line-error -synctex=1 with pdflatex -file-line-error -synctex=1 -shell-escape.

#### 2.3 Ubuntu, Debian or Linux Mint

Sysquake for LaTeX is available as a deb package, with a dependency on the texlive package which will also be installed if it isn't already done. Double-clicking its icon should open the default package management utility for your system. You can then read its description and install it.

Alternatively, in a terminal window, you can type the following command (replace the star and specify the exact filename if you have several versions):

```
sudo dpkg -i sysquakelx-*.deb
```

Should you want to remove Sysquake for LaTeX, you can type the following command:

```
sudo dpkg -r sysquakelx
```

#### 2.4 Manual installation on Linux

Sysquake for LaTeX is also available as a tgz archive, which can run on other Linux distributions and gives you total flexibility for file location. The steps below give you general directions.

 Install the latest release of TeX Live (e.g. TeX Live 2016). On a Linux distribution using rpm packages like RedHat, Fedora or CentOS, you can type

```
sudo yum install texlive tex-live-latex
```

On a Linux distribution using deb packages like Debian, Ubuntu or Linux Mint, you can install TeX Live with the package manager:

```
sudo apt-get install -y texlive-latex-base
```

 Unarchive the distribution of Sysquake for LaTeX for Linux to a suitable location, such as /opt/SysquakeLx:

```
tar xfz sysquakelx-*.tgz
sudo mkdir -p /opt
sudo cp -R SysquakeLx /opt
```

- Have bin/i386-linux/sysquakelatextool in your PATH. There are two ways to do it: put a symbolic link to bin/i386-linux/sysquakelatextool in one of your existing program directories, such as /usr/bin or /usr/local/bin:

```
sudo mkdir -p /usr/local/bin
sudo ln -s /opt/SysquakeLx/bin/i386-linux/sysquakelatextool /usr/local/bi
```

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or add /opt/SysquakeLx/SysquakeLx/bin/i386-linux to your environment variable PATH.

- Create a new environment variable named SYSQUAKELXPATH which contains a list of semicolon-separated paths for the directories containing .lml libraries. The default value is .;./Lib;/opt/SysquakeLx/Lib, which finds standard libraries if Sysquake for LaTeX is installed in /opt/SysquakeLx.
- Check where local TeX files are stored:

kpsewhich -var-value=TEXMFLOCAL

 Add there a symbolic link to the TexFiles directory in SysquakeLx. Assuming you've stored SysquakeLx in /opt and the directory of local TeX files given in the previous step is /usr/local/share/texmf, type

ln -s /opt/SysquakeLx/TexFiles /usr/local/share/texmf/SysquakeLx

Rebuild the TeX file list:

texhash

## **Chapter 3**

# Sysquake for LaTeX Tutorial

Once installed, Sysquake for LaTeX is a LaTeX package which you can use as easily as any other package. In this tutorial, we will create a technical report which analyzes some of the properties of a rational function.

Sysquake for LaTeX works at the level of the TeX program, which, with LaTeX definitions, is usually called latex for creating DVI files, or pdflatex for creating PDF files. You can also choose among different applications which hide the details of running latex and displaying the result, such as emacs on unix platforms, TeXShop on macOS, or Texmaker on Windows and other platforms. Sysquake for LaTeX is perfectly compatible with them; actually it just ignores them. So you can keep your favorite editing environment.

And now, let us begin!

*Remark:* the complete source file of another example comes with Sysquake for LaTeX.

# 3.1 Creating a LaTeX file

Use your usual method to create a LaTeX file and write a few things to check that everything runs correctly. For this tutorial, we will make a report. The LaTeX file, named for example sq-tutorial.tex, contains something like

```
\documentclass[11pt]{report}
\begin{document}

\title{Sysquake for \LaTeX{} tutorial}
\author{Calerga}
```

```
\date{13 August 2007}
\maketitle
\end{document}
```

# 3.2 Add the Sysquake package

Packages are sets of definitions which add new commands to LaTeX. They are listed in the preamble of LaTeX files, between \documentclass and \begin{document}. The Sysquake package is added like any other package:

```
\documentclass[11pt]{report}
\usepackage{sysquake}
\begin{document}
...
```

If there is an error when you typeset this file, the file sysquake.sty is probably not found. Please refer to section Sysquake for LaTeX Installation for more informations.

Assuming there is no error, we can now use LaTeX commands for invoking Sysquake.

## 3.3 Your first Sysquake computation

Since we are anxious to get a result, let us begin with something really simple. We will insert the result of 5+2 as an equation. Add the following line between \maketitle and \end{document}:

```
We all know that $5+2=\sqexpr{5+2}$, don't we?
```

Running LaTeX twice will produce the result you expect. Why twice? Like other features of LaTeX, the Sysquake package needs two phases to do its magic. During the first phase, commands create a file named sq-tutorial.lme (the name is built after the .tex file) which contains Sysquake code, or more exactly code written in the programming language of Sysquake, LME. During the second phase, loading the sysquake package runs the command-line tool sysquakelatextool to interpret this file. The result is written to a file named sq-tutorial.lmeout and, if there are graphics, to EPS files. This file contains the result as LaTeX definitions; for instance, \sqexpr{5+2} above creates the following definition:

```
\def\lmefragi{%
7}
```

Here, the name is \lmefrag followed by the expression number in lowercase roman notation (this might change in future versions). Sysquake commands, in addition to producing sq-tutorial.lme, also invoke the corresponding definition to insert the result in the final typeset document.

Note that typesetting documents is often already performed by running latex twice, or even more times, to resolve cross-references and update the table of contents and the index. So there is nothing new.

#### 3.4 Formatted results

Our first expression could have been computed in plain LaTeX, which gives access to the computation capabilities of TeX. So our next example will use an advanced LME function, magic, which produces a magic square in a matrix. Matrices are displayed in mathematical notation.

Since matrices are too large to fit nicely in a paragraph, we put it in an equation:

```
Magic square $M_3$ of order 3 is
\begin{equation}
M_3 = \sqexpr{magic(3)}
\end{equation}
```

#### 3.5 Programs

\sqexpr is very convenient for simple expressions. However, it is not suited to more complicated programs, for two reasons: first, only a single expression can be evaluated which gives a single result. Second, LME syntax may interfer with LaTeX syntax: the backslash, for instance, is a normal character used for premultiplying with the inverse of a matrix in LME, while it is an escape character in LaTeX. We can always escape special characters in LaTeX, but Sysquake for LaTeX's aim is to make things simpler, not overly complicated.

The answer to \sqexpr's limitations is the sysquake environment which can contain any kind of code. Let us write another magic square with it:

```
Magic square $M_4$ of order 4 is
\begin{equation}
M_4 =
\begin{sysquake}
magic(4)
\end{sysquake}
\end{equation}
```

The sysquake environment can contain any number of lines with any code, unescaped, like verbatim. The only invalid character sequence is \end{sysquake}, which marks the end of Sysquake code. If for an unlikely reason you have to write Sysquake code which contains this sequence of characters, add a space somewhere, or if it is in a string, write a character with an escape sequence, e.g. \end{sysquak\145}.

Code in sysquake environment does not have the restrictions of expressions written with \sqexpr. You can define variables, have conditional executions and loops, use libraries, and more.

Contrary to \sqexpr where a single result is formatted for LaTeX, the sysquake environment only inserts any text produced by the code it contains in the LaTeX source. This text is still processed by LaTeX. To format it, you can either do it yourself, inserting LaTeX commands, or use the Sysquake function sqlxvalue. In the code fragment above, magic(4) without semicolon writes the result as an assignment to variable ans, as raw text:

```
ans =
            2
                   3
                         13
    16
     5
           11
                  10
                          8
     9
            7
                  6
                         12
           14
                  15
                          1
```

In a LaTeX equation, this does not look like a matrix. With sqlxvalue(magic(4)), we obtain

```
\ensuremath{\left[ \begin{array}{cccc}16
&2
&3
&13\\
5
&11
&10
&8\\
9
&7
&6
&12\\
4
&14
&15
&1\end{array}\right]}
```

which is rendered more nicely, like with \sqexpr.

Short code fragments, such as variable assignments, can be included with the \sqeval command. Note that its argument is processed by LaTeX, contrary to the contents of sysquake environment.

## 3.6 Graphics

Creating graphics for LaTeX documents is among the most cumbersome tasks an author is facing. Sysquake for LaTeX makes it as easy as in Sysquake itself.

Graphics are created by adding the size of the EPS image to be created after \begin{sysquake}. The result is inserted automatically in a picture environment of the same size. Package graphicx is used, as well as package epstopdf in PDFLaTeX; they should be imported explicitly with \usepackage:

```
\usepackage{graphicx}
\usepackage{epstopdf}
```

Here is an example of a function plot:

```
\begin{sysquake}(300,200) 
 a = 7.2; 
 fplot(@(x) (x+0.3)^2+a*exp(-3*x^2), [-2,3], 'r') 
 \end{sysquake}
```

In LaTeX, figures are usually floating. This is done with the figure environment:

```
\begin{figure}
\begin{center}
\begin{sysquake}(300,200)
a = 7.2;
fplot(@(x) (x+0.3)^2+a*exp(-3*x^2), [-2,3], 'r')
\end{sysquake}
\caption{Function plot}
\end{center}
\end{figure}
```

#### 3.7 Functions and libraries

Code fragments may rely on the large set of built-in operators and functions as well as on additional functions stored in libraries. Libraries are text files with a *.lml* suffix which contain the source code of sets of related functions. You can use those which come with Sysquake for LaTeX or write develop new ones, which you can share with Sysquake.

To make functions defined in a library available to Sysquake, a use statement should be placed before any call, in a sysquake environment or a \square command.

```
\begin{figure}
\begin{center}
\begin{sysquake}(300,200)
```

```
use polyhedra;
smallstellateddodecahedron;
colormap(blue2yellow2redcm);
plotoption nomargin;
plotoption noframe;
plotoption fill3d;
\end{sysquake}
\caption{Great dodecahedron}
\end{center}
\end{figure}
```

## **Chapter 4**

# **LME Tutorial**

This chapter introduces LME(TM) (Lightweight Math Engine), the interpreter for numeric computing used by Sysquake, and shows you how to perform basic computations. It supposes you can type commands to a command-line interface. You are invited to type the examples as you read this tutorial and to experiment on your own. For a more systematic description of LME, please consult the LME Reference chapter.

In the examples below, we assume that LME displays a prompt >. This is not the case for all applications. You should never type it yourself. Enter what follows the prompt on the same line, hit the Return key (or tap the Eval or Execute button), and observe the result.

#### 4.1 Simple operations

LME interprets what you type at the command prompt and displays the result unless you end the command with a semicolon. Simple expressions follow the syntactic rules of many programming languages.

```
> 2+3*4
ans =
14
> 2+3/4
ans =
2.75
```

As you can see, the evaluation order follows the usual rules which state that the multiplication (denoted with a star) and division (slash) have a higher priority than the addition and subtraction. You can change this order with parenthesis:

```
> (2+3)*4
ans =
20
```

The result of expressions is automatically assigned to variable ans (more about variables later), which you can reuse in the next expression:

```
> 3*ans
ans =
60
```

Power is represented by the ^ symbol:

```
> 2<sup>5</sup>
ans = 32
```

LME has many mathematical functions. Trigonometric functions assume that angles are expressed in radians, and sqrt denotes the square root.

```
> sin(pi/4) * sqrt(2)
ans =
1
```

# 4.2 Complex Numbers

In many computer languages, the square root is defined only for non-negative arguments. However, it is extremely useful to extend the set of numbers to remove this limitation. One defines i such that  $i^2 = -1$ , and applies all the usual algebraic rules. For instance,  $\sqrt{-1} = \sqrt{i^2} = i$ , and  $\sqrt{-4} = \sqrt{4}\sqrt{-1} = 2i$ . Complex numbers of the form a + bi are the sum of a real part a and an imaginary part b. It should be mentioned that i, the symbol used by mathematicians, is called j by engineers. LME accepts both symbols as input, but it always writes it j. You can use it like any function, or stick an i or j after a number:

```
> 2+3*j
ans =
2+3j
> 3j+2
ans =
2+3j
```

Many functions accept complex numbers as argument, and return a complex result when the input requires it even if it is real:

```
> sqrt(-2)
ans =
 0+1.4142i
> exp(3+2j)
```

```
ans =
  -8.3585+18.2637j
> log(-8.3585+18.2637j)
ans =
  3+2j
```

To get the real or imaginary part of a complex number, use the functions real or imag, respectively:

```
> real(2+3j)
ans =
  2
> imag(2+3j)
ans =
  3
```

Complex numbers can be seen as vectors in a plane. Then addition and subtraction of complex numbers correspond to the same operations applied to the vectors. The absolute value of a complex number, also called its magnitude, is the length of the vector:

```
> abs(3+4j)
ans =
5
> sqrt(3^2+4^2)
ans =
5
```

The argument of a complex number is the angle between the x axis ("real axis") and the vector, counterclockwise. It is calculated by the angle function.

```
> angle(2+3j)
ans =
  0.9828
```

The last function specific to complex numbers we will mention here is conj, which calculates the conjugate of a complex number. The conjugate is simply the original number where the sign of the imaginary part is changed.

```
> conj(2+3j)
ans =
  2-3i
```

Real numbers are also complex numbers, with a null imaginary part; hence

```
> abs(3)
ans =
```

```
3 > conj(3) ans = 3 > angle(3) ans = 0 > angle(-3) ans = 3.1416
```

#### 4.3 Vectors and Matrices

LME manipulates vectors and matrices as easily as scalars. To define a matrix, enclose its contents in square brackets and use commas to separate elements on the same row and semicolons to separate the rows themselves:

```
> [1,2;5,3]
ans =
1 2
5 3
```

Column vectors are matrices with one column, and row vectors are matrices with one row. You can also use the colon operator to build a row vector by specifying the start and end values, and optionally the step value. Note that the end value is included only if the range is a multiple of the step. Negative steps are allowed.

```
> 1:5

ans =

1 2 3 4 5

> 0:0.2:1

ans =

0 0.2 0.4 0.6 0.8 1

> 0:-0.3:1

ans =

0 -0.3 -0.6 -0.9
```

There are functions to create special matrices. The zeros, ones, rand, and randn functions create matrices full of zeros, ones, random numbers uniformly distributed between 0 and 1, and random numbers normally distributed with a mean of 0 and a standard deviation of 1, respectively. The eye function creates an identity matrix, i.e. a matrix with ones on the main diagonal and zeros elsewhere. All of these functions can take one scalar argument n to create a square n-by-n matrix, or two arguments m and n to create an m-by-n matrix.

```
> zeros(3)
ans =
0 0 0
0 0 0
0 0 0
> ones(2,3)
ans =
1 1 1
1 1 1
> rand(2)
ans =
0.1386 0.9274
0.3912 0.8219
> randn(2)
ans =
0.2931 1.2931
-2.3011 0.9841
> eye(3)
ans =
 1 0 0
0 1 0
0 0 1
> eye(2,3)
ans =
1 0 0
0 1 0
```

You can use most scalar functions with matrices; functions are applied to each element.

```
> sin([1;2])
ans =
0.8415
0.9093
```

There are also functions which are specific to matrices. For example, det calculates the determinant of a square matrix:

```
> det([1,2;5,3])
ans =
  -7
```

Arithmetic operations can also be applied to matrices, with their usual mathematical behavior. Additions and subtractions are performed on each element. The multiplication symbol  $\ast$  is used for the product of two matrices or a scalar and a matrix.

```
> [1,2;3,4] * [2;7]
ans =
16
34
```

The division symbol / denotes the multiplication by the inverse of the right argument (which must be a square matrix). To multiply by the inverse of the left argument, use the symbol \. This is handy to solve a set of linear equations. For example, to find the values of x and y such that x + 2y = 2 and 3x + 4y = 7, type

```
> [1,2;3,4] \ [2;7]
ans =
3
-0.5
```

Hence x=3 and y=-0.5. Another way to solve this problem is to use the inv function, which return the inverse of its argument. It is sometimes useful to multiply or divide matrices element-wise. The .\*, ./ and .\ operators do exactly that. Note that the + and - operators do not need special dot versions, because they perform element-wise anyway.

```
> [1,2;3,4] * [2,1;5,3]

ans =

12 7

26 15

> [1,2;3,4] .* [2,1;5,3]

ans =

2 2

15 12
```

Some functions change the order of elements. The transpose operator (tick) reverses the columns and the rows:

```
> [1,2;3,4;5,6]'
ans =
1 3 5
2 4 6
```

When applied to complex matrices, the complex conjugate transpose is obtained. Use dot-tick if you just want to reverse the rows and columns. The flipud function flips a matrix upside-down, and fliplr flips a matrix left-right.

```
> flipud([1,2;3,4])
ans =
   3 4
   1 2
> fliplr([1,2;3,4])
ans =
   2 1
   4 3
```

To sort the elements of each column of a matrix, or the elements of a row vector, use the sort function:

```
> sort([2,4,8,7,1,3])
ans =
1 2 3 4 7 8
```

To get the size of a matrix, you can use the size function, which gives you both the number of rows and the number of columns unless you specify which of them you want in the optional second argument:

```
> size(rand(13,17))
ans =
  13 17
> size(rand(13,17), 1)
ans =
  13
> size(rand(13,17), 2)
ans =
  17
```

# 4.4 Polynomials

LME handles mostly numeric values. Therefore, it cannot differentiate functions like  $f(x) = sin(e^x)$ . However, a class of functions has a paramount importance in numeric computing, the polynomials. Polynomials are weighted sums of powers of a variable, such as  $2x^2 + 3x - 5$ . LME stores the coefficients of polynomials in row vectors; i.e.  $2x^2 + 3x - 5$  is represented as [2,3,-5], and  $2x^5 + 3x$  as [2,0,0,0,3,0].

Adding two polynomials would be like adding the coefficient vectors if they had the same size; in the general case, however, you had better use the function addpol, which can also be used for subtraction:

```
> addpol([1,2],[3,7])
ans =
   4 9
> addpol([1,2],[2,4,5])
ans =
   2 5 7
> addpol([1,2],-[2,4,5])
ans =
   -2 -3 -3
```

Multiplication of polynomials corresponds to convolution (no need to understand what it means here) of the coefficient vectors.

```
> conv([1,2],[2,4,5])

ans =

2 8 13 10

Hence (x+2)(2x^2+4x+5) = 2x^3+8x^2+13x+10.
```

## 4.5 Strings

You type strings by delimiting them with single quotes:

```
> 'Hello, World!'
ans =
Hello, World!
```

If you want single quotes in a string, double them:

```
> 'Easy, isn''t it?'
ans =
  Easy, isn't it?
```

Some control characters have a special representation. For example, the line feed, used in LME as an end-of-line character, is \n:

```
> 'Hello,\nWorld!'
ans =
Hello,
World!
```

Strings are actually matrices of characters. You can use commas and semicolons to build larger strings:

```
> ['a','bc';'de','f']
ans =
  abc
  def
```

#### 4.6 Variables

You can store the result of an expression into what is called a variable. You can have as many variables as you want and the memory permits. Each variable has a name to retrieve the value it contains. You can change the value of a variable as often as you want.

```
> a = 3;
> a + 5
ans =
8
> a = 4;
> a + 5
ans =
```

Note that a command terminated by a semicolon does not display its result. To see the result, remove the semicolon, or use a comma if you have several commands on the same line. Implicit assignment to variable ans is not performed when you assign to another variable or when you just display the contents of a variable.

```
> a = 3
a =
3
> a = 7, b = 3 + 2 * a
a =
7
b =
17
```

## 4.7 Loops and Conditional Execution

To repeat the execution of some commands, you can use either a for/end block or a while/end block. With for, you use a variable as a counter:

```
> for i=1:3;i,end
i =
   1
i =
   2
i =
   3
```

With while, the commands are repeated as long as some expression is true:

```
> i = 1; while i < 10; i = 2 * i, end
i =
2
i =
4
i =
8</pre>
```

You can choose to execute some commands only if a condition holds true :

```
> if 2 < 3;'ok',else;'amazing...',end
ans =
  ok</pre>
```

#### 4.8 Functions

LME permits you to extend its set of functions with your own. This is convenient not only when you want to perform the same computation on different values, but also to make you code clearer by dividing the whole task in smaller blocks and giving names to them. To define a

new function, you have to write its code in a file; you cannot do it from the command line. In Sysquake, put them in a function block.

Functions begin with a header which specifies its name, its input arguments (parameters which are provided by the calling expression) and its output arguments (result of the function). The input and output arguments are optional. The function header is followed by the code which is executed when the function is called. This code can use arguments like any other variables.

We will first define a function without any argument, which just displays a magic square, the sum of each line, and the sum of each column:

```
function magicsum3
  magic_3 = magic(3)
  sum_of_each_line = sum(magic_3, 2)
  sum_of_each_column = sum(magic_3, 1)
```

You can call the function just by typing its name in the command line:

```
> magicsum3
magic_3 =
    8 1 6
    3 5 7
    4 9 2
sum_of_each_line =
    15
    15
    sum_of_each_column =
    15 15 15
```

This function is limited to a single size. For more generality, let us add an input argument:

```
function magicsum(n)
  magc = magic(n)
  sum_of_each_line = sum(magc, 2)
  sum_of_each_column = sum(magc, 1)
```

When you call this function, add an argument:

```
> magicsum(2)
magc =
    1 3
    4 2
sum_of_each_line =
    4
    6
sum_of_each_column =
    5 5
```

Note that since there is no 2-by-2 magic square, magic (2) gives something else... Finally, let us define a function which returns the sum of each line and the sum of each column:

```
function (sum_of_each_line, sum_of_each_column) = magicSum(n)
  magc = magic(n);
  sum_of_each_line = sum(magc, 2);
  sum_of_each_column = sum(magc, 1);
```

Since we can obtain the result by other means, we have added semicolons after each statement to suppress any output. Note the uppercase S in the function name: for LME, this function is different from the previous one. To retrieve the results, use the same syntax:

```
> (sl, sc) = magicSum(3)
sl =
   15
   15
   15
   sc =
   15  15  15
```

You do not have to retrieve all the output arguments. To get only the first one, just type

```
> sl = magicSum(3)
sl =
    15
    15
    15
```

When you retrieve only one output argument, you can use it directly in an expression:

```
> magicSum(3) + 3
ans =
   18
   18
   18
```

One of the important benefits of defining function is that the variables have a limited scope. Using a variable inside the function does not make it available from the outside; thus, you can use common names (such as x and y) without worrying about whether they are used in some other part of your whole program. For instance, let us use one of the variables of magicSum:

```
> magc = 77
magc =
77
```

```
> magicSum(3) + magc
ans =
    92
    92
    92
> magc
magc =
    77
```

#### 4.9 Local and Global Variables

When a value is assigned to a variable which has never been referenced, a new variable is created. It is visible only in the current context: the base workspace for assignments made from the command-line interface, or the current function invocation for functions. The variable is discarded when the function returns to its caller.

Variables can also be declared to be global, i.e. to survive the end of the function and to support sharing among several functions and the base workspace. Global variables are declared with keyword global:

```
global x
global y z
```

A global variable is unique if its name is unique, even if it is declared in several functions.

In the following example, we define functions which implement a queue which contains scalar numbers. The queue is stored in a global variable named QUEUE. Elements are added at the front of the vector with function queueput, and retrieved from the end of the vector with function queueget.

```
function queueput(x)
  global QUEUE;
  QUEUE = [x, QUEUE];

function x = queueget
  global QUEUE;
  x = QUEUE(end);
  QUEUE(end) = [];
```

Both functions must declare QUEUE as global; otherwise, the variable would be local, even if there exists also a global variable defined elsewhere. The first time a global variable is defined, its value is set to the empty matrix []. In our case, there is no need to initialized it to another value.

Here is how these functions can be used.

```
> queueput(1);
> queueget
ans =
    1
> queueput(123);
> queueput(2+3j);
> queueget
ans =
    123
> queueget
ans =
    2 + 3j
```

To observe the value of QUEUE from the command-line interface, QUEUE must be declared global there. If a local variable QUEUE already exists, it is discarded.

```
> global QUEUE
> QUEUE
QUEUE =
  []
> queueput(25);
> queueput(17);
> QUEUE
QUEUE =
  17 25
```

## **Chapter 5**

# LaTeX Package Reference

This section documents the LaTeX package sysquake.sty. This package permits to embed in LaTeX source files (.tex suffix) fragments of code written in LME, the programming language of Sysquake. All these fragments are evaluated successively in the same context: definitions and variables can be reused across multiple fragments.

## 5.1 Package options

The package is imported in the main LaTeX source file with \usepackage:

\usepackage{sysquake}

Several options can be specified in a comma-separated list in square brackets; for example

\usepackage[latexingraphics,noseparatelogfile]{sysquake}

Here is a list of all supported options.

latexingraphics By default, text in graphics is embedded in EPS files without further processing. In addition to axis scale labels, graphical commands which display explicit text are label, legend, text, and title.

With option latexingraphics, all the text which appears in graphics in handled by LaTeX. This has two benefits: first, text font and size mix much better with the document text, for a higher overall quality. Second, LaTeX constructs can appear anywhere in graphics, with greek letters, math symbols and equations. A small drawback is that some characters, such as \ and \$, have a special

meaning for LaTeX which make the code incompatible with other versions of Sysquake.

noexec Each time the document is typeset, Sysquake code fragments which have been extracted during the previous iteration are evaluated by a platform-dependant program, sysquakelatextool. This program produces output data fragments which are inserted in the document, and EPS files which contain graphics.

With option noexec, sysquakelatextool is not executed. The typesetting result will still be correct if all its output files (.Imeout and .eps files) already exist and code fragments, and the size of all text strings displayed in graphics if option latexingraphics is used, are not modified.

The main purpose of this option is to let an author make minor modifications on a platform where sysquakelatextool is not supported.

noseparatelogfile By default, standard error channel used when evaluating Sysquake code fragments is redirected to a file whose base name is the same as the main .tex file and the suffix is .lmelog. This file can be opened after typesetting to check that all code fragments were executed correctly. This is especially important when an error occurs, when a code fragment is replaced with ??sysquake?? in the typeset result.

The noseparatelogfile option merges the Sysquake standard error channel with other LaTeX messages.

includelog Instead of storing error and warning messages in a separate file (default) or in the main log file (option noseparatelogfile), with option includelog, this information is appended directly to the typeset output (PDF or DVI file). A reference is added to the code fragment result, which becomes ??sysquake?? (see LME log). If the package hyperref is used, the words LME log are a hypertext link to the error and warning messages.

Additional options with arguments can be set with special LaTeX commands:

\sysquakesize{n} Specifies the amount of memory allocated to Sysquake for LaTeX in megabytes. For special purposes such as large images, a large amount, such as 128 or 256, can be specified.

\sysquakelinewidth{ $\mathbf{w}$ } Specifies the normal line width in EPS files, in 1/72 inch. The default value is 1.

\sysquakepath{**path**} Specifies the list of paths where libraries are searched; each path is separated with a semicolon. By default,

libraries are searched where the installer has placed the standard libraries distributed with Sysquake for LaTeX, and in the same directory as the .tex file being processed.

# 5.2 Other packages

Package sysquake.sty depends on other packages for some tasks. You should also import them with \usepackage if necessary.

**graphicx** Used to include EPS graphics generated by Sysquake for LaTeX.

**epstopdf** Used by pdflatex to convert EPS graphics to PDF. pdflatex cannot include directly EPS files.

**hyperref** Used for hypertext links created by writing text markup to file descriptor 4.

**listings** Used for listings of LME functions inserted with sqlxsource.

The preamble of an article which needs all these packages could be as follows. There is no harm in importing superfluous packages.

```
\documentclass{article}
\usepackage[latexingraphics,noseparatelogfile]{sysquake}
\usepackage{graphicx}
\usepackage{epstopdf}
\usepackage{hyperref}
\usepackage{listings}
\begin{document}
...
\end{document}
```

### 5.3 Commands

Any text output produced by code fragments is inserted in the LaTeX source code. It is subject to further evaluation by LaTeX.

### \sqeval

\sqeval{code} evaluates code fragment code, which is processed as LaTeX code before being evaluated by Sysquake for LaTeX; hence characters aimed at Sysquake which have a meaning in LaTeX, such as \ and \$, must be escaped.

#### **Examples**

```
Simple value:
```

```
The result is \sqeval{disp(1+2);}.
```

Passing a LaTeX counter:

```
\newcounter{n}
\setcounter{n}{10}
The square root of \arabic{n} is \sqeval{disp(sqrt(\arabic{n}));}.
```

### \sqexpr

\sqexpr{expr} evaluates expression expr, formats it for LaTeX and inserts it in the result. The expression should not be terminated by a semicolon or comma.

Currently, the following types are supported: real or complex double, scalar or 2-dim arrays, and strings. Options set by format are used; scientific notation is displayed with the mathematical notation (a multiplication by 10 to some integer power) instead of the classic computer notation.

\sqexpr can be used in text as well as in math mode. It is a fragile command; to use it in a moving argument, such as the argument of \caption, put \protect right before it.

#### **Examples**

Scalar value in an equation:

```
Some small number:
$\frac{\pi}{e^{10}} = \sqexpr{pi/exp(10)}$.

Magic square of size 5:

\[
M_5 = \sqexpr{magic(5)}
\]
```

#### Fragile command:

```
\caption{Simulation with $\alpha=\protect\sqexpr{2*3.9}$}
```

### 5.4 Environment

```
\begin{sysquake} ... \end{sysquake}
```

Text in the sysquake environment is interpreted verbatim as LME code. All LaTeX constructs are ignored, except for the terminating string \end{sysquake}. Standard output produced by functions like disp and fprintf is inserted in the output, where it is processed by LaTeX.

**Remark:** the sysquake environment uses the verbatim package. In the *beamer* class, the [fragile] option should be added to frames where it is used. See below for an example.

#### **Example**

In the fragment below, the eigenvalues of a magic square are computed and displayed in a loop. Function fprintf is used to display symbols with and sqlxvalue to format the numbers.

```
$R$'s eigenvalues $\lambda_i$ are
\begin{sysquake}
R = magic(3);
lambda = eig(R);
for i = 1:length(lambda)
   fprintf('\\[ \\lambda_%d = %s \\]\n', i, sqlxvalue(lambda(i)));
end
\end{sysquake}
```

```
\begin{sysquake}(w,h) ... \end{sysquake}
```

\begin{sysquake}(w,h) starts an environment where sysquake code is evaluated, like \begin{sysquake}. In addition, an EPS file with width w and height h is created with all the output produced by graphical commands the code fragment contains. This EPS is inserted in the LaTeX document in a picture environment with the \includegraphics command of package graphicx.

### **Example**

The fragment below defines a floating figure with the step response of the transfer function  $1/(s^3+2s^2+3s+4)$ . We assume the latexingraphics option of the sysquake package, so that text is processed by LaTeX (see above).

```
\begin{figure}
\begin{center}
\begin{sysquake}(300,200)
step(1, [1,2,3,4]);
```

```
label('$t$','$\\psi$');
legend('$\\psi(t)=\\int_0^\\infty g(\\tau)s(t-\\tau){\\rm d}\\tau$');
\end{sysquake}
\caption{Step response}
\end{center}
\end{figure}
```

Here is an example of a plot in a document with class *beamer*. Note the use of the option [fragile] required by the frame environment.

```
\documentclass{beamer}
\usepackage[latexingraphics]{sysquake}
\usepackage{graphicx}
\usepackage{epstopdf}
\begin{document}
\begin{frame}[fragile] \frametitle{Graphics}
\begin{sysquake}(280,200)
fplot(@(x) (x+0.3)^2+7.2*exp(-3*x^2), [-2,3], 'r');
\end{sysquake}
\end{frame}
\end{document}
```

## **Chapter 6**

# **LME Reference**

This chapter describes LME (Lightweight Math Engine), the interpreter for numeric computing used by Sysquake.

## 6.1 Program format

#### **Statements**

An LME program, or a code fragment typed at a command line, is composed of statements. A statement can be either a simple expression, a variable assignment, or a programming construct. Statements are separated by commas, semicolons, or end of lines. The end of line has the same meaning as a comma, unless the line ends with a semicolon. When simple expressions and assignments are followed by a comma (or an end of line), the result is displayed to the standard output; when they are followed by a semicolon, no output is produced. What follows programming constructs does not matter.

When typed at the command line, the result of simple expressions is assigned to the variable ans; this makes easy reusing intermediate results in successive expressions.

### **Continuation characters**

A statement can span over several lines, provided all the lines but the last one end with three dots. For example,

```
1 + ...
2
```

is equivalent to 1 + 2. After the three dots, the remaining of the line, as well as empty lines and lines which contain only spaces, are ignored.

Inside parenthesis or braces, line breaks are permitted even if they are not escaped by three dots. Inside brackets, line breaks are matrix row separators, like semicolons, unless they follow a comma or a semicolon where they are ignored.

#### **Comments**

Unless when it is part of a string enclosed between single ticks, a single percent character or two slash characters mark the beginning of a comment, which continues until the end of the line and is ignored by LME. Comments must follow continuation characters, if any.

```
a = 2; % comment at the end of a line
x = 5; // another comment
% comment spanning the whole line
b = ... % comment after the continuation characters
    a;
a = 3% no need to put spaces before the percent sign
s = '%'; % percent characters in a string
```

Comments may also be enclosed between /\* and \*/; in that case, they can span several lines.

### **Pragmas**

Pragmas are directives for the LME compiler. They can be placed at the same location as LME statements, i.e. in separate lines or between semicolons or commas. They have the following syntax:

```
_pragma name arguments
```

where name is the pragma name and arguments are additional data whose meaning depends on the pragma.

Currently, only one pragma is defined. Pragmas with unknown names are ignored.

```
Name Arguments Effect
```

```
line n Set the current line number to n
```

\_pragma line 120 sets the current line number as reported by error messages or used by the debugger or profiler to 120. This can be useful when the LME source code has been generated by processing another file, and line numbers displayed in error messages should refer to the original file.

### 6.2 Function Call

Functions are fragments of code which can use *input arguments* as parameters and produce *output arguments* as results. They can be built in LME (*built-in functions*), loaded from optional extensions, or defined with LME statements (*user functions*).

A *function call* is the action of executing a function, maybe with input and/or output arguments. LME supports different syntaxes.

```
fun
fun()
fun(in1)
fun(in1, in2,...)
out1 = fun...
(out1, out2, ...) = fun...
[out1 out2 ...] = fun...
```

Input arguments are enclosed between parenthesis. They are passed to the called function by value, which means that they cannot be modified by the called function. When a function is called without any input argument, parenthesis may be omitted.

Output arguments are assigned to variables or part of variables (structure field, list element, or array element). A single output argument is specified on the left on an equal character. Several output arguments must be enclosed between parenthesis or square brackets (arguments can simply be separated by spaces when they are enclosed in brackets). Parenthesis and square brackets are equivalent as far as LME is concerned; parenthesis are preferred in LME code, but square brackets are available for compatibility with third-party applications.

Output arguments can be discarded without assigning them to variables either by providing a shorter list of variables if the arguments to be discarded are at the end, or by replacing their name with a tilde character. For example to get the index of the maximum value in a vector and to discard the value itself:

```
(\tilde{\ }, index) = max([2, 1, 5, 3]);
```

# 6.3 Named input arguments

Input arguments are usually recognized by their position. Some functions also differentiate them by their data type. This can lead to code which is difficult to write and to maintain. A third method to distinguish the input arguments of a function is to tag them with a name, with a syntax similar to an assignment. Named arguments must follow unnamed arguments.

```
fun(1, [2,3], dim=2, order=1);
```

For some functions, named arguments are an alternative to a sequence of unnamed arguments.

## 6.4 Command syntax

When a function has only literal character strings as input arguments, a simpler syntax can be used. The following conditions must be satisfied:

- No output argument.
- Each input argument must be a literal string
  - without any space, tabulator, comma or semicolon,
  - beginning with a letter, a digit or one of '-/.:\*' (minus, slash, dot, colon, or star),
  - containing at least one letter or digit.

In that case, the following syntax is accepted; left and right columns are equivalent.

```
fun str1 fun('str1')
fun str1 str2 fun('str1','str2')
fun abc,def fun('abc'),def
```

Arguments can also be quoted strings; in that case, they may contain spaces, tabulators, commas, semicolons, and escape sequences beginning with a backslash (see below for a description of the string data type). Quoted and unquoted arguments can be mixed:

```
fun 'a bc\n' fun('a bc\n')
fun strl 'str 2' fun('strl','str 2')
```

The command syntax is especially useful for functions which accept well-known options represented as strings, such as format loose.

### 6.5 Libraries

Libraries are collections of user functions, identified in LME by a name. Typically, they are stored in a file whose name is the library name with a ".lml" suffix (for instance, library stdlib is stored in file "stdlib.lml"). Before a user function can be called, its library must be loaded with the use statement. use statements have an effect only in the context where they are placed, i.e. in a library, or the command-line interface,

or a Sysquake SQ file; this way, different libraries may define functions with the same name provided they are not used in the same context.

In a library, functions can be public or private. Public functions may be called from any context which use the library, while private functions are visible only from the library they are defined in.

# 6.6 Types

### Numerical, logical, and character arrays

The basic type of LME is the two-dimensional array, or matrix. Scalar numbers and row or column vectors are special kinds of matrices. Arrays with more than two dimensions are also supported. All elements have the same type, which are described in the table below. Two non-numeric types exist for character arrays and logical (boolean) arrays. Cell arrays, which contain composite types, are described in a section below.

Туре	Description
double	64-bit IEEE number
complex double	Two 64-bit IEEE numbers
single	32-bit IEEE number
complex single	Two 32-bit IEEE numbers
uint32	32-bit unsigned integer
int32	32-bit signed integer
uint16	16-bit unsigned integer
int16	16-bit signed integer
uint8	8-bit unsigned integer
int8	8-bit signed integer
uint64	64-bit unsigned integer
int64	64-bit signed integer

64-bit integer numbers are not supported by all applications on all platforms.

These basic types can be used to represent many mathematic objects:

**Scalar** One-by-one matrix.

**Vector** n-by-one or one-by-n matrix. Functions which return vectors usually give a column vector, i.e. n-by-one.

**Empty object** 0-by-0 matrix (0-by-n or n-by-0 matrices are always converted to 0-by-0 matrices).

**Polynomial of degree d** 1-by-(d+1) vector containing the coefficients of the polynomial of degree d, highest power first.

**List of n polynomials of same degree d** n-by-(d+1) matrix containing the coefficients of the polynomials, highest power at left.

**List of n roots** n-by-1 matrix.

**List of n roots for m polynomials of same degree n** n-by-m matrix.

**Single index** One-by-one matrix.

**List of indices** Any kind of matrix; the real part of each element taken row by row is used.

**Sets** Numerical array, or list or cell array of strings (see below).

**Boolean value** One-by-one logical array; 0 means false, and any other value (including nan) means true (comparison and logical operators and functions return logical values). In programs and expressions, constant boolean values are entered as false and true. Scalar boolean values are displayed as false or true; in arrays, respectively as F or T.

**String** Usually 1-by-n char array, but any shape of char arrays are also accepted by most functions.

Unless a conversion function is used explicitly, numbers are represented by double or complex values. Most mathematical functions accept as input any type of numeric value and convert them to double; they return a real or complex value according to their mathematical definition.

Basic element-wise arithmetic and comparison operators accept directly integer types ("element-wise" means the operators + - .\* ./ and the functions mod and rem, as well as operators \*/ with a scalar multiplicand or divisor). If their arguments do not have the same type, they are converted to the size of the largest argument size, in the following order:

double > single > uint64 > int64 > uint32 > int32 > uint16
> int16 > uint8 > int8

Literal two-dimensional arrays are enclosed in brackets. Rows are separated with semicolons or line breaks, and row elements with commas or spaces. Here are three different ways to write the same 2-by-3 double array.

```
A = [1, 2, 3; 4, 5, 6];

A = [1 2 3

4 5 6];

A = [1, 2,

3;

4, 5 6];
```

Functions which manipulate arrays (such as reshape which changes their size or repmat which replicates them) preserve their type.

To convert arrays to numeric, char, or logical arrays, use functions + (unary operator), char, or logical respectively. To convert the numeric types, use functions double, single, or uint8 and similar functions.

#### **Numbers**

Double and complex numbers are stored as floating-point numbers, whose finite accuracy depends on the number magnitude. During computations, round-off errors can accumulate and lead to visible artifacts; for example, 2-sqrt(2)\*sqrt(2), which is mathematically 0, yields -4.4409e-16. Integers whose absolute value is smaller than 2°52 (about 4.5e15) have an exact representation, though.

Literal double numbers (constant numbers given by their numeric value) have an optional sign, an integer part, an optional fractional part following a dot, and an optional exponent. The exponent is the power of ten which multiplies the number; it is made of the letter 'e' or 'E' followed by an optional sign and an integer number. Numbers too large to be represented by the floating-point format are changed to plus or minus infinity; too small numbers are changed to 0. Here are some examples (numbers on the same line are equivalent):

```
123 +123 123. 123.00 12300e-2

-2.5 -25e-1 -0.25e1 -0.25e+1

0 0.0 -0 1e-99999

inf 1e999999

-inf -1e999999
```

Literal integer numbers may also be expressed in hexadecimal with prefix 0x, in octal with prefix 0, or in binary with prefix 0b. The four literals below all represent 11, stored as double:

```
0xb
013
0b1011
11
```

Literal integer numbers stored as integers and literal single numbers are followed by a suffix to specify their type, such as 2int16 for the number 2 stored as a two-byte signed number or 0x300uint32 for the number whose decimal representation is 768 stored as a four-byte unsigned number. All the integer types are valid, as well as single. This syntax gives the same result as the call to the corresponding function (e.g. 2int16 is the same as int16(2)), except when the integer number cannot be represented with a double; then the number is rounded

to the nearest value which can be represented with a double. Compare the expressions below:

Expression	Value
uint64(123456789012345678)	123456789012345696
123456789012345678uint64	123456789012345678

Literal complex numbers are written as the sum or difference of a real number and an imaginary number. Literal imaginary numbers are written as double numbers with an i or j suffix, like 2i, 3.7e5j, or 0xffj. Functions i and j can also be used when there are no variables of the same name, but should be avoided for safety reasons.

The suffices for single and imaginary can be combined as isingle or jsingle, in this order only:

```
2jsingle
3single + 4isingle
```

Command format is used to specify how numbers are displayed.

### **Strings**

Strings are stored as arrays (usually row vectors) of 16-bit unsigned numbers. Literal strings are enclosed in single quotes:

```
'Example of string'
```

The second string is empty. For special characters, the following escape sequences are recognized:

Character	Escape seq.	<b>Character code</b>
Null	\0	0
Bell	\a	7
Backspace	\b	8
Horizontal tab	\t	9
Line feed	\n	10
Vertical tab	\v	11
Form feed	\f	12
Carriage return	\r	13
Single tick	\'	39
Single tick	'' (two ')	39
Backslash	\\	92
Hexadecimal number	\xhh	hh
Octal number	\000	000
16-bit UTF-16	\uhhhh	1 UTF-16 code
21-bit UTF-32	\Uhhhhhhhhh	1 or 2 UTF-16 codes

For octal and hexadecimal representations, up to 3 (octal) or 2 (hexadecimal) digits are decoded; the first non-octal or non-hexadecimal

digit marks the end of the sequence. The null character can conveniently be encoded with its octal representation, \0, provided it is not followed by octal digits (it should be written \000 in that case). It is an error when another character is found after the backslash. Single ticks can be represented either by a backslash followed by a single tick, or by two single ticks.

Depending on the application and the operating system, strings can contain directly Unicode characters encoded as UTF-8, or MBCS (multibyte character sequences). 16-bit characters encoded with \uhhhh escape sequences are always accepted and handled correctly by all built-in LME functions (low-level input/output to files and devices which are byte-oriented is an exception; explicit UTF-8 conversion should be performed if necessary).

UTF-32 sequences \Uhhhhhhhh assume UTF-16 encoding. In sequences \uhhhh and \Uhhhhhhhh, up to 4 or 8 hexadecimal digits can be provided, respectively, but the first non-hexadecimal character marks the end of the sequence.

#### Inline data

For large amounts of text or binary data, the syntax described above is impractical. Inline data is a special syntax for storing strings as raw text or uint8 arrays as base64.

Strings (char arrays of dimension 1-by-n) can be defined in the source code as raw text without any escape sequence with the following syntax:

```
@/text marker
text
marker
```

where @/text is that literal sequence of six characters followed or not by spaces and tabs, marker is an arbitrary sequence of characters without spaces, tabs or end-of-lines which does not occur in the text, and text is the text itself. The spaces, tabs and first end-of-line which follow the first marker are ignored. The last marker must be at the beginning of a line; this means that the string always ends with an end-of-line. The whole text inline data is equivalent to a string with the corresponding characters and can be located in an assignment or any expression. End-of-line sequences (\n, \r or \r\n) are replaced by a single linefeed character.

Here is an example of a short fragment of C code, assigned to variable src. The sequence  $\n$  is not interpreted as an escape sequence by LME; it results in the two characters  $\n$  and  $\n$  in src. The trailing semicolon suppresses the display of the assignment, like in any LME expression.

```
src = @/text""
int main() {
  printf("Hello, data!\n");
}
""":
```

Arrays of uint8, of dimension n-by-1 (column vectors), can be defined in the source code in a compact way using the base64 encoding in inline data:

```
@/base64 data
```

where @/base64 is that literal sequence of eight characters, followed by spaces and/or line breaks, and the data encoded with base64 (see RFC 2045). The base64-encoded data can contain lowercase and uppercase letters a-z and A-Z, digits 0-9, and characters / (slash) and + (plus), and is followed by 0, 1 or 2 characters = (equal) for padding. Spaces, tabs and line breaks are ignored. Comments are not allowed.

The first character which is not a valid base64 character signals the end of the inline data and the beginning of the next token of source code. Inline data can be a part of any expression, assignment or function call, like any other literal value. In the case where the inline data is followed by a character which would erroneously be interpreted as more base64 codes (e.g. neither padding with = nor statement terminator and a keyword at the beginning of the following line), it should be enclosed in parenthesis.

Inline data can be generated with the base64encode function. For example, to encode uint8(0:255).' as inline data, you can evaluate

```
base64encode(uint8(0:255))
```

Then copy and paste the result to the source code, for instance as follows to set a variable d (note how the semicolon will be interpreted as the delimiter following the inline data, not the data iteself):

```
d = @/base64
```

AAECAwQFBgcICQoLDA00DxAREhMUFRYXGBkaGxwdHh8gISIjJCUmJygpKiss LS4vMDEyMzQ1Njc4OTo7PD0+P0BBQkNERUZHSElKS0xNTk9QUVJTVFVWV1hZ WltcXV5fYGFiY2RlZmdoaWprbG1ub3BxcnN0dXZ3eHl6e3x9fn+AgYKDhIWG h4iJiouMjY6PkJGSk5SVlpeYmZqbnJ2en6Choq0kpaanqKmqq6ytrq+wsbKz tLW2t7i5uru8vb6/wMHCw8TFxsfIycrLzM30z9DR0tPU1dbX2Nna29zd3t/g 4eLj50Xm5+jp6uvs7e7v8PHy8/T19vf4+fr7/P3+/w== ;

### Lists and cell arrays

Lists are ordered sets of other elements. They may be made of any type, including lists. Literal lists are enclosed in braces; elements are separated with commas.

```
{1,[3,6;2,9],'abc',{1,'xx'}}
Lists can be empty:
{}
```

List's purpose is to collect any kind of data which can be assigned to variables or passed as arguments to functions.

Cell arrays are arrays whose elements (or cells) contain data of any type. They differ from lists only by having more than one dimension. Most functions which expect lists also accept cell arrays; functions which expect cell arrays treat lists of n elements as 1-by-n cell arrays.

To create a cell array with 2 dimensions, cells are written between braces, where rows are separated with semicolons and row elements with commas:

```
{1, 'abc'; 27, true}
```

Since the use of braces without semicolon produces a list, there is no direct way to create a cell array with a single row, or an empty cell array. Most of the time, this is not a problem since lists are accepted where cell arrays are expected. To force the creation of a cell array, the reshape function can be used:

```
reshape({'ab', 'cde'}, 1, 2)
```

#### **Structures**

Like lists and cell arrays, structures are sets of data of any type. While list elements are ordered but unnamed, structure elements, called *fields*, have a name which is used to access them.

There are three ways to make structures: with field assignment syntax inside braces, with the struct function, or by setting each field in an assignment. s.f refers to the value of the field named f in the structure s. Usually, s is the name of a variable; but unless it is in the left part of an assignment, it can be any expression which evaluates to a structure.

With the assignments above, a.os{3} is 'Linux' and c.s.radius is 90.

While the primary way to access structure fields is by name, field order is still preserved, as can be seen by displaying the structure, getting the field names with fieldnames, or converting the structure to a cell array with struct2cell. The fields can be reordered with orderfields.

### Structure arrays

While structure fields can contain any type of array and cell arrays can have structures stored in their cells, structure arrays are arrays where each element has the same named fields. Plain structures are structure arrays of size [1,1], like scalar numbers are arrays of size [1.1].

Values are specified first by indices between parenthesis, then by field name. Braces are invalid to access elements of structure arrays (they can be used to access elements of cell arrays stored in structure array fields).

Structure arrays are created from cell arrays with functions structarray or cell2struct, or by assigning values to fields.

Column struct arrays (1-dimension) can be defined with field assignments inside braces by separating array elements with semicolons. Missing fields are set to the empty array [].

```
D = \{a = 1, b = 2; a = 5, b = 3; b = 8\};
```

### Value sequences

Value sequences are usually written as values separated with commas. They are used as function input arguments or row elements in arrays or lists.

When expressions involving lists, cell arrays or structure arrays evaluate to multiple values, these values are considered as a value sequence, or part of a value sequence, and used as such in context

where value sequences are expected. The number of values can be known only at execution time, and may be zero.

```
L = {1, 2};
v = [L{:}]; // convert L to a row vector
c = complex(L{:}); // convert L to a complex number
```

Value sequences can arise from element access of list or cell arrays with brace indexing, or from structure arrays with field access with or without parenthesis indexing.

#### **Function references**

Function references are equivalent to the name of a function together with the context in which they are created. Their main use is as argument to other functions. They are obtained with operator @.

### Inline and anonymous functions

Inline and anonymous functions encapsulate executable code. They differ only in the way they are created: inline functions are made with function inline, while anonymous functions have special syntax and semantics where the values of variables in the current context can be captured implicitly without being listed as argument. Their main use is as argument to other functions.

#### Sets

Sets are represented with numeric arrays of any type (integer, real or complex double or single, character, or logical), or lists or cell arrays of strings. Members correspond to an element of the array or list. All set-related functions accept sets with multiple values, which are always reduced to unique values with function unique. They implement membership test, union, intersection, difference, and exclusive or. Numerical sets can be mixed; the result has the same type as when mixing numeric types in array concatenation. Numerical sets and list or cell arrays os strings cannot be mixed.

#### Null

Null stands for the lack of data. It is both a data type and the only value it can represent. It can be assigned to a variable, be contained in a list or cell array element or a structure field, or passed as an input or output argument to/from a function.

Null is a recent addition to LME, where the lack of data is usually represented by the empty matrix []. It is especially useful when LME

is interfaced with languages or libraries where the null value has a special meaning, such as SQL (Structured Query Language, used with relational databases) or the DOM (Document Object Model, used with XML).

### **Objects**

Objects are the basis of *Object-Oriented Programming* (OOP), an approach of programming which puts the emphasis on encapsulated data with a known programmatic interface (the objects). Two OOP languages in common use today are C++ and Java.

The exact definition of OOP varies from person to person. Here is what it means when it relates to LME:

**Data encapsulation** Objects contain data, but the data cannot be accessed directly from the outside. All accesses are performed via special functions, called *methods*. What links a particular method to a particular object is a class. Class are identified with a name. When an object is created, its class name is specified. The names of methods able to act on objects of a particular class are prefixed with the class name followed with two colons. Objects are special structures whose contents are accessible only to its methods.

**Function and operator overloading** Methods may have the same name as regular functions. When LME has to call a function, it first checks the type of the input arguments. If one of them is an object, the corresponding method is called, rather than the function defined for non-object arguments. A method which has the same name as a function or another method is said to *overload* it. User functions as well as built-in ones can be overloaded. Operators which have a function name (for instance x+y can also be written plus(x,y)) can also be overloaded. Special functions, called object *constructors*, have the same name as the class and create new objects. They are also methods of the class, even if their input arguments are not necessarily objects.

**Inheritance** A class (*subclass*) may extend the data and methods of another class (*base class* or *parent*). It is said to *inherit* from the parent. In LME, objects from a subclass contain in a special field an object of the parent class; the field name has the same name as the parent class. If LME does not find a method for an object, it tries to find one for its parent, great-parent, etc. if any. An object can also inherit from several parents.

Here is an example of the use of polynom objects, which (as can be guessed from their name) contain polynomials. Statement use polynom imports the definitions of methods for class polynom and others.

```
use polynom;
p = polynom([1,5,0,1])
p =
    x^3+5x^2+1
q = p^2 + 3 * p / polynom([1,0])
q =
    x^6+10x^5+25x^4+2x^3+13x^2+15x+1
```

## 6.7 Input and Output

LME identifies channels for input and output with non-negative integer numbers called *file descriptors*. File descriptors correspond to files, devices such as serial port, network connections, etc. They are used as input argument by most functions related to input and output, such as fprintf for formatted data output or fgets for reading a line of text.

Note that the description below applies to most LME applications. For some of them, files, command prompts, or standard input are irrelevant or disabled; and standard output does not always correspond to the screen.

At least four file descriptors are predefined:

Value	Input/Output	Purpose
0	Input	Standard input from keyboard
1	Output	Standard output to screen
2	Output	Standard error to screen
3	Output	Prompt for commands

You can use these file descriptors without calling any opening function first, and you cannot close them. For instance, to display the value of  $\pi$ , you can use fprintf:

```
fprintf(1, 'pi = %.6f\n', pi);
  pi = 3.141593
```

Some functions use implicitly one of these file descriptors. For instance disp displays a value to file descriptor 1, and warning displays a warning message to file descriptor 2.

File descriptors for files and devices are obtained with specific functions. For instance fopen is used for reading from or writing to a file. These functions have as input arguments values which specify what to open and how (file name, host name on a network, input or output mode, etc.), and as output argument a file descriptor. Such file descriptors are valid until a call to fclose, which closes the file or the connection.

### 6.8 Error Messages

When an error occurs, the execution is interrupted and an error message explaining what happened is displayed, unless the code is enclosed in a try/catch block. The whole error message can look like

```
> use stat
> iqr(123)

Index out of range for variable 'M' (stat/prctile;61)
-> stat/iqr;69
```

The first line contains an error message, the location in the source code where the error occurred, and the name of the function or operator involved. Here stat is the library name, prctile is the function name, and 61 is the line number in the file which contains the library. If the function where the error occurs is called itself by another function, the whole chain of calls is displayed; here, prctile was called by ign at line 69 in library stat.

Here is the list of errors which can occur. For some of them, LME attempts to solve the problem itself, e.g. by allocating more memory for the task.

**Stack overflow** Too complex expression, or too many nested function calls.

**Data stack overflow** Too large objects on the stack (in expressions or in nested function calls).

**Variable overflow** Not enough space to store the contents of a variable.

**Code overflow** Not enough memory for compiling the program.

**Not enough memory** Not enough memory for an operation outside the LME core.

**Algorithm does not converge** A numeric algorithm does not converge to a solution, or does not converge quickly enough. This usually means that the input arguments have invalid values or are ill-conditioned.

**Incompatible size** Size of the arguments of an operator or a function do not agree together.

**Bad size** Size of the arguments of a function are invalid.

**Non-vector array** A row or column vector was expected, but a more general array was found.

**Not a column vector** A column vector was expected, but a more general array was found.

- **Not a row vector** A row vector was expected, but a more general array was found.
- **Non-matrix array** A matrix was expected, but an array with more than 2 dimensions was found.
- **Non-square matrix** A square matrix was expected, but a rectangular matrix was found.
- **Index out of range** Index negative or larger than the size of the array.
- **Wrong type** String or complex array instead of real, etc.
- **Non-integer argument** An argument has a fractional part while an integer is required.
- **Non positive integer argument** An argument is not a positive integer as expected.
- **Argument out of range** An argument is outside the permitted range of values.
- **Non-scalar argument** An argument is an array while a scalar number is required.
- **Non-object argument** An object is required as argument.
- **Not a permutation** The argument is not a permutation of the integers from 1 to n.
- **Bad argument** A numeric argument has the wrong site or the wrong value.
- **Unknown option** A string option has an invalid value.
- **Object too large** An object has a size larger than some fixed limit.
- **Undefined variable** Attempt to retrieve the contents of a variable which has not been defined.
- **Undefined input argument** Attempt to retrieve the contents of an input argument which was neither provided by the caller nor defined in the function.
- **Undefined function** Attempt to call a function not defined.
- **Too few or too many input arguments** Less or more arguments in the call than what the function accepts.

- **Too few or too many output arguments** Less or more left-side variables in an assignment than the function can return.
- **Syntax error** Unspecified compile-time error.
- "function" keyword without function name Incomplete function header.
- **Bad function header** Syntax error in a function header
- **Missing expression** Statement such as if or while without expression.
- **Unexpected expression** Statement such as end or else followed by an expression.
- **Incomplete expression** Additional elements were expected during the compilation of an expression, such as right parenthesis or a sub-expression at the right of an operator.
- "for" not followed by a single assignment for is followed by an expression or an assignment with multiple variables.
- **Bad variable name** The left-hand part of an assignment is not a valid variable name (e.g. 2=3)
- **String without right quote** The left quote of a string was found, but the right quote is missing.
- **Unknown escape character sequence** In a string, the backslash character is not followed by a valid escape sequence.
- **Unexpected right parenthesis** Right parenthesis which does not match a left parenthesis.
- **Unexpected right bracket** Right bracket which does not match a left bracket.
- **Unrecognized or unexpected token** An unexpected character was found during compilation (such as (1+))
- "end" not in an index expression end was used outside of any index sub-expression in an expression.
- "beginning" not in an index expression beginning was used outside of any index sub-expression in an expression.
- "matrixcol" not in an index expression matrixcol was used outside of any index sub-expression in an expression.
- "matrixrow" not in an index expression matrixrow was used outside of any index sub-expression in an expression.

"matrixrow" or "matrixcol" used in the wrong index
matrixrow was used in an index which was not the first one, or
matrixcol was used in an index which was not the only one or
the second one.

- **Compilation overflow** Not enough memory during compilation.
- **Too many nested subexpressions** The number of nested of subexpressions is too high.
- **Variable table overflow** A single statement attempts to define too many new variables at once.
- **Expression too large** Not enough memory to compile a large expression.
- **Too many nested (), [] and {}** The maximum depth of nested subexpressions, function argument lists, arrays and lists is reached.
- **Too many nested programming constructs** Not enough memory to compile that many nested programming constructs such as if, while, switch, etc.
- **Wrong number of input arguments** Too few or too many input arguments for a built-in function during compilation.
- **Wrong number of output arguments** Too few or too many output arguments for a built-in function during compilation.
- **Too many indices** More than two indices for a variable.
- **Variable not found** A variable is referenced, but appears neither in the arguments of the function nor in the left part of an assignment.
- Unbounded language construct if, while, for, switch, or try
   without end.
- **Unexpected "end"** The end statement does not match an if, switch, while, for, or catch block.
- "case" or "otherwise" without "switch" The case or otherwise statement is not inside a switch block.
- "break" or "continue" not in a loop The break or continue statement is not inside a while or for block.

- **Variable name reused** Same variable used twice as input or as output argument.
- **Too many user functions** Not enough memory for that many user functions.
- **Attempt to redefine a function** A function with the same name already exists.
- **Can't find function definition** Cannot find a function definition during compilation.
- **Unexpected end of expression** Missing right parenthesis or square bracket.
- **Unexpected statement** Expression expected, but a statement is found (e.g. if).
- **Null name** Name without any character (when given as a string in functions like feval and struct).
- **Name too long** More than 32 characters in a variable or function name.
- **Unexpected function header** A function header (keyword "function") has been found in an invalid place, for example in the argument of eval.
- **Function header expected** A function header was expected but not found.
- **Bad variable in the left part of an assignment** The left part of an assignment does not contain a variable, a structure field, a list element, or the part of an array which can be assigned to.
- **Bad variable in a for loop** The left part of the assignment of a for loop is not a variable.
- **Source code not found** The source code of a function is not available.
- **File not found** fopen does not find the file specified.
- **Bad file ID** I/O function with a file descriptor which neither is standard nor corresponds to an open file or device.
- **Cannot write to file** Attempt to write to a read-only file.
- **Bad seek** Seek out of range or attempted on a stream file.
- **Too many open files** Attempt to open too many files.
- **End of file** Attempt to read data past the end of a file.

**Timeout** Input or output did not succeed before a too large amount of time elapsed.

**No more OS memory** The operating system cannot allocate more memory.

**Bad context** Call of a function when it should not (application-dependent).

**Not supported** The feature is not supported, at least in the current version.

### 6.9 Character Set

There exist different standards to represent characters. In LME, characters are stored as 16-bit unsigned integer numbers. The mapping between these codes and the actual characters they represent depends on the application and the operating system. Currently, on macOS, Windows and Linux, Sysquake uses the UTF-16 character encoding (i.e. Unicode characters encoded in one or two 16-bit words).

To make the exchange of files possible without manual conversion, all text files used by LME applications can have their character set specified explicitly. In Sysquake, this includes library files (.lml), SQ files (.sq), and SQ data files (.sqd). Versions of Sysquake using Unicode (currently macOS and Linux) convert automatically files with a charset specification.

The character set specification is a comment line with the following format:

```
// charset=charsetname
or
% charset=charsetname
```

Spaces between the comment mark and the keyword charset are ignored. The comment line must be the first or the second line of the text file. The character set *charsetname* must be one of the following:

```
ascii or usascii ASCII
utf-8 or utf8 UTF-8 (unicode)
iso-8859-1 or iso-latin-1 ISO-Latin-1 (Windows 1252)
macintosh or macosroman Mac OS Classic
```

Here are advices about the use of character set specifications, both for the current transition phase where Sysquake for Windows does not use Unicode and for the future.

If you need only ASCII (typically because you work in English, or for files without text or where unaccented letters are acceptable), do not add any character set specification (ASCII is a subset of all supported character sets) or add charset=ascii as an indication that the file should contain only 7-bit characters.

If you need accented characters found in western European languages, use ISO-8859-1 with an explicit character set specification on Windows and other platforms if you need cross-platform compatibility, or any character set with a character set specification otherwise.

If you need another native character set on Windows, do not add any character set specification, and switch to UTF-8 as soon as a unicode version of Sysquake becomes available.

#### 6.10 Formatted text

Like most text-based programming languages, LME primary text format is plain text, without any character or paragraph formatting. To improve readability, it also supports formatted text. Formatting information used to change the character font and size and how paragraphs are aligned is embedded in the text itself as markup, i.e. special sequences of characters unlikely to occur in normal text. This is similar to HTML or LaTeX, but with a simpler syntax similar to what is used in wikis and blogs. The markup which has been chosen for LME is Creole, a collaborative effort to create a common markup language to be used across different wikis, and more precisely the open-source NME implementation.

### Formatted text output channel

In addition to the standard output channel (file descriptor 1) and the standard error channel (file descriptor 2), LME has a special channel for formatted output (file descriptor 4). Anything written to that channel is parsed so that markup constructs are interpreted; the result is displayed in the command window. Each write command should contain a whole block of text with markup; there is no buffering across output commands.

Not all versions of LME support formatted output, and how formatted output is displayed depends on the application and the platform. For instance, in shell applications, word-wrap is applied to paragraphs, but bold or italic text is rendered as plain text.

#### Example

```
T = 27.3;
fprintf(4, '=Report=\nTemperature is %.1f\n', T);
```

The same markup is used in LME applications at other places. For example, in Sysquake, the version and help information can contain markup.

### Markup reference

Text with markup is plain text with a few character sequences which are recognized by the markup processor and change the format of the result. The result is styled text with titles, paragraphs with justification, bold and italic faces, etc.

There are two kinds of markup constructs: blocks and inline. Blocks are paragraphs (indented or not), headings, list items, table cells, horizontal rules and block extensions. Inline constructs are character styles, verbatim text and links; they are part of blocks. Except in preformatted blocks, sequences of spaces or tabs are replaced with single spaces.

#### **Paragraphs**

Paragraphs are made of lines whose first character is not one of \*#:;= nor sequence  $\{\{\{, ---, or <<<.\} Lines can begin with sequences <math>**$  and ##.

Paragraphs end with blank lines or next heading, list, table, indented paragraph, preformatted block of text, horizontal rule or block extension. They can contain styled text, links, spans of verbatim text, and inline extensions.

#### Example

This is a paragraph written in two lines.

This is another paragraph.

### Indenting

Indented paragraphs are paragraphs which begin with a colon. Multiple colons define the level of indenting. Indented paragraphs can contain styled text, links, spans of verbatim text, and inline extensions.

### Example

This is a normal paragraph. :This is an indented

```
paragraph in two lines.
::This is more indented.
```

#### **Headings**

Headings are made of a single line beginning with an equal character. The number of equal characters defines the level of title, from main title (=) to sub-sub-subtitle (====). Headings may end with a sequence of equal characters.

#### Examples

```
=Level 1 heading=
Paragraph
=Another level 1 heading, without trailing equal character
==Level 2 heading==
===Level 3 heading===
```

#### Lists

Lists are collections of items. There are three kinds of lists: unnumbered lists, numbered lists, and definition lists. Lists can be nested; they end with the next heading, indented paragraph, table, or blank line.

Unnumbered lists are represented as indented paragraphs with a bullet. Each item begins with a star character (\*); it can span multiple lines.

Numbered lists are represented as indented paragraphs with a number. Items are numbered automatically with consecutive integers starting at 1. Each item begins with a sharp character (#); it can span multiple lines.

Definition lists are made of two kinds of items: title, typically displayed in bold font, and definition, typically displayed indented. Titles begin with a semicolon at the beginning of a line. Definitions either follow the title, separated with a colon; or they begin on a new line beginning with a colon.

List nesting can combine different kinds of lists. Sublist items begin with multiple markers, the first one(s) corresponding to the enclosing list(s). For instance, items of an unnumbered list nested in a numbered list should start with #\* at the beginning of the line, without any preceding space. List markers must be used in a consistent way; for example, ## at the beginning of a line in an unnumbered list is not interpreted as a second-level numbered list item, but as monospace style (see below).

#### **Examples**

```
* First item of unnumbered list.
* Second
item, in two lines.
*Third item without space (spaces are optional).

# First item of numbered list.
# Second item.
#* First item on unnumbered sublist.
#* Second item.
# Thirst item of top-level numbered list.

; First title of definition list
: Definition of first item.
; Second title: Second definition
beginning on the same line.
```

Paragraph separated with a blank line.

#### **Tables**

Tables are rectangular array of cells. They are made of one line per row. Each cell begins with character |. Heading cells (typically displayed in bold face) begin with |=. Rows may end with a trailing |.

### Example

In the table below, the first row and the first column contain headings. The very first cell is empty.

```
||=First column|=Second column
|=First row|Cell 1,1|Cell 1,2
|=Second row|Cell 2,1|Cell 2,2
```

#### **Preformatted**

Preformatted text is a block of text displayed literally, including line feeds. Preformatted text is preceded with a line containing {{{ and is followed by a line containing }}}.

### Example

```
This is some C code:
{{{
  double f(double x)
  {
    return 2 * x; // f(x) = 2x
```

```
}
}}}
```

In preformatted text, lines which begin with }}}, with leading spaces or not, must have an additional space which is discarded in the output.

#### Horizontal rules

Horizontal rules are single lines containing four hyphens.

#### Example

```
Paragraph.
---
Paragraph following a horizontal rule.
```

#### **Extensions**

Sequences << and <<< are reserved for extensions.

#### **Character style**

Inside paragraphs, indented paragraphs, headings, list elements, and table cells, the following two-character sequences toggle on or off the corresponding style. It is not mandatory to nest spans of styled characters between matching style sequences. Style is reset at the end of text block.

Markup	Style
**	Bold
//	Italic
##	Monospace
" (two commas)	Subscript
^	Superscript
(two underscores)	Underlined

Double stars and double sharps are interpreted as the beginning of list items when they are the first characters of a line in the context of lists. To be interpreted as style markup, they can be preceded by a space.

#### Example

```
This is //italic text//, **bold text**, and //**bold italic text**//.
```

#### **Escape character**

The tilde character, when followed by any character except space, tab or line feed, is an escape character; it is not displayed and the next character loses its special meaning, if any.

#### Example

```
Two stars: "*" or "*" or *"; tilde: "".

is rendered as "Two stars: ** or ** or **; tilde: "."
```

#### **Verbatim**

Verbatim text is a sequence of characters enclosed between {{{ and }}}}. After {{{, all characters are output verbatim, without any markup interpreting, until the next }}} or the end of text block. Multiple spaces and tabs and single line feeds are still converted to single spaces, though. Verbatim text is an alternative to the escape character; it is more convenient for sequences of characters.

### Example

```
\{\{\{**\}\}\}\ //\{\{\{\{\{xx\}\}\}\}\}\}// is rendered as "** \{\{\{xx\}\}\}\}".
```

#### Line break

Except in preformatted blocks, line breaks are not preserved. The sequence \\ forces a line break.

### Example

```
The next line of this paragraph begins...\\here!
```

#### Links

Hypertext links (URLs) are enclosed between [[ and ]]. The text displayed as the link is either the same as the URL itself if there is no | character, or it is what follows |. No markup is recognized in the URL part; what follows | can contain styled text and verbatim text. Spaces surrounding | are ignored.

#### Examples

```
* Simple link: [[https://calerga.com]]
```

- \* Link with link text: [[https://calerga.com | Calerga]]
- \* Link with styled link text: [[https://calerga.com | \*\*Calerga\*\*]]

# 6.11 List of Commands, Functions, and **Operators**

### **Programming keywords**

break for function case alobal catch hideimplementation clear continue define otherwise endfunction persistent else private elseif public error repeat

rethrow return switch trv until use

useifexists

while

## **Programming operators and functions**

assert fun2str sandbox Variable assignment inline sandboxtrust Operator () isdefined str2fun Operator @ isfun str2obi builtin isglobal subsasgn deal lasterr subsref dumpvar lasterror varargin eval namedargin varargout feval nargin fevalx nargout

### **Platform**

exist	iskeyword	lookfor
help	ismac	variables
info	ispc	which
inmem	isunix	

# **Arrays**

[]	inthist	permute
,	ipermute	rand
;	isempty	randi
:	length	randn
arrayfun	linspace	repmat
beginning	logspace	reshape
cat	magic	rng
diag	matrixcol	rot90
end	matrixrow	size
eye	meshgrid	sort
find	ndgrid	squeeze
flipdim	ndims	sub2ind
fliplr	nnz	unique
flipud	numel	unwrap
ind2sub	ones	zeros

# **Strings**

base64decode lo base64encode ma char ma deblank ischar isdigit isempty isletter	ength ower othml othmlpoly otstr olit orintf ocanf orcmp	strmatch strrep strtok strtrim unicodeclass upper utf32decode utf32encode utf8decode utf8encode
---	--	--

#### Hash

hmac shal md5 shal

#### Lists

{} islist num2list
apply length replist
join list2num
isempty map

### **Cell arrays**

cell iscell cellfun num2cell

### **Structures and structure arrays**

cell2struct isstruct struct2cell
cellfun orderfields structarray
fieldnames rmfield structmerge
getfield setfield
isfield struct

### **Null value**

isnull null

### **Objects**

class isobject superiorto inferiorto methods

isa superclasses

### **Logical operators**

### **Logical functions**

all isfinite isprime isfloat isrow any false isinf isscalar find isinteger isspace isletter ischar isvector iscolumn islogical logical isdigit ismatrix true isempty isnan xor isequal isnumeric

**Bitwise functions** 

bitall	bitget	bitxor
bitand	bitor	graycode
bitany	bitset	igraycode
bitcmp	bitshift	

# **Integer functions**

int8	int64	uint16
int16	map2int	uint32
int32	uint8	uint64

### **Set functions**

intersect	setdiff	union
ismember	setxor	unique

### **Constants**

eps	inf	рi
false	intmax	realmax
flintmax	intmin	realmin
goldenratio	j	true
i	nan	

### **Arithmetic functions**

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# Trigonometric functions in radians

acos	atan	sec
acot	atan2	sin
acsc	cos	tan
asec	cot	
asin	CSC	

# **Trigonometric functions in degrees**

acosd	atand	secd
acotd	atan2d	sind
acscd	cosd	tand
asecd	cotd	
asind	cscd	

# **Hyperbolic functions**

acosh	asinh	csch
acoth	atanh	sech
acsch	cosh	sinh
asech	coth	tanh

# Other scalar math functions

abs	erfcinv	log
angle	erfcx	log10
beta	erfinv	log1p
betainc	exp	log2
betaln	expm1	nchoosek
conj	factor	nthroot
diln	factorial	rat
ellipam	gamma	real
ellipe	gammainc	reallog
ellipf	gammaln	realpow
ellipj	gcd	realsqrt
ellipke	hypot	sign
erf	imag	sinc
erfc	lcm	sqrt

# **Type conversion functions**

cast	fix	single
ceil	floor	swapbytes
complex	round	typecast
double	roundn	

# **Matrix math functions**

,	fft	null
. '	funm	orth
balance	hess	pinv
care	householder	qr
chol	householderapply	rank
cond	ifft	schur
conv2	inv	sqrtm
dare	linprog	svd
det	logm	trace
dlyap	lu	tril
eig	lyap	triu
expm	norm	

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# **Geometry functions**

cart2pol cross pol2cart cart2sph dot sph2cart

# **Probability distribution functions**

cdf pdf
icdf random

# Statistic functions

cov max moment
cummax mean skewness
cummin median std
kurtosis min var

# **Polynomial math functions**

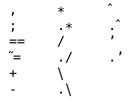
addpol filter polyint conv poly polyval deconv polyder roots

# Interpolation and triangulation functions

delaunay interp1 voronoi
delaunayn interpn voronoin
griddata tsearch

griddata tsearch griddatan tsearchn

# **Quaternion operators**



# **Quaternion math functions**

abs	q2mat	real
conj	q2rpy	rpy2q
cos	q2str	sign
cumsum	qimag	sin
diff	qinv	sqrt
exp	qnorm	sum
log	qslerp	
mean	quaternion	

# Other quaternion functions

beginning	fliplr	permute
cat	flipud	repmat
char	ipermute	reshape
disp	isempty	rot90
dumpvar	isquaternion	size
double	length	squeeze
end	ndims	subsasgn
flipdim	numel	subsref

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# Non-linear numerical functions

fminbnd	integral	ode45
fminsearch	lsqcurvefit	odeset
fsolve	lsqnonlin	optimset
fzero	ode23	-

# **Dynamical systems functions**

c2dm	margin	zp2ss
d2cm	ss2tf	
dmargin	tf2ss	

# Input/output

bwrite	format	redirect
disp	fprintf	sprintf
error	fread	sread
fclose	frewind	sscanf
feof	fscanf	swrite
fgetl	fseek	warning
fgets	ftell	
fionread	fwrite	

# **Files**

fopen

# **Path manipulation**

fileparts filesep fullfile

#### **XML**

getElementById getElementsByTagName saxcurrentline saxcurrentpos

saxnew saxnext saxrelease xmlread xmlreadstring xmlrelease

# **Basic graphics**

activeregion
altscale
area
bar
barh
circle
colormap
contour
figurestyle

image
label
legend
line
math
pcolor
plot
plotoption
plotset

fplot

polar
quiver
scale
scalefactor
subplotstyle
text
tickformat
ticks
otion title

# **Graphics for dynamical systems**

dsigma

bodemag bodephase dbodephase dimpulse dinitial dlsim dnichols dnyquist

fontset

dstep erlocus hgrid hstep impulse initial lsim ngrid nichols nyquist plotroots rlocus sgrid sigma step zgrid

# **Data compression**

deflate inflate

# Image input/output

imageread imageset imagewrite

### **Date and time**

```
cal2julian julian2cal tic clock posixtime toc
```

# Unix

```
cd getenv sleep
cputime pwd unix
```

# **Sysquake Remote**

```
beginfigure htmlspecialchars urldecode endfigure http urlencode figurelist httpheader getclick httpvars
```

# **6.12 Variable Assignment and Subscripting**

# Variable assignment

Assignment to a variable or to some elements of a matrix variable.

# **Syntax**

```
var = expr
(var1, var2, ...) = function(...)
```

# **Description**

var = expr assigns the result of the expression expr to the variable var. When the expression is a naked function call, (var1,var2,...) = function(...) assigns the value of the output arguments of the function to the different variables. Usually, providing less variables than the function can provide just discards the superfluous output arguments; however, the function can also choose to perform in a different way (an example of such a function is size, which returns the number of rows and the number of columns of a matrix either as two numbers if there are two output arguments, or as a 1-by-2 vector if there is a single output argument). Providing more variables than what the function can provide is an error.

Variables can store any kind of contents dynamically: the size and type can change from assignment to assignment.

A subpart of a matrix variable can be replaced with the use of parenthesis. In this case, the size of the variable is expanded when required; padding elements are 0 for numeric arrays and empty arrays [] for cell arrays and lists.

#### See also

Operator (), operator {}, clear, exist, for, subsasgn

# beginning

First index of an array.

# Syntax

```
v(...beginning...)
A(...beginning...)
function e = C::beginning(obj, i, n)
```

# **Description**

In an expression used as an index to access some elements of an array, beginning gives the index of the first element (line or column, depending of the context). It is always 1 for native arrays.

beginning can be overloaded for objects of used-defined classes. Its definition should be have a header equivalent to function e=C::beginning(obj,i,n), where C is the name of the class, obj is the object to be indexed, i is the position of the index expression where beginning is used, and n is the total number of index expressions.

#### See also

Operator (), operator {}, beginning, end, matrixcol, matrixrow

#### end

Last index of an array.

# **Syntax**

```
v(...end...)
A(...end...)
function e = C::end(obj, i, n)
```

# **Description**

In an expression used as an index to access some elements of an array, end gives the index of the last element (line or column, depending of the context).

end can be overloaded for objects of used-defined classes. Its definition should be have a header equivalent to function e=C::end(obj,i,n), where C is the name of the class, obj is the object to be indexed, i is the position of the index expression where end is used, n is the total number of index expressions.

#### **Examples**

Last 2 elements of a vector:

```
a = 1:5; a(end-1:end)
4 5
```

Assignment to the last element of a vector:

```
a(end) = 99
a =
1 2 3 4 99
```

Extension of a vector:

```
a(end + 1) = 100
a =
1 2 3 4 99 100
```

#### See also

Operator (), operator {}, size, length, beginning, matrixcol, matrixrow

# global persistent

Declaration of global or persistent variables.

#### **Syntax**

```
global x y ...
persistent x y ...
```

### **Description**

By default, all variables are *local* and created the first time they are assigned to. Local variables can be accessed only from the body of the function where they are defined, but not by any other function, even the ones they call. They are deleted when the function exits. If the function is called recursively (i.e. if it calls itself, directly or indirectly), distinct variables are defined for each call. Similarly, local variables defined in the workspace using the command-line interface cannot be referred to in functions.

On the other hand, global variables can be accessed by multiple functions and continue to exist even after the function which created them exits. Global variables must be declared with global in each function which uses them. They can also be declared in the workspace. There exists only a single variable for each different name.

Declaring a global variable has the following result:

- If a previous local variable with the same name exists, it is deleted.
- If the global variable does not exist, it is created and initialized with the empty array [].
- Every access which follows the declaration in the same function or workspace uses the global variable.

Like global variables, *persistent variables* are preserved between function calls; but they cannot be shared between different functions. They are declared with persistent. They cannot be declared outside a function. Different persistent variables can have the same name in different functions.

# **Examples**

Functions to reset and increment a counter:

```
function reset
  global counter;
  counter = 0;

function value = increment
  global counter;
  counter = counter + 1;
  value = counter;
```

Here is how the counter can be used:

```
reset;
i = increment
i =
    1
j = increment
j =
    2
```

#### See also

function

### matrixcol

First index in a subscript expression.

# Syntax

```
A(...matrixcol...)
function e = C::matrixcol(obj, i, n)
```

# **Description**

In an expression used as a single subscript to access some elements of an array A(expr), matrixcol gives an array of the same size as A where each element is the column index. For instance for a 2-by-3 matrix, matrixcol gives the 2-by-3 matrix [1,2,3;1,2,3].

In an expression used as the second of multiple subscripts to access some elements of an array A(...,expr) or A(...,expr,...), matrixcol gives a row vector of length size(A,2) whose elements are the indices of each column. It is equivalent to the range (beginning:end).

matrixcol is useful in boolean expressions to select some elements of an array.

matrixcol can be overloaded for objects of used-defined classes. Its definition should have a header equivalent to function

e=C::matrixcol(obj,i,n), where C is the name of the class, obj is the object to be indexed, i is the position of the index expression where matrixcol is used, and n is the total number of index expressions.

#### **Example**

Set to 0 the NaN values which are not in the first column:

```
A = [1, nan, 5; nan, 7, 2; 3, 1, 2];
A(matrixcol > 1 & isnan(A)) = 0
A =

1 0 5

nan 7 2

3 1 2
```

#### See also

matrixrow, beginning, end

#### matrixrow

First index in a subscript expression.

# **Syntax**

```
A(...matrixrow...)
function e = C::matrixrow(obj, i, n)
```

### **Description**

In an expression used as a single subscript to access some elements of an array A(expr), matrixrow gives an array of the same size as A where each element is the row index. For instance for a 2-by-3 matrix, matrixrow gives the 2-by-3 matrix [1,1,1;2,2,2].

In an expression used as the first of multiple subscripts to access some elements of an array A(expr,...), matrixrow gives a row vector of length size(A,1) whose elements are the indices of each row. It is equivalent to the range (beginning:end).

matrixrow is useful in boolean expressions to select some elements of an array.

matrixrow can be overloaded for objects of used-defined classes. Its definition should be have a header equivalent to function e=C::matrixrow(obj,i,n), where C is the name of the class, obj is the object to be indexed, i is the position of the index expression where matrixrow is used, and n is the total number of index expressions.

#### See also

matrixcol, beginning, end

# subsasgn

Assignment to a part of an array, list, or structure.

### **Syntax**

```
A = subsasqn(A, s, B)
```

# **Description**

When an assignment is made to a subscripted part of an object in a statement like A(s1,s2,...)=B, LME executes A=subsasgn(A,s,B), where subsasgn is a method of the class of variable A and s is a structure with two fields: s.type which is '()', and s.subs which is the list of subscripts  $\{s1,s2,...\}$ . If a subscript is the colon character which stands for all elements along the corresponding dimensions, it is represented with the string ':' in s.subs.

When an assignment is made to a subscripted part of an object in a statement like  $A\{s\}=B$ , LME executes A=subsasgn(A,s,B), where subsasgn is a method of the class of variable A and s is a structure with two fields: s.type which is ' $\{\}$ ', and s.subs which is the list containing the single subscript  $\{s\}$ .

When an assignment is made to the field of an object in a statement like A.f=B, LME executes A=subsasgn(A,s,B), where s is a structure with two fields: s.type which is '.', and s.subs which is the name of the field ('f' in this case).

While the primary purpose of subsasgn is to permit the use of subscripts with objects, a built-in implementation of subsasgn is provided for arrays when s.type is '()', for lists and cell arrays when s.type is a list or a cell array, and for structures when s.type is '.'. In that case, the second argument s can be reduced to the list of subscripts or the field name; and a single subscripts can be given directly instead of a list of length 1.

# **Examples**

```
A = [1,2;3,4];
subsasgn(A, {type='()',subs={1,':'}}, 999)
999 999
3  4
subsasgn(A, {type='()',subs={':',1}}, [])
```

2

Same result when the indices are given directly as the second argument:

```
subsasgn(A, {1,':'}, 999)
    999 999
    3    4
s = {a=2, b=1:5};
subsasgn(s, 'b', 'abc')
    a: 2
    b: 'abc'
```

#### See also

Operator (), operator {}, subsref, beginning, end

#### subsref

Reference to a part of an array, list, or structure.

# **Syntax**

```
B = subsref(A, s)
```

# Description

When an object variable is subscripted in an expression like A(s1,s2,...), LME evaluates subsref(A,s), where subsref is a method of the class of variable A and s is a structure with two fields: s.type which is '()', and s.subs which is the list of subscripts {s1,s2,...}. If a subscript is the colon character which stands for all elements along the corresponding dimensions, it is represented with the string ':' in s.subs.

When an object variable is subscripted in an expression like  $A\{s\}$ , LME evaluates subsref(A,s), where subsref is a method of the class of variable A and s is a structure with two fields: s.type which is '{}', and s.subs which is the list containing the single subscript {s}.

When the field of an object variable is retrieved in an expression like A.f, LME executes subsref(A,s), where s is a structure with two fields: s.type which is '.', and s.subs which is the name of the field ('f' in this case).

While the primary purpose of subsref is to permit the use of subscripts with objects, a built-in implementation of subsref is provided for arrays when s.type is '()', for lists when s.type is '{}', and for

structures when s.type is '.'. In that case, the second argument s can be reduced to the list of subscripts or the field name; and a single subscripts can be given directly instead of a list of length 1.

### **Examples**

```
A = [1,2;3,4];
subsref(A, {type='()',subs={1,':'}})
1 2
```

Same result when the indices are given directly as the second argument:

```
subsref(A, {1,':'})
    1 2
s = {a='abc', b=1:5};
subsref(s, 'b')
    1 2 3 4 5
```

#### See also

Operator (), operator {}, subsasgn, beginning, end

# **6.13 Programming Constructs**

Programming constructs are the backbone of any LME program. Except for the variable assignment, all of them use reserved keywords which may not be used to name variables or functions. In addition to the constructs described below, the following keywords are reserved for future use:

```
classdef parfor
goto spmd
```

# break

Terminate loop immediately.

# **Syntax**

break

# **Description**

When a break statement is executed in the scope of a loop construct (while, repeat or for), the loop is terminated. Execution continues at the statement which follows end. Only the innermost loop where break is located is terminated.

The loop must be in the same function as break. It is an error to execute break outside any loop.

#### See also

while, repeat, for, continue, return

#### case

Conditional execution of statements depending on a number or a string.

#### See also

switch. otherwise

#### catch

Error recovery.

#### See also

try

#### continue

Continue loop from beginning.

#### **Syntax**

continue

# **Description**

When a continue statement is executed in the scope of a loop construct (while, repeat or for), statements following continue are ignored and a new loop is performed if the loop termination criterion is not fulfilled.

The loop must be in the same function as continue. It is an error to execute continue outside any loop.

#### See also

```
while, repeat, for, break
```

#### define

Definition of a constant.

# **Syntax**

```
define c = expr
define c = expr;
```

# **Description**

define c=expr assign permanently expression expr to c. It is equivalent to

```
function y = c
y = expr;
```

Since c does not have any input argument, the expression is usually constant. A semicolon may follow the definition, but it does not have any effect. define must be the first element of the line (spaces and comments are skipped).

# **Examples**

```
define e = exp(1);
define g = 9.81;
define c = 299792458;
define G = 6.672659e-11;
```

#### See also

function

#### for

Loop controlled by a variable which takes successively the value of the elements of a vector or a list.

```
for v = vect
    s1
    ...
end

for v = list
    s1
    ...
end
```

### **Description**

The statements between the for statement and the corresponding end are executed repeatedly with the control variable v taking successively every column of vect or every element of list list. Typically, vect is a row vector defined with the range operator.

You can change the value of the control variable in the loop; however, next time the loop is repeated, that value is discarded and the next column of vect is fetched.

# **Examples**

```
for i = 1:3; i, end
  i =
  i =
    2
  i =
for i = (1:3)'; i, end
  i =
    1
    2
for i = 1:2:5; end; i
for i = 1:3; break; end; i
  i =
for el = \{1, 'abc', \{2,5\}\}; el, end
  el =
    1
  el =
    abc
  el =
    {2,5}
```

#### See also

while, repeat, break, continue, variable assignment

# function endfunction

Definition of a function, operator, or method.

# **Syntax**

```
function f
    statements

function f(x1, x2, ...)
    statements

function f(x1, x2 = expr2, ...)
    statements

function y = f(x1, x2, ...)
    statements

function (y1,y2,...) = f(x1,x2,...)
    statements

function ... class::method ...
    statements

function ...
    statements

endfunction ...
    statements
```

# **Description**

New functions can be written to extend the capabilities of LME. They begin with a line containing the keyword function, followed by the list of output arguments (if any), the function name, and the list of input arguments between parenthesis (if any). The output arguments must be enclosed between parenthesis or square brackets if they are several. One or more variable can be shared in the list of input and output arguments. When the execution of the function terminates (either after the last statement or because of the command return), the current value of the output arguments, as set by the function's statements, is given back to the caller. All variables used in the function's statements are local; their value is undefined before the first assignment (and it is illegal to use them in an expression), and is not shared with variables in other functions or with recursive calls of the same

function. Different kinds of variables can be declared explicitly with global and persistent.

When multiple functions are defined in the same code source (for instance in a library), the body of a function spans from its header to the next function or until the endfunction keyword, whichever comes first. Function definitions cannot be nested. endfunction is required only when the function definition is followed by code to be executed outside the scope of any function. This includes mixed code and function definitions entered in one large entry in a command-line interface, or applications where code is mainly provided as statements, but where function definitions can help and separate libraries are not wished (note that libraries cannot contain code outside function definitions; they do not require endfunction). Both function and endfunction appear usually at the beginning of a line, but are also permitted after a semicolon or a comma.

### Variable number of arguments

Not all of the input and output arguments are necessarily specified by the caller. The caller fixes the number of input and output arguments, which can be obtained by the called function with nargin and nargout, respectively. Unspecified input arguments (from nargin+1 to the last one) are undefined, unless a default value is provided in the function definition: with the definition function f(x,y=2), y is 2 when f is called with a single input argument. Unused output arguments (from nargout+1 to the last one) do not have to be set, but may be.

Functions which accept an unspecified number of input and/or output arguments can use the special variables varargin and varargout, which are lists of values corresponding to remaining input and output arguments, respectively.

# Named arguments

The caller can pass some or all of the input arguments by name, such as f(x=2). Named arguments must follow unnamed ones. Their order does not have to match the order of the input arguments in the function declaration, and some arguments can be missing. Missing arguments are set to their default value if it exists, or left undefined. Undefined arguments can be detected with isdefined, or the error caused by their use caught by try.

Functions which accept unspecified named arguments or which do not want to expose the argument names used in their implementation can use the special variable namedargin, which is a structure containing all named arguments passed by the caller.

### **Unused arguments**

Character stands for an unused argument. It can be used as a placeholder for an input argument name in the function definition, or in the list of output arguments specified for the function call.

If function f is defined with function header function  $f(x,\tilde{\ })$ , it accepts two input arguments, the first one assigned to x and the second one discarded. This can be useful if f is called by code which expects a function with two input arguments.

In  $(a, \tilde{\ }, c)=f$ , function f is called to provide three output arguments (nargout==3), but the second output argument is discarded.

### Operator overloading

To redefine an operator (which is especially useful for object methods; see below), use the equivalent function, such as plus for operator +. The complete list is given in the section about operators.

To define a method which is executed when one of the input arguments is an object of class class (or a child in the classes hierarchy), add class:: before the method (function) name. To call it, use only the method name, not the class name.

# **Examples**

Function with optional input and output arguments:

```
function (Sum, Prod) = calcSumAndProd(x, y)
  if nargout == 0
                     % nothing to be computed
    return;
  if nargin == 0 % make something to be computed...
    x = 0;
 end
  if nargin <= 1
                      % sum of elements of x
    Sum = sum(x);
                      % sum of x and y
 else
    Sum = x + y;
 end
  if nargout == 2
                      % also compute the product
    if nargin == 1
                      % product of elements of x
      Prod = prod(x);
                      % product of x and y
      Prod = x \cdot * y;
    end
 end
```

Two equivalent definitions:

```
function S = area(a, b = a, ellipse = false)
  S = ellipse ? pi * a * b / 4 : a * b;
function S = area(a, b, ellipse)
  if ~isdefined(b)
    b = a;
  end
  if ~isdefined(ellipse)
    ellipse = false;
  end
  S = ellipse ? pi * a * b / 4 : a * b;
```

With unnamed arguments only, area can be called with values for a only, a and b, or a, b and ellipse. By naming ellipse, the second argument can be omitted:

```
S = area(2, ellipse=true)
S =
    3.1416
```

Function max can return the index of the maximum value in a vector. In the following call, the maximum itself is discarded.

```
(~, maxIndex) = max([2,7,3,5])
maxIndex =
2
```

#### See also

return, nargin, nargout, isdefined, varargin, varargout, namedargin, define, inline, global, persistent

# hideimplementation

Hide the implementation of remaining functions in a library.

# **Syntax**

hideimplementation

# **Description**

In a library, functions which are defined after the hideimplementation keyword have their implementation hidden: for errors occuring when they are executed, the error message is the same as if the function was a native function (it does not contain information about the error location in the function or subfunctions), and during debugging, dbstep in steps over the function call.

hideimplementation may not be placed in the same line of source code as any other command (comments are possible, though).

#### See also

public, private, function, use, error, dbstep

# if elseif else end

Conditional execution depending on the value of one or more boolean expressions.

# **Syntax**

```
if expr
    s1
end
if expr
    s1
    . . .
else
    s2
end
if expr1
    s1
    . . .
elseif expr2
    s2
else
    s3
end
```

# **Description**

If the expression following if is true (nonempty and all elements different from 0 and false), the statements which follow are executed. Otherwise, the expressions following elseif are evaluated, until one of them is true. If all expressions are false, the statements following else are executed. Both elseif and else are optional.

# **Example**

```
if x > 2
  disp('large');
elseif x > 1
```

```
disp('medium');
else
  disp('small');
end
```

#### See also

switch, while

#### include

Include libraries.

### **Syntax**

include lib

# **Description**

include lib inserts the contents of the library file lib. Its effect is similar to the use statement, except that the functions and constants in lib are defined in the same context as the library where include is located. Its main purpose is to permit to define large libraries in multiple files in a transparent way for the user. include statements must not follow other statements on the same line, and can reference only one library which is searched at the same locations as use. They can be used only in libraries.

Since LME replaces include with the contents of lib, one should be cautious about the public or private context which is preserved between the libraries. It is possible to include a fragment of function without a function header.

#### See also

use, includeifexists, private, public

#### includeifexists

Include library if it exists.

#### **Syntax**

includeifexists lib

# **Description**

includeifexists lib inserts the contents of the library file lib if it exists; if the library does not exists, it does nothing.

#### See also

include, useifexists, private, public

# otherwise

Conditional execution of statements depending on a number or a string.

#### See also

switch. case

# private

Mark the beginning of a sequence of private function definitions in a library.

### **Syntax**

private

# **Description**

In a library, functions which are defined after the private keyword are private. private may not be placed in the same line of source code as any other command (comments are possible, though).

In a library, functions are either public or private. Private functions can only be called from the same library, while public functions can also be called from contexts where the library has been imported with a use command. Functions are public by default.

# **Example**

Here is a library for computing the roots of a second-order polynomial. Only function roots2 can be called from the outside of the library.

```
private
function d = discr(a, b, c)
  d = b^2 - 4 * a * c;
public
```

```
function r = roots2(p)
  a = p(1);
  b = p(2);
  c = p(3);
  d = discr(a, b, c);
  r = [-b+sqrt(d); -b-sqrt(d)] / (2 * a);
```

#### See also

public, function, use

# public

Mark the beginning of a sequence of public function definitions in a library.

#### **Syntax**

public

### **Description**

In a library, functions which are defined after the public keyword are public. public may not be placed in the same line of source code as any other command (comments are possible, though).

In a library, functions are either public or private. Private functions can only be called from the same library, while public functions can also be called from contexts where the library has been imported with a use command. Functions are public by default: the public keyword is not required at the beginning of the library.

#### See also

private, function, use

# repeat

Loop controlled by a boolean expression.

# **Syntax**

```
repeat
s1
...
until expr
```

# **Description**

The statements between the repeat statement and the corresponding until are executed repeatedly (at least once) until the expression of the until statement yields true (nonempty and all elements different from 0 and false).

### **Example**

```
v = [];
repeat
  v = [v, sum(v)+1];
until v(end) > 100;
v
  1  2  4  8  16  32  64  128
```

#### See also

while, for, break, continue

#### return

Early return from a function.

# **Syntax**

return

# **Description**

return stops the execution of the current function and returns to the calling function. The current value of the output arguments, if any, is returned. return can be used in any control structure, such as if, while, or try, or at the top level.

# **Example**

```
function dispFactTable(n)
  % display the table of factorials from 1 to n
  if n == 0
    return; % nothing to display
  end
  fwrite(' i i!\n');
  for i = 1:n
    fwrite('%2d %3d\n', i, prod(1:i));
  end
```

#### See also

function

#### switch

Conditional execution of statements depending on a number or a string.

# **Syntax**

```
switch expr
    case e1
         s1
    case [e2,e3,...]
         s23
         . . .
    case {e4,e5,...}
         s45
    otherwise
         S0
         . . .
end
switch string
    case str1
         s1
         . . .
    case str2
         s2
    case {str3,str4,...}
         s34
         . . .
    otherwise
         S0
         . . .
end
```

# **Description**

The expression of the switch statement is evaluated. If it yields a number, it is compared successively to the result of the expressions of the case statements, until it matches one; then the statements which follow the case are executed until the next case, otherwise or end. If the case expression yields a vector or a list, a match occurs if the

switch expression is equal to any of the elements of the case expression. If no match is found, but otherwise is present, the statements following otherwise are executed. If the switch expression yields a string, a match occurs only in case of equality with a case string expression or any element of a case list expression.

### **Example**

```
switch option
  case 'arithmetic'
    m = mean(data);
  case 'geometric'
    m = prod(data)^(1/length(data));
  otherwise
    error('unknown option');
end
```

#### See also

case, otherwise, if

# try

Error recovery.

# **Syntax**

```
try
...
end

try
...
catch
...
end

try
...
catch e
...
end
```

# **Description**

The statements after try are executed. If an error occurs, execution is switched to the statements following catch, if any, or to the state-

ments following end. If catch is followed by a variable name, a structure describing the error (the result of lasterror) is assigned to this variable; otherwise, the error message can be retrieved with lasterr or lasterror. If no error occurs, the statements between try and end are ignored.

try ignores two errors:

- the interrupt key (Control-Break on Windows, Command-. on macOS, Control-C on other operating systems with a keyboard, timeout in Sysquake Remote);
- an attempt to execute an untrusted function in a sandbox. The error can be handled only outside the sandbox.

### **Examples**

```
a = 1;
a(2), 555
   Index out of range 'a'
try, a(2), end, 555
   555
try, a(2), catch, 333, end, 555
   333
   555
try, a, catch, 333, end, 555
a =
   1
   555
```

#### See also

lasterr, lasterror, error

#### until

End of repeat/until loop.

#### See also

repeat

#### use

Import libraries.

```
use lib
use lib1, lib2, ...
```

### **Description**

Functions can be defined in separate files, called *libraries*. use makes them available in the current context, so that they can be called by the functions or statements which follow. Using a library does not make available functions defined in its sublibraries; however, libraries can be used multiple times, in each context where their functions are referenced.

All use statements are parsed before execution begins. They can be placed anywhere in the code, typically before the first function. They cannot be skipped by placing them after an if statement. Likewise, try/catch cannot be used to catch errors; useifexists should be used if the absence of the library is to be ignored.

#### See also

useifexists, include, function, private, public, info

# useifexists

Import libraries if they exist.

# **Syntax**

```
useifexists lib useifexists lib1, lib2, ...
```

# Description

useifexists has the same syntax and effect as use, except that libraries which are not found are ignored without error.

#### See also

use, include, function, private, public, info

#### while

Loop controlled by a boolean expression.

```
while expr
s1
...
```

# **Description**

The statements between the while statement and the corresponding end are executed repeatedly as long as the expression of the while statement yields true (nonempty and all elements different from 0 and false).

If a break statement is executed in the scope of the while loop (i.e. not in an enclosed loop), the loop is terminated.

If a continue statement is executed in the scope of the while loop, statements following continue are ignored and a new loop is performed if the while statement yields true.

### **Example**

```
e = 1;
i = 2;
while true % forever
  eNew = (1 + 1/i) ^ i;
  if abs(e - eNew) < 0.001
    break;
  end
  e = eNew;
  i = 2 * i;
end
e
2.717</pre>
```

#### See also

repeat, for, break, continue, if

# 6.14 Miscellaneous Functions

This section describes functions related to programming: function arguments, error processing, evaluation, memory.

#### assert

Check that an assertion is true.

```
assert(expr)
assert(expr, str)
assert(expr, format, arg1, arg2, ...)
assert(expr, identifier, format, arg1, arg2, ...)
```

### **Description**

assert(expr) checks that expr is true and throws an error otherwise. Expression expr is considered to be true if it is a non-empty array whose elements are all non-zero.

With more input arguments, assert checks that expr is true and throws the error specified by remaining arguments otherwise. These arguments are the same as those expected by function error.

When the intermediate code is optimized, assert can be ignored. It should be used only to produce errors at an early stage or as a debugging aid, not to trigger the try/catch mechanism. The expression should not have side effects. The most common use of assert is to check the validity of input arguments.

### **Example**

```
function y = fact(n)
  assert(length(n)==1 && isreal(n) && n==round(n), 'LME:nonIntArg');
  y = prod(1:n);
```

#### See also

error, warning, try

#### builtin

Built-in function evaluation.

# **Syntax**

```
(argout1, ...) = builtin(fun, argin1, ...)
```

# **Description**

(y1, y2, ...)=builtin(fun,x1,x2,...) evaluates the built-in function fun with input arguments x1, x2, etc. Output arguments are assigned to y1, y2, etc. Function fun is specified by its name as a string.

builtin is useful to execute a built-in function which has been redefined.

#### **Example**

Here is the definition of operator plus so that it can be used with character strings to concatenate them.

```
function r = plus(a, b)
  if ischar(a) && ischar(b)
    r = [a, b];
  else
    r = builtin('plus', a, b);
  end
```

The original meaning of plus for numbers is preserved:

```
1 + 2
3
'ab' + 'cdef'
abcdef
```

#### See also

feval

# clear

Discard the contents of a variable.

# **Syntax**

```
clear
clear(v1, v2, ...)
clear -functions
```

# **Description**

Without argument, clear discards the contents of all the local variables, including input arguments. With string input arguments, clear(v1,v2,...) discards the contents of the enumerated variables. Note that the variables are specified by strings; clear is a normal function which evaluates its arguments if they are enclosed between parenthesis. You can also omit parenthesis and quotes and use command syntax.

clear is usually not necessary, because local variables are automatically discarded when the function returns. It may be useful if a large variable is used only at the beginning of a function, or at the command-line interface.

clear -functions or clear -f removes the definition of all functions. It can be used only from the command-line interface, not in a function.

### **Examples**

In the example below, clear(b) evaluates its argument and clears the variable whose name is 'a'; clear b, without parenthesis and quotes, does not evaluate it; the argument is the literal string 'b'.

```
a = 2;
b = 'a';
clear(b)
a
   Undefined variable 'a'
b
   a
clear b
b
   Undefined variable b
```

#### See also

variable assignment, isdefined

### deal

Copy input arguments to output arguments.

# **Syntax**

```
(v1, v2, ...) = deal(e)
(v1, v2, ...) = deal(e1, e2, ...)
```

# Description

With a single input argument, deal provides a copy of it to all its output arguments. With multiple input arguments, deal provides them as output arguments in the same order.

deal can be used to assign a value to multiple variables, to swap the contents of two variables, or to assign the elements of a list to different variables.

# **Examples**

Swap variable a and b:

```
a = 2;
b = 'abc';
(a, b) = deal(b, a)
a =
```

```
abc
b =
```

Copy the same random matrix to variables x, y, and z:

```
(x, y, z) = deal(rand(5));
```

Assign the elements of list 1 to variables v1, v2, and v3:

```
l = {1, 'abc', 3:5};
(v1, v2, v3) = deal(l{:})
v1 =
    1
    v2 =
    abc
    v3 =
    3 4 5
```

#### See also

varargin, varargout, operator {}

# dumpvar

Dump the value of an expression as an assignment to a variable.

# **Syntax**

```
dumpvar(value)
dumpvar(name, value)
dumpvar(fd, name, value)
str = dumpvar(value)
str = dumpvar(name, value)
... = dumpvar(..., fd=fd, NPrec=nPrec)
```

# **Description**

dumpvar(fd, name, value) writes to the channel fd (the standard output by default) a string which would set the variable name to value, if it was evaluated by LME. If name is omitted, only the textual representation of value is written. A file descriptor can also be specified as a named argument fd.

With an output argument, dumpvar stores result into a string and produces no output.

In addition to fd, dumpvar also accepts named argument NPrec for the maximum number of digits in floating-point numbers.

### **Examples**

```
dumpvar(2+3)
5
a = 6; dumpvar('a', a)
a = 6;
s = 'abc'; dumpvar('string', s)
    string = 'abc';
dumpvar('x', 1/3, NPrec=5)
    x = 0.33333;
```

### See also

```
fprintf, sprintf, str2obj
```

### error

Display an error message and abort the current computation.

### **Syntax**

```
error(str)
error(format, arg1, arg2, ...)
error(identifier, format, arg1, arg2, ...)
error(identifier)
error(..., throwAsCaller=b)
```

# **Description**

Outside a try block, error(str) displays string str as an error message and the computation is aborted. With more arguments, error use the first argument as a format string and displays remaining arguments accordingly, like fprintf.

In a try block, error(str) throws a user error without displaying anything.

An error identifier can be added in front of other arguments. It is a string made of at least two segments separated by semicolons. Each segment has the same syntax as variable or function name (i.e. it begins with a letter or an underscore, and it continues with letters, digits and underscores.) The identifier can be retrieved with lasterr or lasterror in the catch part of a try/catch construct and helps to identify the error. For errors thrown by LME built-in functions, the first segment is always LME.

The identifier of an internal error (an error which can be thrown by an LME builti-in function, such as 'LME:indexOutOfRange'), can be used as the only argument; then the standard error message is displayed.

error also accepts a boolean named argument throwAsCaller. If it is true, the context of the error is changed so that the function calling error appears to throw the error itself. It is useful for fully debugged functions whose internal operation can be hidden. Keyword hideimplementation has a similar effect at the level of a library, by hiding the internal error handling in all its functions.

## **Examples**

```
error('Invalid argument.');
   Invalid argument.
o = 'ground';
error('robot:hit', 'The robot is going to hit %s', o);
   The robot is going to hit ground
lasterror
   message: 'The robot is going to hit ground'
   identifier: 'robot:hit'
```

Definition of a function which checks its input arguments, and a test function which calls it:

```
function xmax = largestRoot(a, b, c)
  // largest root of a x^2 + b x + c = 0
  if b^2 - 4 * a * c < 0
    error('No real root', throwAsCaller=true);
  end
  xmax = (-b + sqrt(b^2 - 4 * a * c)) / (2 * a);
function test
  a = largestRoot(1,1,1);</pre>
```

Error message:

```
test
No real root (test;8)
```

Error message without throwAsCaller=true in the definition of largestRoot:

```
test
No real root (largestRoot;4)
-> test;8
```

#### See also

warning, try, lasterr, lasterror, assert, fprintf, hideimplementation

### eval

Evaluate the contents of a string as an expression or statements.

### **Syntax**

```
x = eval(str_expression)
eval(str_statement)
```

## **Description**

If eval has output argument(s), the input argument is evaluated as an expression whose result(s) is returned. Without output arguments, the input argument is evaluated as statement(s). eval can evaluate and assign to existing variables, but cannot create new ones.

## **Examples**

```
eval('1+2')
3
a = eval('1+2')
a = 3
eval('a=2+3')
a = 5
```

### See also

feval

### exist

Existence of a function or variable.

# **Syntax**

```
b = exist(name)
b = exist(name, type)
```

# **Description**

exist returns true if its argument is the name of an existing function or variable, or false otherwise. A second argument can restrict the lookup to builtin functions ('builtin'), user functions ('function'), or variables ('variable').

# **Examples**

```
exist('sin')
  true
exist('cos', 'function')
  false
```

info, isdefined

## feval

Function evaluation.

## **Syntax**

```
(argout1,...) = feval(fun,argin1,...)
```

### **Description**

(y1, y2, ...)=feval(fun, x1, x2, ...) evaluates function fun with input arguments x1, x2, etc. Output arguments are assigned to y1, y2, etc. Function fun is specified by either its name as a string, a function reference, or an anonymous or inline function.

If a variable f contains a function reference or an anonymous or inline function, f(arguments) is equivalent to feval(f,arguments).

### **Examples**

```
y = feval('sin', 3:5)
y =
    0.1411 -0.7568 -0.9589
y = feval(@(x) sin(2*x), 3:5)
y =
    -0.2794 0.9894 -0.544
fun = @(x) sin(2*x);
y = fun(3:5)
y =
    -0.2794 0.9894 -0.544
```

### See also

builtin, eval, fevalx, apply, inline, operator @

## fun2str

Name of a function given by reference or source code of an inline function.

### **Syntax**

```
str = fun2str(funref)
str = fun2str(inlinefun)
```

## **Description**

fun2str(funref) gives the name of the function whose reference is funref.

fun2str(inlinefun) gives the source code of the inline function
inlinefun.

## **Examples**

```
fun2str(@sin)
  sin
fun2str(inline('x+2*y', 'x', 'y'))
  function y=f(x,y);y=x+2*y;
```

### See also

operator @, str2fun

### info

Information about LME.

## **Syntax**

```
info
info builtin
info errors
info functions
info global
info libraries
info methods
info operators
info persistent
info size
info threads
info usedlibraries
info variables
info(kind, fd=fd)
str = info
SA = info(kind)
```

### **Description**

info displays the language version. With an output argument, the language version is given as a string.

info builtin displays the list of built-in functions with their module name (modules are subsets of built-in functions). A letter u is displayed after each untrusted function (functions which cannot be executed in the sandbox). With an output argument, info('builtin') gives a structure array which describes each built-in function, with the following fields:

name function name module module name

trusted true if the function is trusted

info operators displays the list of operators. With an output argument, info('operators') gives a list of structures, like info('builtin').

info functions displays the list of user-defined functions with the library where they are defined and the line number in the source code. Parenthesis denote functions known by LME, but not loaded; they also indicate spelling errors in function or variable names. With an output argument, info('functions') gives a structure array which describes each user-defined function, with the following fields:

library library name name function name loaded true if loaded

line line number if available, or []

info methods displays the list of methods. With an output argument, info('methods') gives a structure array which describes each method, with the following fields:

library library name class class name name function name loaded true if loaded

line line number if available, or []

info variables displays the list of variables with their type and size. With an output argument, info('variables') gives a structure array which describes each variable, with the following fields:

name function name defined true if defined

info global displays the list of all global variables. With an output argument, info('global') gives the list of the global variable names.

info persistent displays the list of all persistent variables. With an output argument, info('persistent') gives the list of the persis-

tent variable names.

info libraries displays the list of all loaded libraries with the libraries they have loaded with use. The base context in which direct commands are evaluated is displayed as (base); it is not an actual library and contains no function definition. With an output argument, info('libraries') gives a structure array with the following fields:

```
library library name, or '(base)' sublibraries list of sublibraries
```

info usedlibraries displays the list of libraries available in the current context. With an output argument, info('usedlibraries') gives the list of the names of these libraries.

info errors displays the list of error messages. With an output argument, info('errors') gives a structure array which describes each error message, with the following fields:

```
id error ID msg error message
```

info size displays the size in bytes of integer numbers (as used for indices and most internal computations), double numbers, single numbers, and pointers; the byte ordering in multibyte values (little-endian if the least-significant byte comes first, else big-endian), and whether arrays are stores column-wise or row-wise. With an output argument, info('size') gives them in a structure of six fields:

```
int integer size
double double size
single single size (or 0)
ptr pointer size
be true if big-endian
columnwise true for column-wise array layout
```

info threads displays the ID of all threads. With an output argument, info('threads') gives a structure array which describes each thread, with the following fields:

```
id thread ID
totaltime execution time in seconds
```

Only the first character of the argument is meaningful; info b is equivalent to info builtin.

A named argument fd can specify the output channel; in that case, the command syntax cannot be used.

# **Examples**

```
info
LME 5.2
info s
```

```
int: 4 bytes
double: 8 bytes
ptr: 4 bytes
little endian
array layout: row-wise
info b
  LME/abs
  LME/acos
  LME/acosh
  (etc.)
info v
  ans (1x1 complex)
vars = info('v')
  var =
  2x1 struct array (2 fields)
```

List of variables displayed on channel 2 (standard error channel):

```
info('v', fd=2)
```

Library hierarchy in the command-line interface:

```
use lti
info l
  (base): _cli, lti
  _cli: lti
  lti: polynom
  polynom
```

The meaning is as follows: (base) is the context where commands are evaluated; functions defined from the command-line interface, stored in \_cli, and in lti can be called from there. Functions defined from the command-line interface also have access to the definitions of lti. Library lti uses library polynom, but functions defined in polynom cannot be called directly from commands (polynom does not appear as a sublibrary of (base) or \_cli). Finally, library polynom does not import a sublibrary itself.

#### See also

inmem, which, exist, use

# isequal

Comparison.

### Syntax

```
b = isequal(a, b, ...)
```

### **Description**

isequal compares its input arguments and returns true if all of them are equal, and false otherwise. Two numeric, logical and/or char arrays are considered to be equal if they have the same size and if their corresponding elements have the same value; an array which has at least one NaN (not a number) element is not equal to any other array. Two lists, cell arrays, structures or structure arrays are equal if the corresponding elements or fields are equal. Structure fields do not have to be in the same order.

isequal differs from operator == in that it results in a scalar logical value and arrays do not have to have the same size. It differs from operator === in that it does not require the type or the structure field order to agree, and in the way NaN is interpreted.

### See also

```
operator ==, operator ===
```

### inline

Creation of inline function.

# Syntax

```
fun = inline(funstr)
fun = inline(expr)
fun = inline(expr, arg1, ...)
fun = inline(funstr, param)
fun = inline(expr, arg1, ..., paramstruct)
fun = inline(expr, ..., true)
```

# **Description**

Inline function are LME objects which can be evaluated to give a result as a function of their input arguments. Contrary to functions declared with the function keyword, inline functions can be assigned to variables, passed as arguments, and built dynamically. Evaluating them with feval is faster than using eval with a string, because they are compiled only once to an intermediate code. They can also be used as the argument of functions such as fzero and fmin.

inline(funstr) returns an inline function whose source code is funstr. Input argument funstr follows the same syntax as a plain function. The function name is ignored.

inline(expr) returns an inline function with one implicit input argument and one result. The input argument expr is a string which

evaluates to the result. The implicit input argument of the inline function is a symbol made of a single lower-case letter different from i and j, such as x or t, which is found in expr. If several such symbols are found, the one closer to x in alphabetical order is picked.

inline(expr,arg1,...) returns an inline function with one result and the specified arguments arg1 etc. These arguments are also given as strings.

Inline functions also accept an additional input argument which correspond to fixed parameters provided when the function is executed. inline(funstr,param), where funstr is a string which contains the source code of a function, stores param together with the function. When the function is called, param is prepended to the list of input arguments.

inline(expr,args...,paramstruct) is a simplified way to create an inline function when the code consists of a single expression. args is the names of the arguments which must be supplied when the inline function is called, as strings; paramstruct is a structure whose fields define fixed parameters.

inline(expr,...,true) defines a function which can return as many output arguments as what feval (or other functions which call the inline function) expects. Argument expr must be a function call itself.

Anonymous functions created with operator @ are an alternative, often easier way of creating inline functions. The result is the same. Since inline is a normal function, it must be used in contexts where fixed parameters cannot be created as separate variables.

## **Examples**

A simple expression, evaluated at x=1 and x=2:

```
fun = inline('cos(x)*exp(-x)');
y = feval(fun, 2)
y =
     -5.6319e-2
y = feval(fun, 5)
y =
     1.9113e-3
```

A function of x and y:

```
fun = inline('exp(-x^2-y^2)', 'x', 'y');
```

A function with two output arguments (the string is broken in three lines to have a nice program layout):

```
'b=prod(v)^(1/length(v));']);
(am, gm) = feval(fun, 1:10)
   am =
     5.5
   gm =
     4.5287
```

Simple expression with fixed parameter a:

```
fun = inline('cos(a*x)', 'x', struct('a',2));
feval(fun, 3)
   0.9602
```

An equivalent function where the source code of a complete function is provided:

```
fun = inline('function y=f(a,x); y=cos(a*x);', 2);
feval(fun, 3)
  0.9602
```

The same function created with the anonymous function syntax:

```
a = 2;
fun = @(x) cos(a*x);
```

A function with two fixed parameters a and b whose values are provided in a list:

```
inline('function y=f(p,x);(a,b)=deal(p{:});y=a*x+b;',{2,3})
```

An inline function with a variable number of output arguments:

```
fun = inline('eig(exp(x))',true);
e = feval(fun, magic(2))
  e =
    -28.1440
    38.2514
(V,D) = feval(fun, magic(2))
  V =
    -0.5455    -0.4921
    0.8381    -0.8705
D =
    -28.1440     0.0000
    0.0000     38.2514
```

### See also

function, operator @, feval, eval

### inmem

List of functions loaded in memory.

## **Syntax**

```
inmem
SA = inmem
```

## **Description**

inmem displays the list of user-defined functions loaded in memory with the library where they are defined. With an output argument, inmem gives the result as a structure array which describes each user-defined function loaded in memory, with the following fields:

```
library library name
class class name ('' for functions)
name function name
```

### See also

info, which

### isdefined

Check if a variable is defined.

## Syntax

```
isdefined(var)
```

# **Description**

isdefined(var) returns true if variable var is defined, and false otherwise. Unlike ordinary functions, isdefined's argument must be a variable known to LME, referenced by name without quotes, and not an arbitrary expression. A variable is undefined in the following circumstances:

- function input argument when the function call does not supply enough values;
- function output argument which has not been assigned to, in the function itself, not in a function call;
- function local variable before its first assignment;

 function local variable after it has been cleared with function clear.

At command-line interface, clear usually discards completely variables.

## **Example**

Let function f be defined as

```
function f(x)
  if isdefined(x)
    disp(x);
  else
    disp('Argument x is not defined.');
  end

Then

f
  Argument x is not defined.
f(3)
  3
```

### See also

nargin, exist, which, clear, function

### isfun

Test for an inline function or function reference.

# **Syntax**

```
b = isfun(obj)
```

# **Description**

isfun(obj) returns true if obj is an inline function or a function reference, or false otherwise.

### See also

```
isa, class, fun2str
```

# isglobal

Test for the existence of a global variable.

### **Syntax**

```
b = isglobal(str)
```

### **Description**

isglobal(str) returns true if the string str is the name of a global variable, defined as such in the current context.

### See also

info, exist, isdefined, which

# iskeyword

Test for a keyword name.

## **Syntax**

```
b = iskeyword(str)
list = iskeyword
```

# **Description**

iskeyword(str) returns true if the string str is a reserved keyword which cannot be used as a function or variable name, or false otherwise. Keywords include if and global, but not the name of built-in functions like sin or i.

Without input argument, iskeyword gives the list of all keywords.

## **Examples**

```
iskeyword('otherwise')
  true
iskeyword
  {'break','case','catch','continue','else','elseif',
   'end','endfunction','for','function','global',
   'hideimplementation','if','otherwise','persistent',
   'private','public','repeat','return','switch','try',
   'until','use','useifexists','while'}
```

info, which

### ismac

Check whether computer runs under macOS.

## **Syntax**

b = ismac

## **Description**

ismac returns true on macOS, false on other platforms.

### See also

isunix, ispc

# ispc

Check whether platform is a PC.

# **Syntax**

b = ispc

# **Description**

ispc returns true on Windows, false on other platforms.

## See also

isunix, ismac

### isunix

Check whether computer runs under unix.

# **Syntax**

b = isunix

## **Description**

isunix returns true on unix platforms (including Mac OS X and unix-like), false on other platforms.

### See also

```
ispc, ismac
```

### lasterr

Last error message.

# **Syntax**

```
msg = lasterr
(msg, identifier) = lasterr
```

## **Description**

lasterr returns a string which describes the last error. With two output arguments, it also gives the error identifier. It can be used in the catch part of the try construct.

## **Example**

```
x = 2;
x(3)
  Index out of range
(msg, identifier) = lasterr
  msg =
    Index out of range
  identifier =
    LME:indexOutOfRange
```

### See also

lasterror, try, error

## lasterror

Last error structure.

### **Syntax**

```
s = lasterror
```

### **Description**

lasterror returns a structure which describes the last error. It contains the following fields:

```
identifier string short tag which identifies the error message string error message
```

The structure can be used as argument to rethrow in the catch part of a try/catch construct to propagate the error further.

## **Example**

```
x = 2;
x(3)
  Index out of range
lasterror
  message: 'Index out of range'
  identifier: 'LME:indexOutOfRange'
```

### See also

lasterr, try, rethrow, error

# namedargin

Named input arguments.

# **Syntax**

```
function \dots = fun(\dots, namedargin)
```

# Description

namedargin is a special variable which can be used to collect named input arguments. In the function declaration, it must be used as the last (or unique) input argument. When the function is called with named arguments, all of them are collected and stored in namedargin as a structure, where field names correspond to the argument names. With namedargin, there is no matching between the named arguments and the argument names in the function declaration. If the function is called without any named argument, namedargin is set to an empty structure.

In the body of the function, namedargin is a normal variable. Its fields can be accessed with the dot notation namedargin.name or namedargin.(name). All functions using structures can be used, such

as fieldnames or isfield. namedargin can also be modified or assigned to any value of any type.

When both varargin (for a variable number of unnamed arguments) and namedargin are used in the same function, they must be the last-but-one and the last arguments in the function declaration, respectively.

## **Example**

Here is a function which calculates the volume of a solid of revolution defined by a function y=f(x) between x=a and x=b, rotating around y=0. It accepts the same options as integral, given as a single option argument, as named values or both.

```
function V = solidRevVolume(fun, a, b, opt=struct, namedargin)
  opt = structmerge(opt, namedargin);
  V = pi * integral(@(x) fun(x)^2, a, b, opt);
```

It can be called without any option (opt is set to its default value, an empty structure):

```
cyl = solidRevVolume(@(x) 1, 0, 1)

cyl = 3.1416

cone = solidRevVolume(@(x) x, 0, 2, RelTol=1e-4)

cone = 8.3776
```

### See also

varargin, function, struct, fieldnames, structmerge, operator.

# nargin

Number of input arguments.

# Syntax

```
n = nargin
n = nargin(fun)
```

# **Description**

Calling a function with less arguments than what the function expects is permitted. In this case, the trailing variables are not defined. The function can use the nargin function to know how many arguments were passed by the caller to avoid accessing the undefined variables. Named arguments (arguments passed as name=value by the caller) are not included in the count.

Note that if you want to have an optional argument before the end of the list, you have to interpret the meaning of the variables yourself. LME always sets the nargin first arguments.

There are two other ways to let a function accept a variable number of input arguments: to define default values directly in the function header, or to call varargin to collect some or all of the input arguments in a list.

With one argument, nargin(fun) returns the (maximum) number of input arguments a function accepts. fun can be the name of a built-in or user function, a function reference, or an inline function. Functions with a variable number of input arguments (such as fprintf) give -1.

### **Examples**

A function with a default value (pi) for its second argument:

```
function x = multiplyByScalar(a,k)
if nargin < 2 % multiplyByScalar(x)
   k = pi; % same as multiplyByScalar(x,pi)
end
x = k * a;</pre>
```

A function with a default value (standard output) for its first argument. Note how you have to interpret the arguments.

```
function fprintstars(fd,n)
if nargin == 1 % fprintstars(n) to standard output
  fprintf(repmat('*',1,fd)); % n is actually stored in fd
else
  fprintf(fd, repmat('*',1,n));
end
```

Number of input arguments of function plus (usually called as the infix operator "+"):

```
nargin('plus')
2
```

#### See also

nargout, varargin, isdefined, function

# nargout

Number of output arguments.

### **Syntax**

```
n = nargout
n = nargout(fun)
```

### **Description**

A function can be called with between 0 and the number of output arguments listed in the function definition. The function can use nargout to check whether some output arguments are not used, so that it can avoid computing them or do something else.

With one argument, nargout (fun) returns the (maximum) number of output arguments a function can provide. fun can be the name of a built-in or user function, a function reference, or an inline function. Functions with a variable number of output arguments (such as feval) give -1.

## Example

A function which prints nicely its result when it is not assigned or used in an expression:

```
function y = multiplyByTwo(x)
if nargout > 0
  y = 2 * x;
else
  fprintf('The double of %f is %f\n', x, 2*x);
end
```

Maximum number of output arguments of svd:

```
nargout('svd')
3
```

### See also

nargin, varargout, function

## rethrow

Throw an error described by a structure.

# Syntax

```
rethrow(s)
rethrow(s, throwAsCaller=b)
```

### **Description**

rethrow(s) throws an error described by structure s, which contains the same fields as the output of lasterror. rethrow is typically used in the catch part of a try/catch construct to propagate further an error; but it can also be used to initiate an error, like error.

rethrow also accepts a boolean named argument throwAsCaller. If it is true, the context of the error is changed so that the function calling rethrow appears to throw the error itself. It is useful for fully debugged functions whose internal operation can be hidden.

## **Example**

The error whose identifier is 'LME:indexOutOfRange' is handled by catch; other errors are not.

```
try
...
catch
  err = lasterror;
  if err.identifier === 'LME:indexOutOfRange'
    ...
  else
    rethrow(err);
  end
end
```

### See also

lasterror, try, error

### str2fun

Function reference.

# **Syntax**

```
funref = str2fun(str)
```

# **Description**

str2fun(funref) gives a function reference to the function whose name is given in string str. It has the same effect as operator @, which is preferred when the function name is fixed.

### **Examples**

```
str2fun('sin')
    @sin
@sin
    @sin
a = 'cos';
str2fun(a)
    @cos
```

### See also

operator @, fun2str

# str2obj

Convert to an object its string representation.

### **Syntax**

```
obj = str2obj(str)
```

# Description

str2obj(str) evaluates string str and gives its result. It has the inverse effect as dumpvar with one argument. It differs from eval by restricting the syntax it accepts to literal values and to the basic constructs for creating complex numbers, arrays, lists, structures, objects, and other built-in types.

## **Examples**

eval, dumpvar

# varargin

Remaining input arguments.

### **Syntax**

```
function ... = fun(..., varargin)
function ... = fun(..., varargin, namedargin)
l = varargin
```

## **Description**

varargin is a special variable which can be used to collect input arguments. In the function declaration, it must be used after the normal input arguments; if namedargin is also present, varargin immediately precedes it. When the function is called with more arguments than what can be assigned to the other arguments, remaining ones are collected in a list and assigned to varargin. In the body of the function, varargin is a normal variable. Its elements can be accessed with the brace notation varargin{i}. nargin is always the total number of arguments passed to the function by the caller.

When the function is called with fewer arguments than what is declared, varargin is set to the empty list, {}.

## **Example**

Here is a function which accepts any number of square matrices and builds a block-diagonal matrix:

```
function M = blockdiag(varargin)
M = [];
for block = varargin
   // block takes the value of each input argument
   (m, n) = size(block);
   M(end+1:end+m,end+1:end+n) = block;
end
```

In the call below, varargin contains the list {ones(3),2\*ones(2),3}.

```
blockdiag(ones(3), 2*ones(2), 3)
     1
            1
                   1
                          0
                                 0
                                       0
                   1
     1
            1
                          0
                                 0
                                       0
     1
            1
                   1
                                        0
                          0
```

0	0	0	2	2	0
0 0 0	0	0	2 2 0	2	0 0 3
0	0	0	0	0	3

nargin, namedargin, varargout, function

## varargout

Remaining output arguments.

## **Syntax**

```
function (..., varargout) = fun(...)
varargout = ...
```

## **Description**

varargout is a special variable which can be used to dispatch output arguments. In the function declaration, it must be used as the last (or unique) output argument. When the function is called with more output arguments than what can be obtained from the other arguments, remaining ones are extracted from the list varargout. In the body of the function, varargout is a normal variable. Its value can be set globally with the brace notation {...} or element by element with varargout{i}. nargout can be used to know how many output arguments to produce.

# **Example**

Here is a function which differentiates a vector of values as many times as there are output arguments:

```
function varargout = multidiff(v)
  for i = 1:nargout
    v = diff(v);
    varargout{i} = v;
  end
```

In the call below, [1,3,7,2,5,3,1,8] is differentiated four times.

```
(v1, v2, v3, v4) = multidiff([1,3,7,2,5,3,1,8])
v1 =
     2
           4
                - 5
                       3
                            - 2
                                  - 2
                                         7
v2 =
     2
          - 9
                8
                      - 5
                             0
                                   9
```

nargout, varargin, function

## variables

Contents of the variables as a structure.

### **Syntax**

```
v = variables
```

## **Description**

variables returns a structure whose fields contain the variables defined in the current context.

## **Example**

```
a = 3;
b = 1:5;
variables
a: 3
b: real 1x5
```

### See also

info

# warning

Write a warning to the standard error channel.

# Syntax

```
warning(msg)
warning(format, arg1, arg2, ...)
```

## **Description**

warning(msg) displays the string msg. It should be used to notify the user about potential problems, *not* as a general-purpose display function.

With more arguments, warning uses the first argument as a format string and displays remaining arguments accordingly, like fprintf.

### **Example**

```
warning('Doesn\'t converge.');
```

error, disp, fprintf

### which

Library where a function is defined.

## **Syntax**

```
fullname = which(name)
```

# **Description**

which(name) returns an indication of where function name is defined. If name is a user function or a method prefixed with its class and two colons, the result is name prefixed with the library name and a slash. If name is a built-in function, it is prefixed with (builtin); a variable, with (var); and a keyword, with (keyword). If name is unknown, which returns the empty string.

# **Examples**

```
which logspace
  stdlib/logspace
which polynom::plus
  polynom/polynom::plus
which sin
  (builtin)/sin
x = 2;
which x
  (var)/x
```

info, isdefined

# 6.15 Sandbox Function

### sandbox

Execute untrusted code in a secure environment.

## **Syntax**

```
sandbox(str)
sandbox(str, varin)
varout = sandbox(str)
varout = sandbox(str, varin)
```

## **Description**

sandbox(str) executes the statements in string str. Functions which might do harm if used improperly are disabled; they include those related to the file system, to devices and to the network. Global and persistent variables are forbidden as well; but local variables can be created. The same restrictions apply to functions called directly or indirectly by statements in str. The purpose of sandbox is to permit the evaluation of code which comes from untrusted sources, such as the Internet

sandbox(str,varin) evaluates the statements in string str in a context with local variables equal to the fields of structure varin.

With an output argument, sandbox collects the contents of all variables in the fields of a single structure.

An error is thrown when the argument of sandbox attempts to execute one of the functions which are disabled. This error can be caught by a try/catch construct outside sandbox, but not inside its argument, so that unsuccessful attempts to circumvent the sandbox are always reported to the appropriate level.

# **Examples**

Evaluation of two assignments; the second value is displayed, and the variables are discarded at the end of the evaluation.

```
sandbox('a=2; b=3:5');
b =
   3 4 5
```

Evaluation of two assignments; the contents of the variables are stored in structure result.

```
result = sandbox('a=2; b=3:5;')
  result =
    a: 2
    b: real 1x3
```

Evaluation with local variables x and y initialized with the field of a structure. Variable z is local to the sandbox.

```
in.x = 12;
in.y = 1:10;
sandbox('z = x + y', in);
z =
    13 14 15 16 17 18 19 20 21 22
```

Attempt to execute the untrusted function fopen and to hide it from the outside. Both attempts fail: fopen is trapped and the security violation error is propagated outside the sandbox.

```
sandbox('try; fd=fopen('/etc/passwd'); end');
Security violation 'fopen'
```

### See also

sandboxtrust, eval, variables

### sandboxtrust

Escape the sandbox restrictions.

### **Syntax**

```
sandboxtrust(fun)
```

## **Description**

sandboxtrust(fun) sets a flag associated with function fun so that fun is executed without restriction, even when called from a sandbox. All functions called directly or indirectly from a trusted function are executed without restriction, except if a nested call to sandbox is performed. Argument fun can be a function reference or the name of a function as a string; the function must be a user function, not a built-in one.

The purpose of sandboxtrust is to give back some of the capabilities of unrestricted code to code executed in a sandbox. For instance.

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if unsecure code must be able to read the contents of a specific file, a trusted function should be written for that. It is very important for the trusted function to check carefully its arguments, such as file paths or URL.

### **Example**

Function which reads the contents of file 'data.txt':

```
function data = readFile
  fd = fopen('data.txt');
  data = fread(fd, inf, '*char');
  fclose(fd);
```

Execution of unsecure code which may read this file:

```
sandboxtrust(@readFile);
sandbox('d = readFile;');
```

#### See also

sandbox

# 6.16 Operators

Operators are special functions with a syntax which mimics mathematical arithmetic operations like the addition and the multiplication. They can be infix (such as x+y), separating their two arguments (called operands); prefix (such as -x), placed before their unique operand; or postfix (such as M'), placed after their unique operand. In Sysquake, their arguments are always evaluated from left to right. Since they do not require parenthesis or comma, their priority matters. Priority specifies when subexpressions are considered as a whole, as the argument of some operator. For instance, in the expression a+b\*c, where \* denotes the multiplication, the evaluation could result in (a+b)\*c or a+(b\*c); however, since operator \*'s priority is higher than operator +'s, the expression yields a+(b\*c) without ambiguity.

Here is the list of operators, from higher to lower priority:

Most operators have also a functional syntax; for instance, a+b can also be written plus(a,b). This enables their overriding with new definitions and their use in function references or functions such as feval which take the name of a function as an argument.

Here is the correspondence between operators and functions:

```
[a;b]
                               minus(a,b)
      vertcat(a,b)
                         a-b
[a,b]
      horzcat(a,b)
                               mtimes(a,b)
                         a∗b
a:b
       colon(a,b)
                         a/b
                               mrdivide(a,b)
a:b:c colon(a,b,c)
                         a\b
                               mldivide(a,b)
alb
       or(a.b)
                         a.*b
                               times(a,b)
                         a./b
a&b
       and(a,b)
                               rdivide(a,b)
a \le b le(a,b)
                         a.\b
                               ldivide(a,b)
a<b
      lt(a,b)
                         a^b
                               mpower(a,b)
a>=b
      qe(a,b)
                         a.^b
                               power(a,b)
                               not(a)
a>b
      qt(a,b)
                         ~a
                               uminus(a)
a==b
      eq(a,b)
                         - a
a~=b ne(a,b)
                               uplus(a)
                         +a
a===b same(a,b)
                         a'
                               ctranspose(a)
                         a.'
a~==b
      unsame(a,b)
                               transpose(a)
a+b
       plus(a,b)
```

Operator which do *not* have a corresponding function are ?:, && and || because unlike functions, they do not always evaluate all of their operands.

# Operator ()

Parenthesis.

### **Syntax**

```
(expr)
v(:)
v(index)
v(index1, index2)
v(:, index)
v(index, :)
v(select)
v(select1, select2)
v(:,:)
```

### **Description**

A pair of parenthesis can be used to change the order of evaluation. The subexpression it encloses is evaluated as a whole and used as if it was a single object. Parenthesis serve also to indicate a list of input or output parameters; see the description of the function keyword.

The last use of parenthesis is for specifying some elements of an array or list variable.

**Arrays:** In LME, any numeric object is considered as an array of two dimensions or more. Therefore, at least two indices are required to specify a single element; the first index specifies the row, the second the column, and so on. In some circumstances, however, it is convenient to consider an array as a vector, be it a column vector, a row vector, or even a matrix whose elements are indexed row-wise (or on some platforms). For this way of handling arrays, a single index is specified.

The first valid value of an index is always 1. The array whose elements are extracted is usually a variable, but can be any expression: an expression like [1,2;3,4](1,2) is valid and gives the 2nd element of the first row, i.e. 3.

In all indexing operations, several indices can be specified simultaneously to extract more than one element along a dimension. A single colon means all the elements along the corresponding dimension.

Instead of indices, the elements to be extracted can be selected by the true values in a logical array of the same size as the variable (the result is a column vector), or in a logical vector of the same size as the corresponding dimension. Calculating a boolean expression based on the variable itself used as a whole is the easiest way to get a logical array.

Variable indexing can be used in an expression or in the left hand side of an assignment. In this latter case, the right hand size can be one of the following:

- An array of the same size as the extracted elements.

- A scalar, which is assigned to each selected element of the variable.
- An empty matrix [], which means that the selected elements should be deleted. Only whole rows or columns (or (hyper)planes for arrays of more dimensions) can be deleted; i.e. a(2:5,:) = [] and b([3,6:8]) = [] (if b is a row or column vector) are legal, while c(2,3) = [] is not.

When indices are larger than the dimensions of the variable, the variable is expanded; new elements are set to 0 for numeric arrays, false for logical arrays, the nul character for character array, and the empty array [] for cell arrays.

**Lists:** In LME, lists have one dimension; thus a single index is required. Be it with a single index or a vector of indices, indexed elements are grouped in a list. New elements, also provided in a list, can be assigned to indexed elements; if the list to be assigned has a single element, the element is assigned to every indexed element of the variable.

**Cell arrays:** cell arrays are subscripted like other arrays. The result, or the right-hand side of an assignment, is also a cell array, or a list for the syntax v(select) (lists are to cell arrays what column vectors are to non-cell arrays). To create a single logical array for selecting some elements, function cellfun may be useful. To remove cells, the right-hand side of the assignment can be the empty list {} or the empty array [].

**Structure arrays:** access to structure array fields combines subscripting with parenthesis and structure field access with dot notation. When the field is not specified, parenthesis indexing returns a structure or structure array. When indexing results in multiple elements and a field is specified, the result is a value sequence.

### **Examples**

Ordering evaluation:

```
(1+2)*3
9
```

Extracting a single element, a row, and a column:

```
a = [1,2,3; 4,5,6];
a(2,3)
6
a(2,:)
4 5 6
a(:,3)
3
6
```

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Extracting a sub-array with contiguous rows and non-contiguous columns:

```
a(1:2,[1,3])
1 3
4 6
```

Array elements as a vector:

```
a(3:5)
3
4
5
a(:)
1
2
3
4
5
```

Selections of elements where a logical expression is true:

```
a(a>=5)
5
6
a(:, sum(a,1) > 6)
2 3
5 6
```

Assignment:

```
a(1,5) = 99
a =
1 2 3 0 99
4 5 6 0 0
```

Extraction and assignment of elements in a list:

```
a = {1,[2,7,3],'abc',magic(3),'x'};
a([2,5])
    {[2,7,3],'x'}
a([2,5]) = {'ab','cde'}
a =
    {1,'ab','abc',[8,1,6;3,5,7;4,9,2],'cde'}
a([2,5]) = {[3,9]}
a =
    {1,[3,9],'abc',[8,1,6;3,5,7;4,9,2],[3,9]}
```

Removing elements in a list ({} and [] have the same effect here):

```
a(4) = {}

a =

{1,[3,9],'abc',[3,9]}

a([1, 3]) = []

a =

{[3,9],[3,9]}
```

Replacing NaN with empty arrays in a cell array:

```
C = {'abc', nan; 2, false};
C(cellfun(@(x) any(isnan(x(:))), C)) = {[]};
```

Element in a structure array:

```
SA = structarray('a',{1,[2,3]},'b',{'ab','cde'});
SA(1).a
   2 3
SA(2).b = 'X';
```

When assigning a new field and/or a new element of a structure array, the new field is added to each element and the size of the array is expanded; fields are initialized to the empty array [].

```
SA(3).c = true;
SA(1).c
[]
```

#### See also

Operator {}, operator ., end, reshape, variable assignment, operator [], subsref, subsasgn, cellfun

# Operator []

Brackets.

## **Syntax**

```
[matrix_elements]
```

# **Description**

A pair of brackets is used to define a 2-d array given by its elements or by submatrices. The operator , (or spaces) is used to separate elements on the same row, and the operator ; (or newline) is used to separate rows. Since the space is considered as a separator when it is in the direct scope of brackets, it should not be used at the top level

of expressions; as long as this rule is observed, each element can be given by an expression.

Inside brackets, commas and semicolons are interpreted as calls to horzcat and vertcat. Brackets themselves have no other effect than changing the meaning of commas, semicolons, spaces, and new lines: the expression [1], for instance, is strictly equivalent to 1. The empty array [] is a special case.

Since horzcat and vertcat also accept cell arrays, brackets can be used to concatenate cell arrays, too.

## **Examples**

```
[1, 2, 3+5]
  1 2 8
[1:3; 25, 9]
  1 2 3
  2 5 9
[5-2, 3]
  3 3
[5 - 2, 3]
  5 -2 3
[(5 - 2), 3]
  3 3
[1 2
3 41
  1 2
  3 4
[]
  []
```

Concatenation of two cell arrays:

```
C1 = {1; 2};
C2 = {'ab'; false};
[C1, C2]
2x2 cell array
```

Compare this with the effect of braces, where elements are not concatenated but used as cells:

```
{C1, C2}
 1x2 cell array
```

### See also

```
Operator {}, operator (), operator ,, operator ;
```

# Operator {}

Braces.

### **Syntax**

```
{list_elements}
{cells}
{struct_elements}
v{index}
v{index1, index2, ...}
v{index} = expr
fun(...,v{:},...)
```

## **Description**

A pair of braces is used to define a list, a cell array, a struct, or an n-by-1 struct array given by its elements. When no element has a name (a named element is written name=value where value can be any expression), the result is a list or a celle array; when all elements have a name, the result is a struct or a struct array.

In a list, the operator , is used to separate elements. In a cell array, the operator , is used to separate cells on the same row; the operator ; is used to separate rows. Braces without semicolons produce a list; braces with semicolon(s) produce a cell array.

In a struct, the operator , is used to separate fields. In a struct array, the operator ; is used to separate elements.

 $v\{index\}$  is the element of list variable v whose index is given. index must be an integer between 1 (for the first element) and length(v) (for the last element).  $v\{index\}$  may be used in an expression to extract an element, or on the left hand-side of the equal sign to assign a new value to an element. Unless it is the target of an assignment, v may also be the result of an expression. If v is a cell array,  $v\{index\}$  is the element number index.

v{index1,index2,...} gives the specified cell of a cell array.

v itself may be an element or a field in a larger variable, provided it is a list; i.e. complicated assignments like  $a\{2\}.f\{3\}(2,5)=3$  are accepted. In an assignment, when the index (or indices) are larger than the list or cell array size, the variable is expanded with empty arrays [].

In the list of the input arguments of a function call,  $v\{:\}$  is replaced with its elements. v may be a list variable or the result of an expression.

### **Examples**

```
x = {1, 'abc', [3,5;7,1]}
x =
    {1,string,real 2x2}
x{3}
    3 5
```

```
7 1
x{2} = 2+3i
 x =
   {1,2+3j,real 2x2}
x{3} = {2}
 x =
    {1,2+3j,list}
x\{end+1\} = 123
    {1,2+3j,list,123}
C = \{1, false; 'ab', magic(3)\}
  2x2 cell array
C\{2, 1\}
  ab
a = \{1, 3:5\};
fprintf('%d', a{:}, 99);
  1 3 4 5 99
s = \{a=1, b='abc'\};
s.a
  1
S = \{a=1, b='abc'; a=false, b=1:5\};
size(S)
 2 1
S(2).b
 1 2 3 4 5
S = \{a=1; b=2\};
S(1).b
  []
```

#### See also

operator ,, operator [], operator (), operator ;, operator ., subsref, subsasgn

# Operator . (dot)

Structure field access.

# **Syntax**

```
v.field
v.field = expr
```

# Description

A dot is used to access a field in a structure. In v.field, v is the name of a variable which contains a structure, and field is the name of the field. In expressions, v.field gives the value of the field; it is an error

if it does not exist. As the target of an assignment, the value of the field is replaced if it exists, or a new field is added otherwise; if v itself is not defined, a structure is created from scratch.

v itself may be an element or a field in a larger variable, provided it is a structure (or does not exists in an assignment); i.e. complicated assignments like  $a\{2\}.f\{3\}(2,5)=3$  are accepted.

If V is a structure array, V. field is a value sequence which contains the specified field of each element of V.

The syntax v.(expr) permits to specify the field name dynamically at run-time, as the result of evaluating expression expr. v('f') is equivalent to v.f. This syntax is more elegant than functions getfield and setfield.

## **Examples**

```
s.f = 2
    s =
        f: 2
s.g = 'hello'
    s =
        f: 2
        s: string
s.f = 1:s.f
s =
        f: real 1x2
        g: string
```

#### See also

Operator (), operator {}, getfield setfield, subsref, subsasgn

# Operator +

Addition.

```
x + y
M1 + M2
M + x
plus(x, y)
+x
+M
uplus(x)
```

## **Description**

With two operands, both operands are added together. If both operands are matrices with a size different from 1-by-1, their size must be equal; the addition is performed element-wise. If one operand is a scalar, it is added to each element of the other operand.

With one operand, no operation is performed, except that the result is converted to a number if it was a string or a logical value, like with all mathematical operators and functions. For strings, each character is replaced with its numeric encoding. The prefix + is actually a synonym of double.

plus(x,y) is equivalent to x+y, and uplus(x) to +x. They can be used to redefine these operators for objects.

## **Example**

```
2 + 3

5

[1 2] + [3 5]

4 7

[3 4] + 2

5 6
```

#### See also

operator -, sum, addpol, double

# Operator -

Subtraction or negation.

## **Syntax**

```
x - y
M1 - M2
M - x
minus(x, y)
-x
-M
uminus(x)
```

# **Description**

With two operands, the second operand is subtracted from the first operand. If both operands are matrices with a size different from 1-by-1, their size must be equal; the subtraction is performed element-

wise. If one operand is a scalar, it is repeated to match the size of the other operand.

With one operand, the sign of each element is changed.

minus(x,y) is equivalent to x-y, and uminus(x) to -x. They can be used to redefine these operators for objects.

## **Example**

```
2 - 3

-1

[1 2] - [3 5]

-2 -3

[3 4] - 2

1 2

-[2 2-3j]

-2 -2+3j
```

#### See also

operator +, conj

## Operator \*

Matrix multiplication.

## **Syntax**

```
X * y
M1 * M2
M * x
mtimes(x, y)
```

# **Description**

x\*y multiplies the operands together. Operands can be scalars (plain arithmetic product), matrices (matrix product), or mixed scalar and matrix.

mtimes(x,y) is equivalent to x\*y. It can be used to redefine this operator for objects.

## **Example**

```
2 * 3
6
[1,2;3,4] * [3;5]
```

#### See also

operator .\*, operator /, prod

## Operator .\*

Scalar multiplication.

## **Syntax**

```
X .* y
M1 .* M2
M .* x
times(x, y)
```

## **Description**

x.\*y is the element-wise multiplication. If both operands are matrices with a size different from 1-by-1, their size must be equal; the multiplication is performed element-wise. If one operand is a scalar, it multiplies each element of the other operand.

times(x,y) is equivalent to x.\*y. It can be used to redefine this operator for objects.

## **Example**

```
[1 2] .* [3 5]
3 10
[3 4] .* 2
6 8
```

#### See also

```
operator *, operator ./, operator .^
```

# Operator /

Matrix right division.

## **Syntax**

```
a / b
A / B
A / b
mrdivide(a, b)
```

## **Description**

a/b divides the first operand by the second operand. If the second operand is a scalar, it divides each element of the first operand. Otherwise, it must be a square matrix; M1/M2 is equivalent to M1\*inv(M2). mrdivide(x,y) is equivalent to x/y. It can be used to redefine this operator for objects.

## **Example**

```
9 / 3

3

[2,6] / [1,2;3,4]

5 -1

[4 10] / 2

2 5
```

#### See also

operator \, inv, operator ./, deconv

# Operator ./

Scalar right division.

## **Syntax**

```
x ./ y
M1 ./ M2
M ./ x
x ./ M
rdivide(x, y)
```

# **Description**

The first operand is divided by the second operand. If both operands are matrices with a size different from 1-by-1, their size must be equal; the division is performed element-wise. If one operand is a scalar, it is repeated to match the size of the other operand.

rdivide(x,y) is equivalent to x./y. It can be used to redefine this operator for objects.

### **Examples**

```
[3 10] ./ [3 5]
1 2
[4 8] ./ 2
2 4
10 ./ [5 2]
2 5
```

#### See also

operator /, operator .\*, operator .^

# **Operator** \

Matrix left division.

## **Syntax**

```
x \ y
M1 \ M2
x \ M
mldivide(x, y)
```

## **Description**

x\y divides the second operand by the first operand. If the first operand is a scalar, it divides each element of the second operand. Otherwise, it must be a square matrix;  $M1\M2$  is equivalent to inv(M1)\*M2.

mldivide(x,y) is equivalent to  $x\y$ . It can be used to redefine this operator for objects.

## **Examples**

```
3 \ 9

3

[1,2;3,4] \ [2;6]

2

0

2 \ [4 10]

2 5
```

### See also

```
operator /, inv, operator .\
```

# Operator .\

Scalar left division.

## **Syntax**

```
M1 .\ M2
M1 .\ x
ldivide(x, y)
```

## **Description**

The second operand is divided by the first operand. If both operands are matrices with a size different from 1-by-1, their size must be equal; the division is performed element-wise. If one operand is a scalar, it is repeated to match the size of the other operand.

ldivide(x,y) is equivalent to x.\y. It can be used to redefine this operator for objects.

## **Example**

```
[1 2 3] .\ [10 11 12]
10 5.5 4
```

#### See also

operator \, operator ./

# Operator ^

Matrix power.

```
x ^ y
M ^ y
x ^ M
mpower(x, y)
```

## **Description**

 $x^y$  calculates x to the y power, provided that either

- both operands are scalar;
- the first operand is a square matrix and the second operand is a scalar:
- or the first operand is a scalar and the second operand is a square matrix.

Other cases yield an error.

mpower(x,y) is equivalent to  $x^y$ . It can be used to redefine this operator for objects.

## **Examples**

```
2 ^ 3

8

[1,2;3,4] ^ 2

7 10

15 22

2 ^ [1,2;3,4]

10.4827 14.1519

21.2278 31.7106
```

## **Algorithms**

If the first operand is a scalar and the second a square matrix, the matrix exponential is used. The result is expm(log(x)\*M).

If the first operand is a square matrix and the second a scalar, unless for small real integers, the same algorithm as for matrix functions is used, i.e. a complex Schur decomposition followed by the Parlett method. The result is  $funm(M, @(x) x^{\circ}y)$ .

#### See also

```
operator .^, expm, funm
```

# Operator .^

Scalar power.

## **Syntax**

```
M1 .^ M2
x .^ M
M .^ x
power(x, y)
```

## **Description**

M1. M2 calculates M1 to the M2 power, element-wise. Both arguments must have the same size, unless one of them is a scalar.

power(x,y) is equivalent to x. $\hat{y}$ . It can be used to redefine this operator for objects.

## **Examples**

```
[1,2;3,4].<sup>2</sup>
1 4
9 16
[1,2,3].<sup>2</sup>[5,4,3]
1 16 27
```

#### See also

```
operator ^, exp
```

# Operator '

Complex conjugate transpose.

## **Syntax**

```
M'
ctranspose(M)
```

# **Description**

M' is the transpose of the real matrix M, i.e. columns and rows are permuted. If M is complex, the result is the complex conjugate transpose of M. If M is an array with multiple dimensions, the first two dimensions are permuted.

ctranspose(M) is equivalent to M'. It can be used to redefine this operator for objects.

## **Examples**

```
[1,2;3,4]'
1 3
2 4
[1+2j, 3-4j]'
1-2j
3+4j
```

#### See also

```
operator .', conj
```

# Operator .'

Transpose.

## **Syntax**

```
M.'
transpose(M)
```

## **Description**

M. ' is the transpose of the matrix M, i.e. columns and rows are permuted. M can be real or complex. If M is an array with multiple dimensions, the first two dimensions are permuted.

transpose(M) is equivalent to M.'. It can be used to redefine this operator for objects.

## **Example**

```
[1,2;3,4].'
1 3
2 4
[1+2j, 3-4j].'
1+2j
3-4j
```

#### See also

```
operator', permute, fliplr, flipud, rot90
```

# Operator ==

Equality.

## **Syntax**

```
x == y
eq(x, y)
```

## Description

x == y is true if x is equal to y, and false otherwise. Comparing NaN (not a number) to any number yields false, including to NaN. If x and/or y is an array, the comparison is performed element-wise and the result has the same size.

eq(x,y) is equivalent to x==y. It can be used to redefine this operator for objects.

## **Example**

```
1 == 1
    true
1 == 1 + eps
    false
1 == 1 + eps / 2
    true
inf == inf
    true
nan == nan
    false
[1,2,3] == [1,3,3]
    T F T
```

#### See also

operator ~=, operator <=, operator >=, operator >=, operator >=, operator ===, operator ~==, strcmp

# Operator ===

Object equality.

```
a === b
same(a, b)
```

## **Description**

a === b is true if a is the same as b, and false otherwise. a and b must have exactly the same internal representation to be considered as equal; with IEEE floating-point numbers, nan===nan is true and  $\theta$ ===-0 is false. Contrary to the equality operator ==, === returns a single boolean even if its operands are arrays.

same(a,b) is equivalent to a===b.

## **Example**

```
(1:5) === (1:5)
    true
(1:5) == (1:5)
    T T T T
[1,2,3] === [4,5]
    false
[1,2,3] == [4,5]
    Incompatible size
nan === nan
    true
nan == nan
    false
```

#### See also

operator ~==, operator ==, operator ~=, operator <=, operator >=, operator >=, operator ~=, strcmp

# Operator ~=

Inequality.

## **Syntax**

```
x ~= y
ne(x, y)
```

# **Description**

x = y is true if x is not equal to y, and false otherwise. Comparing NaN (not a number) to any number yields true, including to NaN. If x and/or y is an array, the comparison is performed element-wise and the result has the same size.

ne(x,y) is equivalent to  $\tilde{x}=y$ . It can be used to redefine this operator for objects.

### **Example**

```
1 ~= 1
  false
inf ~= inf
  false
nan ~= nan
  true
[1,2,3] ~= [1,3,3]
  F T F
```

#### See also

operator ==, operator <, operator <=, operator >=, operator >=, operator ===, operator ===, strcmp

# Operator ~==

Object inequality.

## **Syntax**

```
a ~== b
unsame(a, b)
```

## Description

a  $\tilde{}==$  b is true if a is not the same as b, and false otherwise. a and b must have exactly the same internal representation to be considered as equal; with IEEE numbers, nan $\tilde{}==$ nan is false and 0 $\tilde{}==-0$  is true. Contrary to the inequality operator,  $\tilde{}==$  returns a single boolean even if its operands are arrays.

unsame(a,b) is equivalent to  $\tilde{a} == b$ .

## **Example**

```
(1:5) ~== (1:5)
  false
(1:5) ~= (1:5)
  F F F F F
[1,2,3] ~== [4,5]
  true
[1,2,3] ~= [4,5]
  Incompatible size
nan ~== nan
  false
nan ~= nan
  true
```

#### See also

operator ===, operator ==, operator <, operator <=, operator >=, strcmp

# Operator <

Less than.

## **Syntax**

## **Description**

x < y is true if x is less than y, and false otherwise. Comparing NaN (not a number) to any number yields false, including to NaN. If x and/or y is an array, the comparison is performed element-wise and the result has the same size.

lt(x,y) is equivalent to x < y. It can be used to redefine this operator for objects.

## Example

#### See also

operator ==, operator ~=, operator >=, operator >=

# Operator >

Greater than.

## **Description**

x > y is true if x is greater than y, and false otherwise. Comparing NaN (not a number) to any number yields false, including to NaN. If x and/or y is an array, the comparison is performed element-wise and the result has the same size.

gt(x,y) is equivalent to x>y. It can be used to redefine this operator for objects.

### **Example**

$$[2,3,4] > [2,4,2]$$
  
F F T

#### See also

operator ==, operator ~=, operator <=, operator >=

## Operator <=

Less or equal to.

#### **Syntax**

$$x \le y$$
 le(x, y)

## **Description**

 $x \le y$  is true if x is less than or equal to y, and false otherwise. Comparing NaN (not a number) to any number yields false, including to NaN. If x and/or y is an array, the comparison is performed elementwise and the result has the same size.

le(x,y) is equivalent to  $x \le y$ . It can be used to redefine this operator for objects.

## **Example**

$$[2,3,4] <= [2,4,2]$$
T T F

#### See also

operator ==, operator ~=, operator >, operator >=

# Operator >=

Greater or equal to.

## **Syntax**

```
x >= y qe(x, y)
```

## **Description**

x >= y is true if x is greater than or equal to y, and false otherwise. Comparing NaN (not a number) to any number yields false, including to NaN. If x and/or y is an array, the comparison is performed elementwise and the result has the same size.

ge(x,y) is equivalent to x>=y. It can be used to redefine this operator for objects.

## **Example**

$$[2,3,4] >= [2,4,2]$$
  
T F T

#### See also

operator ==, operator ~=, operator <=, operator >=

# Operator ~

Not.

## **Syntax**

```
~b
not(b)
```

# **Description**

b is false (logical 0) if b is different from 0 or false, and true otherwise. If b is an array, the operation is performed on each element.

not(b) is equivalent to ~b. It can be used to redefine this operator for objects.

Character ~ can also be used as a placeholder for unused arguments.

### **Examples**

```
~true
  false
~[1,0,3,false]
  F T F T
```

#### See also

operator ~=, bitcmp, function (unused arguments)

# Operator &

And.

## **Syntax**

```
b1 & b2
and(b1, b2)
```

## **Description**

b1&b2 performs the logical AND operation between the corresponding elements of b1 and b2; the result is true (logical 1) if both operands are different from false or 0, and false (logical 0) otherwise.

and (b1, b2) is equivalent to b1&b2. It can be used to redefine this operator for objects.

## **Example**

```
[false, false, true, true] & [false, true, false, true]
    F F T
```

#### See also

```
operator |, xor, operator ~, operator &&, all
```

# Operator &&

And with lazy evaluation.

## Syntax

b1 && b2

## **Description**

b1&&b2 is b1 if b1 is false, and b2 otherwise. Like with if and while statements, b1 is true if it is a nonempty array with only non-zero elements. b2 is evaluated only if b1 is true.

b1&&b2&&...&&bn returns the last operand which is false (remaining operands are not evaluated), or the last one.

### **Example**

Boolean value which is true if the vector v is made of pairs of equal values:

```
mod(length(v),2) == 0 \&\& v(1:2:end) == v(2:2:end)
```

The second operand of && is evaluated only if the length is even.

#### See also

operator ||, operator ?, operator &, if

## Operator |

Or.

### **Syntax**

```
b1 | b2
or(b1, b2)
```

## **Description**

b1|b2 performs the logical OR operation between the corresponding elements of b1 and b2; the result is false (logical 0) if both operands are false or 0, and true (logical 1) otherwise.

or (b1,b2) is equivalent to b1|b2. It can be used to redefine this operator for objects.

## **Example**

```
[false, false, true, true] | [false, true, false, true]
   F T T T
```

#### See also

operator &, xor, operator ~, operator ||, any

# Operator ||

Or with lazy evaluation.

### **Syntax**

b1 || b2

## **Description**

b1||b2 is b1 if b1 is true, and b2 otherwise. Like with if and while statements, b1 is true if it is a nonempty array with only non-zero elements. b2 is evaluated only if b1 is false.

b1||b2||...||bn returns the last operand which is true (remaining operands are not evaluated), or the last one.

## Example

Boolean value which is true if the vector v is empty or if its first element is NaN:

```
isempty(v) || isnan(v(1))
```

#### See also

operator &&, operator ?, operator |, if

# **Operator**?

Alternative with lazy evaluation.

#### **Syntax**

b ? x : y

# Description

b?x:y is x if b is true, and y otherwise. Like with if and while statements, b is true if it is a nonempty array with only non-zero elements. Only one of x and y is evaluated depending on b.

Operators ? and : have the same priority; parenthesis or brackets should be used if e.g. x or y is a range.

## **Example**

Element of a vector v, or default value 5 if the index ind is out of range:

```
ind < 1 \mid ind > length(v) ? 5 : v(ind)
```

#### See also

operator &&, operator ||, if

## Operator,

Horizontal matrix concatenation.

## **Syntax**

```
[M1, M2, ...]
[M1 M2 ...]
horzcat(M1, M2, ...)
```

## **Description**

Between brackets, the comma is used to separate elements on the same row in a matrix. Elements can be scalars, vector, arrays, cell arrays, or structures; their number of rows must be the same, unless one of them is an empty array. For arrays with more than 2 dimensions, all dimensions except dimension 2 (number of columns) must match.

Outside brackets or between parenthesis, the comma is used to separate statements or the arguments of functions.

horzcat(M1,M2,...) is equivalent to [M1,M2,...]. It can be used to redefine this operator for objects. It accepts any number of input arguments; horzcat() is the real double empty array [], and horzcat(M) is M.

Between braces, the comma separates cells on the same row.

## **Examples**

```
[1,2]

1 2

[[3;5],ones(2)]

3 1 1

5 1 1

['abc','def']

abcdef
```

### See also

```
operator [], operator ;, cat, join, operator {}
```

## Operator;

Vertical matrix concatenation.

### **Syntax**

```
[M1; M2]
vertcat(M1, M2)
```

## **Description**

Between brackets, the semicolon is used to separate rows in a matrix. Rows can be scalars, vector, arrays, cell arrays, or structures; their number of columns must be the same, unless one of them is an empty array. For arrays with more than 2 dimensions, all dimensions except dimension 1 (number of rows) must match.

Outside brackets, the comma is used to separate statements; they loose any meaning between parenthesis and give a syntax error.

vertcat(M1,M2) is equivalent to [M1;M2]. It can be used to redefine this operator for objects.

Between braces, the semicolon separates rows of cells.

## **Examples**

```
[1;2]
1
2
[1:5;3,2,4,5,1]
1 2 3 4 5
3 2 4 5 1
['abc';'def']
abc
def
```

### See also

```
operator [], operator ,, join, operator {}
```

# Operator:

Range.

## **Syntax**

```
x1:x2
x1:step:x2
colon(x1,x2)
colon(x1,step,x2)
```

## **Description**

x1:x2 gives a row vector with the elements x1, x1+1, x1+2, etc. until x2. The last element is equal to x2 only if x2-x1 is an integer, and smaller otherwise. If x2 < x1, the result is an empty matrix.

x1:step:x2 gives a row vector with the elements x1, x1+step, x1+2\*step, etc. until x2. The last element is equal to x2 only if (x2-x1)/step is an integer. With fractional numbers, rounding errors may cause x2 to be discarded even if (x2-x1)/step is "almost" an integer. If x2\*sign(step)<x1\*sign(step), the result is an empty matrix.

If x1 or step is complex, a complex vector is produced, with the expected contents. The following algorithm is used to generate each element:

```
z = x1
while real((x2 - z) * conj(step)) >= 0
    append z to the result
    z = z + step
end
```

Values are added until they go beyond the projection of x2 onto the straight line defined by x1 and direction step. If x2-x1 and step are orthogonal, it is attempted to produce an infinite number of elements, which will obviously trigger an out of memory error. This is similar to having a null step in the real case.

Note that the default step value is always 1 for consistency with real values. Choosing for instance sign(x2-x1) would have made the generation of lists of indices more difficult. Hence for a vector of purely imaginary numbers, always specify a step.

colon(x1,x2) is equivalent to x1:x2, and colon(x1,step,x2) to x1:step:x2. It can be used to redefine this operator for objects.

The colon character is also used to separate the alternatives of a conditional expression b?x:y.

### **Example**

```
2:5
2 3 4 5
2:5.3
2 3 4 5
```

```
3:3
3
3:2
[]
2:2:8
2 4 6 8
5:-1:2
5 4 3 2
0:1j:10j
0 1j 2j 3j 4j 5j 6j 7j 8j 9j 10j
1:1+1j:5+4j
1 2+1j 3+2j 4+3j 5+4j
0:1+1j:5
0 1+1j 2+2j 3+3j 4+4j 5+5j
```

#### See also

repmat, operator?

# Operator @

Function reference or anonymous function.

## Syntax

```
@fun
@(arguments) expression
```

## **Description**

@fun gives a reference to function fun which can be used wherever an inline function can. Its main use is as the argument of functions like feval or integral, but it may also be stored in lists, cell arrays, or structures. A reference cannot be cast to a number (unlike characters or logical values), nor can it be stored in a numeric array. The function reference of an operator must use its function name, such as @plus.

Anonymous functions are an alternative, more compact syntax for inline functions. In @(args) expr, args is a list of input arguments and expr is an expression which contains two kinds of variables:

- input arguments, provided when the anonymous expression is executed:
- captured variables (all variables which do not appear in the list of input arguments), which have the value of variables of the same name existing when and where the anonymous function is created. These values are fixed.

If the top-level element of the anonymous function is itself a function, multiple output arguments can be specified for the call of the anonymous function, as if a direct call was performed. Anonymous functions which do not return any output are also valid.

Anonymous functions may not have input arguments with default values (@(x=2)x+5 is invalid).

Anonymous functions are a convenient way to provide the glue between functions like fzero and ode45 and the function they accept as argument. Additional parameters can be passed directly in the anonymous function with captured variables, instead of being supplied as additional arguments; the code becomes clearer.

### **Examples**

Function reference:

```
integral(@sin, 0, pi)
2
```

Anonymous function:

```
a = 2;
fun = @(x) sin(a * x);
fun(3)
  -0.2794
integral(fun, 0, 2)
  0.8268
```

Without anonymous function, parameter a should be passed as an additional argument after all the input arguments defined for integral, including those which are optional when parameters are missing:

```
integral(inline('sin(a * x)', 'x', 'a'), 0, 2, [], false, a) 0.8268
```

Anonymous functions are actually stored as inline functions with all captured variables:

```
dumpvar(fun)
  inline('function y=f(a,x);y=sin(a*x);',2)
```

Anonymous function with multiple output arguments:

```
fun = @(A) size(A);
s = fun(ones(2,3))
s =
    2 3
(m, n) = fun(ones(2,3))
m =
    2
n =
    3
```

#### See also

fun2str, str2fun, inline, feval, apply

# 6.17 Mathematical Functions

### abs

Absolute value.

## **Syntax**

```
x = abs(z)
```

## **Description**

abs takes the absolute value of each element of its argument. The result is an array of the same size as the argument; each element is non-negative.

## **Example**

```
abs([2,-3,0,3+4j]
2 3 0 5
```

#### See also

angle, sign, real, imag, hypot

#### acos

Arc cosine.

#### **Syntax**

$$y = acos(x)$$

# **Description**

acos(x) gives the arc cosine of x, which is complex if x is complex or if abs(x)>1.

## **Examples**

```
acos(2)
0+1.3170j
acos([0,1+2j])
1.5708 1.1437-1.5286j
```

#### See also

cos, asin, acosh

# acosd acotd acscd asecd asind atand atan2d

Inverse trigonometric functions with angles in degrees.

## **Syntax**

```
y = acosd(x)
y = acotd(x)
y = acscd(x)
y = asecd(x)
y = asind(x)
y = atand(x)
z = atan2d(y, x)
```

# **Description**

Inverse trigonometric functions whose name ends with a d give a result expressed in degrees instead of radians.

# **Examples**

```
acosd(0.5)
60.0000
acos(0.5) * 180 / pi
60.0000
```

#### See also

cosd, cotd, cscd, secd, sind, tand, acos, acot, acsc, asec, asin,
atan, atan2

### acosh

Inverse hyperbolic cosine.

## **Syntax**

```
y = acosh(x)
```

## **Description**

acosh(x) gives the inverse hyperbolic cosine of x, which is complex if x is complex or if x<1.

## **Examples**

```
acosh(2)
1.3170
acosh([0,1+2j])
0+1.5708j 1.5286+1.1437j
```

#### See also

cosh, asinh, acos

### acot

Inverse cotangent.

## **Syntax**

```
y = acot(x)
```

# **Description**

acot(x) gives the inverse cotangent of x, which is complex if x is.

#### See also

```
cot, acoth, cos
```

#### acoth

Inverse hyperbolic cotangent.

```
y = acoth(x)
```

## **Description**

acoth(x) gives the inverse hyperbolic cotangent of x, which is complex if x is complex or is in the range (-1,1).

#### See also

coth, acot, atanh

#### acsc

Inverse cosecant.

## **Syntax**

```
y = acsc(x)
```

## **Description**

acsc(x) gives the inverse cosecant of x, which is complex if x is complex or is in the range (-1,1).

#### See also

csc, acsch, asin

### acsch

Inverse hyperbolic cosecant.

## **Syntax**

```
y = acsch(x)
```

## **Description**

 $\operatorname{acsch}(x)$  gives the inverse hyperbolic cosecant of x, which is complex if x is.

#### See also

csc, acsc, asinh

# angle

Phase angle of a complex number.

## **Syntax**

```
phi = angle(z)
```

## **Description**

angle(z) gives the phase of the complex number z, i.e. the angle between the positive real axis and a line joining the origin to z. angle(0) is 0.

## **Examples**

```
angle(1+3j)
1.2490
angle([0,1,-1])
0 0 3.1416
```

#### See also

abs, sign, atan2

#### asec

Inverse secant.

## **Syntax**

```
y = asec(x)
```

# **Description**

asec(x) gives the inverse secant of x, which is complex if x is complex or is in the range (-1,1).

#### See also

sec, asech, acos

## asech

Inverse hyperbolic secant.

## **Syntax**

```
y = asech(x)
```

## **Description**

asech(x) gives the inverse hyperbolic secant of x, which is complex if x is complex or strictly negative.

#### See also

```
sech, asec, acosh
```

### asin

Arc sine.

## **Syntax**

```
y = asin(x)
```

## **Description**

asin(x) gives the arc sine of x, which is complex if x is complex or if abs(x)>1.

## **Examples**

```
asin(0.5)
0.5236
asin(2)
1.5708-1.317j
```

#### See also

sin, acos, asinh

## asinh

Inverse hyperbolic sine.

```
y = asinh(x)
```

## **Description**

asinh(x) gives the inverse hyperbolic sine of x, which is complex if x is complex.

## **Examples**

```
asinh(2)
1.4436
asinh([0,1+2j])
0 1.8055+1.7359j
```

#### See also

sinh, acosh, asin

#### atan

Arc tangent.

## **Syntax**

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

## **Description**

atan(x) gives the arc tangent of x, which is complex if x is complex.

# **Example**

```
atan(1)
0.7854
```

#### See also

tan, asin, acos, atan2, atanh

#### atan2

Direction of a point given by its Cartesian coordinates.

```
phi = atan2(y,x)
```

## **Description**

atan2(y,x) gives the direction of a point given by its Cartesian coordinates x and y. Imaginary component of complex numbers is ignored. atan2(y,x) is equivalent to atan(y/x) if x>0.

## **Examples**

```
atan2(1, 1)
0.7854
atan2(-1, -1)
-2.3562
atan2(0, 0)
```

#### See also

atan, angle

## atanh

Inverse hyperbolic tangent.

# **Syntax**

```
y = atanh(x)
```

# **Description**

atan(x) gives the inverse hyperbolic tangent of x, which is complex if x is complex or if abs(x)>1.

# **Examples**

```
atanh(0.5)
0.5493
atanh(2)
0.5493 + 1.5708j
```

#### See also

asinh, acosh, atan

## beta

Beta function.

## **Syntax**

$$y = beta(z,w)$$

## **Description**

beta(z,w) gives the beta function of z and w. Arguments and result are real (imaginary part is discarded). The beta function is defined as

$$B(z, w) = \int_0^1 t^{z-1} (1-t)^{w-1} dt$$

### **Example**

### See also

gamma, betaln, betainc

### betainc

Incomplete beta function.

### **Syntax**

$$y = betainc(x,z,w)$$

# **Description**

betainc(x,z,w) gives the incomplete beta function of x, z and w. Arguments and result are real (imaginary part is discarded). x must be between 0 and 1. The incomplete beta function is defined as

$$I_X(z, w) = \frac{1}{B(z, w)} \int_0^x t^{z-1} (1-t)^{w-1} dt$$

## **Example**

#### See also

beta, betaln, gammainc

### betaln

Logarithm of beta function.

## **Syntax**

```
y = betaln(z, w)
```

## **Description**

betaln(z,w) gives the logarithm of the beta function of z and w. Arguments and result are real (imaginary part is discarded).

## **Example**

```
betaln(0.5,2)
0.2877
```

#### See also

beta, betainc, gammaln

# cart2pol

Cartesian to polar coordinates transform.

# Syntax

```
(phi, r) = cart2pol(x, y)

(phi, r, z) = cart2pol(x, y, z)
```

# **Description**

(phi,r)=cart2pol(x,y) transforms Cartesian coordinates x and y to polar coordinates phi and r such that  $x = r\cos(\varphi)$  and  $x = r\sin(\varphi)$ . (phi,r,z)=cart2pol(x,y,z) transform Cartesian coordinates to cylindrical coordinates, leaving z unchanged.

# **Example**

```
(phi, r) = cart2pol(1, 2)
  phi =
    1.1071
  r =
    2.2361
```

#### See also

cart2sph, pol2cart, sph2cart, abs, angle

## cart2sph

Cartesian to spherical coordinates transform.

## **Syntax**

```
(phi, theta, r) = cart2sph(x, y, z)
```

## **Description**

(phi,theta,r)=cart2sph(x,y,z) transforms Cartesian coordinates x, y, and z to polar coordinates phi, theta, and r such that  $x = r\cos(\varphi)\cos(\vartheta)$ ,  $y = r\sin(\varphi)\cos(\vartheta)$ , and  $z = r\sin(\vartheta)$ .

# Example

```
(phi, theta, r) = cart2sph(1, 2, 3)
phi =
    1.1071
theta =
    0.9303
r =
    3.7417
```

#### See also

cart2pol, pol2cart, sph2cart

#### cast

Type conversion.

```
Y = cast(X, type)
```

### **Description**

cast (X, type) converts the numeric array X to the type given by string type, which can be 'double', 'single', 'int8' or any other signed or unsigned integer type, 'char', or 'logical'. The number value is preserved when possible; conversion to integer types discards most significant bytes. If X is an array, conversion is performed on each element; the result has the same size. The imaginary part, if any, is discarded only with conversions to integer types.

## **Example**

```
cast(pi, 'int8')
  3int8
```

### See also

uint8 and related functions, double, single, typecast

#### cdf

Cumulative distribution function.

# Syntax

```
y = cdf(distribution,x)
y = cdf(distribution,x,a1)
y = cdf(distribution,x,a1,a2)
```

# **Description**

cdf(distribution,x) calculates the integral of a probability density function from  $-\infty$  to x. The distribution is specified with the first argument, a string; case is ignored ('t' and 'T' are equivalent). Additional arguments must be provided for some distributions. The distributions are given in the table below. Default values for the parameters, when mentioned, mean that the parameter may be omitted.

Distribution	Name	<b>Parameters</b>
beta	beta	a and b
Cauchy	cauchy	a and b
χ	chi	deg. of freedom $ u$
$\chi^2$	chi2	deg. of freedom $ u$
γ	gamma	shape $lpha$ and $\lambda$
exponential	exp	mean
F	f	deg. of freedom $v_1$ and $v_2$
half-normal	half-normal	9
Laplace	laplace	mean and scale
lognormal	logn	mean (0) and st. dev. (1)
Nakagami	nakagami	$\mu$ and $\omega$
normal	norm	mean (0) and st. dev. (1)
Rayleigh	rayl	b
Student's T	t	deg. of freedom $ u$
uniform	unif	limits of the range (0 and 1)
Weibull	weib	shape k and scale $\lambda$

```
cdf('chi2', 2.5, 3)
  0.5247
integral(@(x) pdf('chi2',x,3), 0, 2.5, AbsTol=1e-4)
  0.5247
```

## See also

pdf, icdf, random, erf

### ceil

Rounding towards +infinity.

## **Syntax**

```
y = ceil(x)
```

# **Description**

 $\mathtt{ceil}(x)$  gives the smallest integer larger than or equal to x. If the argument is a complex number, the real and imaginary parts are handled separately.

```
ceil(2.3)
   3
ceil(-2.3)
   -2
ceil(2.3-4.5j)
   3-4j
```

#### See also

floor, fix, round, roundn

# complex

Make a complex number.

### **Syntax**

```
z = complex(x, y)
```

# **Description**

complex (x,y) makes a complex number from its real part x and imaginary part y. The imaginary part of its input arguments is ignored.

## **Examples**

```
complex(2, 3)
2 + 3j
complex(1:5, 2)
1+2j 2+2j 3+2j 4+2j 5+2j
```

#### See also

```
real, imag, i
```

# conj

Complex conjugate value.

# **Syntax**

```
w = conj(z)
```

## **Description**

conj(z) changes the sign of the imaginary part of the complex number z.

### **Example**

```
conj([1+2j,-3-5j,4,0])
1-2j -3+5j 4 0
```

### See also

```
imag, angle, j, operator -
```

#### cos

Cosine.

### **Syntax**

$$y = cos(x)$$

# Description

cos(x) gives the cosine of x, which is complex if x is complex.

# Example

```
cos([0, 1+2j])
1 2.0327-3.0519j
```

## See also

sin, acos, cosh

# cosd cotd cscd secd sind tand

Trigonometric functions with angles in degrees.

# **Syntax**

```
y = cosd(x)
y = cotd(x)
y = cscd(x)
y = secd(x)
y = sind(x)
y = tand(x)
```

## **Description**

Trigonometric functions whose name ends with a d have an argument expressed in degrees instead of radians.

## **Examples**

```
cosd(20)
  0.9397
cos(20 * pi / 180)
  0.9397
```

### See also

acosd, acotd, acscd, asecd, asind, atand, atan2d, cos, cot, csc, sec, sin, tan

### cosh

Hyperbolic cosine.

## **Syntax**

```
y = \cosh(x)
```

# **Description**

cos(x) gives the hyperbolic cosine of x, which is complex if x is complex.

### **Example**

```
cosh([0, 1+2j])
1 -0.6421+1.0686j
```

sinh, acosh, cos

### cot

Cotangent.

# **Syntax**

$$y = cot(x)$$

# **Description**

cot(x) gives the cotangent of x, which is complex if x is.

## See also

acot, coth, tan

## coth

Hyperbolic cotangent.

### **Syntax**

$$y = coth(x)$$

# **Description**

coth(x) gives the hyperbolic cotangent of x, which is complex if x is.

### See also

acoth, cot, tanh

#### **CSC**

Cosecant.

# **Syntax**

$$y = csc(x)$$

## **Description**

csc(x) gives the cosecant of x, which is complex if x is.

### See also

acsc, csch, sin

### csch

Hyperbolic cosecant.

### **Syntax**

y = csch(x)

## **Description**

csch(x) gives the hyperbolic cosecant of x, which is complex if x is.

#### See also

acsch, csc, sinh

### diln

Dilogarithm.

### **Syntax**

$$y = diln(x)$$

# **Description**

diln(x) gives the dilogarithm, or Spence's integral, of x. Argument and result are real (imaginary part is discarded). The dilogarithm is defined as

$$diln(x) = \int_{1}^{x} \frac{\log(t)}{t - 1} dt$$

# **Example**

### double

Conversion to double-precision numbers.

## **Syntax**

```
B = double(A)
```

## **Description**

double(A) converts number or array A to double precision. A can be any kind of numeric value (real, complex, or integer), or a character or logical array.

To keep the integer type of logical and character arrays, the unitary operator + should be used instead.

### **Examples**

```
double(uint8(3))
  3
double('AB')
  65 66
islogical(double(1>2))
  false
```

#### See also

uint8 and related functions, single, cast, operator +, setstr, char, logical

# ellipam

Jacobi elliptic amplitude.

# **Syntax**

```
phi = ellipam(u, m)
phi = ellipam(u, m, tol)
```

# Description

ellipam(u,m) gives the Jacobi elliptic amplitude phi. Parameter m must be in [0,1]. The Jacobi elliptic amplitude is the inverse of the Jacobi integral of the first kind, such that  $u = F(\varphi|m)$ .

ellipam(u,m,tol) uses tolerance tol; the default tolerance is eps.

```
phi = ellipam(2.7, 0.6)
  phi =
     2.0713
u = ellipf(phi, 0.6)
u =
     2.7
```

### See also

ellipf, ellipj

# ellipe

Jacobi elliptic integral of the second kind.

## **Syntax**

```
u = ellipe(phi, m)
```

# **Description**

ellipe(phi,m) gives the Jacobi elliptic integral of the second kind, defined as

$$E(\varphi|m) = \int_0^{\varphi} \sqrt{1 - m \sin^2 t} \, \mathrm{d}t$$

Complete elliptic integrals of first and second kinds, with phi=pi/2, can be obtained with ellipke.

#### See also

ellipf, ellipke

# ellipf

Jacobi elliptic integral of the first kind.

# **Syntax**

```
u = ellipf(phi, m)
```

## **Description**

ellipf(phi,m) gives the Jacobi elliptic integral of the first kind, defined as

$$F(\varphi|m) = \int_0^{\varphi} \frac{\mathrm{d}t}{\sqrt{1 - m\sin^2 t}}$$

Complete elliptic integrals of first and second kinds, with phi=pi/2, can be obtained with ellipke.

#### See also

ellipe, ellipke, ellipam

# ellipj

Jacobi elliptic functions.

### **Syntax**

# **Description**

ellipj(u,m) gives the Jacobi elliptic function sn, cn, and dn. Parameter m must be in [0,1]. These functions are based on the Jacobi elliptic amplitude  $\varphi$ , the inverse of the Jacobi elliptic integral of the first kind which can be obtained with ellipam):

$$u=F(\varphi|m)$$

$$\operatorname{sn}(u|m) = \sin(\varphi)$$

$$cn(u|m) = cos(\varphi)$$

$$dn(u|m) = \sqrt{1 - m \sin^2 \varphi}$$

ellipj(u,m,tol) uses tolerance tol; the default tolerance is eps.

```
(sn, cn, dn) = ellipj(2.7, 0.6)
  sn =
    0.8773
  cn =
    -0.4799
  dn =
    0.7336
  sin(ellipam(2.7, 0.6))
    0.8773
ellipj(0:5, 0.3)
    0.0000    0.8188    0.9713    0.4114   -0.5341   -0.9930
```

### See also

ellipam, ellipke

# ellipke

Complete elliptic integral.

## **Syntax**

```
(K, E) = ellipke(m)
(K, E) = ellipke(m, tol)
```

# **Description**

(K,E)=ellipke(m) gives the complete elliptic integrals of the first kind K=F(m) and second kind E=E(m), defined as

$$F(m) = \int_0^{\pi/2} \frac{\mathrm{d}t}{\sqrt{1 - m\sin^2 t}}$$

$$E(m) = \int_0^{\pi/2} \sqrt{1 - m \sin^2 t} \, dt$$

Parameter m must be in [0,1]. ellipke(m,tol) uses tolerance tol; the default tolerance is eps.

# **Example**

```
ellipj
```

## eps

Difference between 1 and the smallest number x such that x > 1.

### **Syntax**

```
e = eps
e = eps(x)
e = eps(type)
```

### **Description**

Because of the floating-point encoding of "real" numbers, the absolute precision depends on the magnitude of the numbers. The relative precision is characterized by the number given by eps, which is the smallest double positive number such that 1+eps can be distinguished from 1.

eps(x) gives the smallest number e such that x+e has the same sign as x and can be distinguished from x. It takes into account whether x is a double or a single number. If x is an array, the result has the same size; each element corresponds to an element of the input.

eps('single') gives the smallest single positive number e such that 1single+e can be distinguished from 1single. eps('double') gives the same value as eps without input argument.

## **Examples**

```
eps
2.2204e-16
1 + eps - 1
2.2204e-16
eps / 2
1.1102e-16
1 + eps / 2 - 1
```

#### See also

```
inf, realmin, pi, i, j
```

### erf

Error function.

## **Syntax**

$$y = erf(x)$$

# **Description**

erf(x) gives the error function of x. Argument and result are real (imaginary part is discarded). The error function is defined as

$$\operatorname{erf}(x) = \frac{2}{\sqrt{\pi}} \int_0^x e^{-t^2} \, \mathrm{d}t$$

## **Example**

### See also

erfc, erfinv

### erfc

Complementary error function.

# Syntax

$$y = erfc(x)$$

# **Description**

erfc(x) gives the complementary error function of x. Argument and result are real (imaginary part is discarded). The complementary error function is defined as

$$\operatorname{erfc}(x) = 1 - \operatorname{erf}(x) = \frac{2}{\sqrt{\pi}} \int_{x}^{\infty} e^{-t^2} dt$$

# **Example**

```
erf, erfcx, erfcinv
```

### erfciny

Inverse complementary error function.

## **Syntax**

```
x = erfcinv(y)
```

# Description

erfcinv(y) gives the value x such that y=erfc(x). Argument and result are real (imaginary part is discarded). y must be in the range [0,2]; values outside this range give nan.

## **Example**

```
y = erfc(0.8)
y =
    0.2579
erfcinv(y)
    0.8
```

#### See also

erfc, erfinv

# erfcx

Scaled complementary error function.

### **Syntax**

```
y = erfcx(x)
```

# **Description**

erfcx(x) gives the scaled complementary error function of x, defined as  $exp(x^2)*erfc(x)$ . Argument and result are real (imaginary part is discarded).

```
erfcx(1)
0.4276
```

#### See also

erfc

### erfinv

Inverse error function.

# **Syntax**

```
x = erfinv(y)
```

## **Description**

erfinv(y) gives the value x such that y=erf(x). Argument and result are real (imaginary part is discarded). y must be in the range [-1,1]; values outside this range give nan.

# Example

```
y = erf(0.8)
y =
    0.7421
erfinv(y)
    0.8
```

#### See also

erf, erfcinv

# exp

Exponential.

## **Syntax**

```
y = exp(x)
```

# **Description**

 $\exp(x)$  is the exponential of x, i.e. 2.7182818284590446... $^{2}x$ .

```
exp([0,1,0.5j*pi])
1 2.7183 1j
```

### See also

log, expm1, expm, operator .^

# expm1

Exponential minus one.

### **Syntax**

```
y = expm1(x)
```

# **Description**

expm1(x) is exp(x)-1 with improved precision for small x.

# **Example**

```
expm1(1e-15)
1e-15
exp(1e-15)-1
1.1102e-15
```

#### See also

exp, log1p

### factor

Prime factors.

# **Syntax**

```
v = factor(n)
```

# **Description**

factor(n) gives a row vector which contains the prime factors of n in ascending order. Multiple prime factors are repeated.

```
factor(350)
2 5 5 7
```

#### See also

isprime

### factorial

Factorial.

### **Syntax**

```
y = factorial(n)
```

# **Description**

factorial (n) gives the factorial n! of nonnegative integer n. If the input argument is negative or noninteger, the result is NaN. The imaginary part is ignored.

## **Examples**

```
factorial(5)
  120
factorial([-1,0,1,2,3,3.14])
  nan  1   1   2   6   nan
```

#### See also

gamma, nchoosek

#### fix

Rounding towards 0.

## **Syntax**

```
y = fix(x)
```

# **Description**

fix(x) truncates the fractional part of x. If the argument is a complex number, the real and imaginary parts are handled separately.

```
fix(2.3)
2
fix(-2.6)
-2
```

#### See also

floor, ceil, round

### flintmax

Largest of the set of consecutive integers stored as floating point.

## **Syntax**

```
x = flintmax
x = flintmax(type)
```

## **Description**

flintmax gives the largest positive integer number in double precision such that all smaller integers can be represented in double precision.

flintmax(type) gives the largest positive integer number in double precision if type is 'double', or in single precision if type is 'single'. flintmax is 2^53 and flintmax('single') is 2^24.

## **Examples**

```
flintmax
9007199254740992
flintmax - 1
9007199254740991
flintmax + 1
9007199254740992
flintmax + 2
9007199254740994
```

#### See also

realmax, intmax

### floor

Rounding towards -infinity.

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### **Syntax**

$$y = floor(x)$$

## **Description**

floor(x) gives the largest integer smaller than or equal to x. If the argument is a complex number, the real and imaginary parts are handled separately.

## **Examples**

```
floor(2.3)
2
floor(-2.3)
-3
```

#### See also

ceil, fix, round, roundn

# gamma

Gamma function.

#### **Syntax**

```
y = qamma(x)
```

## **Description**

gamma(x) gives the gamma function of x. Argument and result are real (imaginary part is discarded). The gamma function is defined as

$$\Gamma(x) = \int_0^\infty t^{x-1} e^{-t} \, \mathrm{d}t$$

For positive integer values,  $\Gamma(n) = (n-1)!$ .

## **Examples**

```
gamma(5)
24
gamma(-3)
inf
gamma(-3.5)
0.2701
```

beta, gammaln, gammainc, factorial

# gammainc

Incomplete gamma function.

## **Syntax**

```
y = gammainc(x,a)
```

# **Description**

gammainc(x,a) gives the incomplete gamma function of x and a. Arguments and result are real (imaginary part is discarded). x must be nonnegative. The incomplete gamma function is defined as

gammainc(x, a) = 
$$\frac{1}{\Gamma(a)} \int_0^x t^{a-1} e^{-t} dt$$

## **Example**

```
gammainc(2,1.5)
0.7385
```

#### See also

gamma, gammaln, betainc

# gammaln

Logarithm of gamma function.

# **Syntax**

$$y = gammaln(x)$$

# Description

gammaln(x) gives the logarithm of the gamma function of x. Argument and result are real (imaginary part is discarded). gammaln does not rely on the computation of the gamma function to avoid overflows for large numbers.

```
gammaln(1000)
5905.2204
gamma(1000)
inf
```

### See also

gamma, gammainc, betaln

# gcd

Greatest common divisor.

# **Syntax**

```
q = gcd(a, b)
```

## **Description**

gcd(a,b) gives the greatest common divisor of integer numbers a and b.

# **Example**

```
gcd(72, 56)
8
```

### See also

lcm

# goldenratio

Golden ratio constant.

# **Syntax**

```
x = goldenratio
```

# **Description**

goldenratio is the golden ration  $(\sqrt{5} + 1)/2$ , up to the precision of its floating-point representation.

```
goldenratio
1.6180
```

### See also

```
pi, eps
```

# hypot

Hypotenuse.

### **Syntax**

```
c = hypot(a, b)
```

# **Description**

hypot(a,b) gives the square root of the square of a and b, or of their absolute value if they are complex. The result is always real. hypot avoids overflow when the result itself does not overflow.

## **Examples**

```
hypot(3, 4)
5
hypot([1,2,3+4j,inf], 5)
5.099 5.3852 5.831 inf
```

#### See also

```
sqrt, abs, norm
```



Imaginary unit.

### **Syntax**

```
i
j
1.23e4i
1.23e4j
```

### **Description**

i or j are the imaginary unit, i.e. the pure imaginary number whose square is -1. i and j are equivalent.

Used as a suffix appended without space to a number, i or j mark an imaginary number. They must follow the fractional part and the exponent, if any; for single-precision numbers, they must precede the single suffix.

To obtain a complex number i, you can write either i or 1i (or j or 1j). The second way is safer, because variables i and j are often used as indices and would hide the meaning of the built-in functions. The expression 1i is always interpreted as an imaginary constant number.

Imaginary numbers are displayed with i or j depending on the option set with the format command.

### **Examples**

```
i
   1j
format i
2i
   2i
2single + 5jsingle
2+5i (single)
```

### See also

imag, complex

# icdf

Inverse cumulative distribution function.

# **Syntax**

```
x = icdf(distribution,p)
x = icdf(distribution,p,a1)
x = icdf(distribution,p,a1,a2)
```

# **Description**

icdf(distribution,p) calculates the value of x such that cdf(distribution,x) is p. The distribution is specified with the first argument, a string; case is ignored ('t' and 'T' are equivalent). Additional arguments must be provided for some distributions. The distributions are given in the table below. Default values for the

parameters, when mentioned, mean that the parameter may be omitted.

Distribution	Name	<b>Parameters</b>
beta	beta	a and b
$\chi^2$	chi2	deg. of freedom $ u$
γ	gamma	shape $lpha$ and scale $\lambda$
F	f	deg. of freedom $\nu_1$ and $\nu_2$
lognormal	logn	mean (0) and st. dev. (1)
normal	norm	mean (0) and st. dev. (1)
Student's T	t	deg. of freedom $ u$
uniform	unif	limits of the range (0 and 1)

# **Example**

```
x = icdf('chi2', 0.6, 3)
x =
   2.9462
cdf('chi2', x, 3)
   0.6000
```

### See also

cdf, pdf, random

# imag

Imaginary part of a complex number.

# **Syntax**

```
im = imag(z)
```

# **Description**

imag(z) is the imaginary part of the complex number z, or 0 if z is real.

# **Examples**

```
imag(1+2j)
   2
imag(1)
   0
```

```
real, complex, i, j
```

### inf

Infinity.

### **Syntax**

```
x = inf
x = Inf
x = inf(n)
x = inf(n1,n2,...)
x = inf([n1,n2,...])
x = inf(..., type)
```

### **Description**

inf is the number which represents infinity. Most mathematical functions accept infinity as input argument and yield an infinite result if appropriate. Infinity and minus infinity are two different quantities.

With integer non-negative arguments, inf creates arrays whose elements are infinity. Arguments are interpreted the same way as zeros and ones.

The last argument of inf can be a string to specify the type of the result: 'double' for double-precision (default), or 'single' for single-precision.

### **Examples**

```
1/inf
0
-inf
-inf
```

#### See also

```
isfinite, isinf, nan, zeros, ones
```

### iscolumn

Test for a column vector.

### **Syntax**

```
b = iscolumn(x)
```

### **Description**

iscolumn(x) is true if the input argument is a column vector (real or complex 2-dimension array of any floating-point or integer type, character or logical value with second dimension equal to 1, or empty array), and false otherwise.

## **Examples**

```
iscolumn([1, 2, 3])
  false
iscolumn([1; 2])
  true
iscolumn(7)
  true
iscolumn([1, 2; 3, 4])
  false
```

#### See also

isrow, ismatrix, isscalar, isnumeric, size, ndims, length

# isfinite

Test for finiteness.

## **Syntax**

```
b = isfinite(x)
```

# **Description**

isfinite(x) is true if the input argument is a finite number (neither infinite nor nan), and false otherwise. The result is performed on each element of the input argument, and the result has the same size.

# **Example**

```
isfinite([0,1,nan,inf])
   T T F F
```

isinf, isnan

## isfloat

Test for a floating-point object.

### **Syntax**

```
b = isfloat(x)
```

## **Description**

isfloat(x) is true if the input argument is a floating-point type (double or single), and false otherwise.

### **Examples**

```
isfloat(2)
  true
isfloat(2int32)
  false
```

#### See also

isnumeric, isinteger

### isinf

Test for infinity.

## **Syntax**

```
b = isinf(x)
```

# **Description**

isinf(x) is true if the input argument is infinite (neither finite nor nan), and false otherwise. The result is performed on each element of the input argument, and the result has the same size.

# **Example**

```
isinf([0,1,nan,inf])
    F F F T
```

isfinite, isnan, inf

# isinteger

Test for an integer object.

## **Syntax**

```
b = isinteger(x)
```

## **Description**

isinteger(x) is true if the input argument is an integer type (including char and logical), and false otherwise.

### **Examples**

```
isinteger(2int16)
   true
isinteger(false)
   true
isinteger('abc')
   true
isinteger(3)
   false
```

#### See also

isnumeric, isfloat

### ismatrix

Test for a matrix.

## Syntax

```
b = ismatrix(x)
```

# **Description**

ismatrix(x) is true if the input argument is a matrix (real or complex 2-dimension array of any floating-point or integer type, character or logical value with, or empty array), and false otherwise.

```
ismatrix([1, 2, 3])
    true
ismatrix([1; 2])
    true
ismatrix(7)
    true
ismatrix([1, 2; 3, 4])
    true
ismatrix(ones([2,2,1])
    true
ismatrix(ones([1,2,2])
    false
```

#### See also

isrow, iscolumn, isscalar, isnumeric, isscalar, size, ndims, length

## isnan

Test for Not a Number.

## **Syntax**

```
b = isnan(x)
```

# **Description**

isnan(x) is true if the input argument is nan (not a number), and false otherwise. The result is performed on each element of the input argument, and the result has the same size.

# **Example**

```
isnan([0,1,nan,inf])
   F F T F
```

#### See also

isinf, nan

### isnumeric

Test for a numeric object.

## **Syntax**

```
b = isnumeric(x)
```

## **Description**

isnumeric(x) is true if the input argument is numeric (real or complex scalar, vector, or array), and false otherwise.

## **Examples**

```
isnumeric(pi)
  true
isnumeric('abc')
  false
```

#### See also

ischar, isfloat, isinteger, isscalar

# isprime

Prime number test.

# **Syntax**

```
b = isprime(n)
```

# **Description**

isprime(n) returns true if n is a prime number, or false otherwise. If n is a matrix, the test is applied to each element and the result is a matrix the same size.

# **Examples**

```
isprime(7)
  true
isprime([0, 2, 10])
  F T F
```

#### See also

factor

### isrow

Test for a row vector.

### **Syntax**

```
b = isrow(x)
```

# **Description**

isrow(x) is true if the input argument is a row vector (real or complex 2-dimension array of any floating-point or integer type, character or logical value with first dimension equal to 1, or empty array), and false otherwise.

### **Examples**

```
isrow([1, 2, 3])
   true
isrow([1; 2])
   false
isrow(7)
   true
isrow([1, 2; 3, 4])
   false
```

#### See also

iscolumn, ismatrix, isscalar, isnumeric, size, ndims, length

### isscalar

Test for a scalar number.

# **Syntax**

```
b = isscalar(x)
```

# **Description**

isscalar(x) is true if the input argument is scalar (real or complex number of any floating-point or integer type, character or logical value), and false otherwise.

```
isscalar(2)
  true
isscalar([1, 2, 5])
  false
```

### See also

isnumeric, isvector, ismatrix, size

### isvector

Test for a vector.

### **Syntax**

```
b = isvector(x)
```

## **Description**

isvector(x) is true if the input argument is a row or column vector (real or complex 2-dimension array of any floating-point or integer type, character or logical value with one dimension equal to 1, or empty array), and false otherwise.

## **Examples**

```
isvector([1, 2, 3])
   true
isvector([1; 2])
   true
isvector(7)
   true
isvector([1, 2; 3, 4])
  false
```

### See also

isnumeric, isscalar, iscolumn, isrow, size, ndims, length

### **Icm**

Least common multiple.

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## **Syntax**

```
q = lcm(a, b)
```

## **Description**

lcm(a,b) gives the least common multiple of integer numbers a and b.

## **Example**

```
lcm(72, 56)
504
```

#### See also

gcd

# log

Natural (base e) logarithm.

# **Syntax**

```
y = log(x)
```

# **Description**

log(x) gives the natural logarithm of x. It is the inverse of exp. The result is complex if x is not real positive.

# Example

```
log([-1,0,1,10,1+2j])
0+3.1416j inf 0 2.3026 0.8047+1.1071j
```

### See also

```
log10, log2, log1p, reallog, exp
```

# log10

Decimal logarithm.

## **Syntax**

```
y = log10(x)
```

## **Description**

log10(x) gives the decimal logarithm of x, defined by log10(x) = log(x)/log(10). The result is complex if x is not real positive.

## **Example**

```
log10([-1,0,1,10,1+2j])
0+1.3644j inf 0 1 0.3495+0.4808j
```

### See also

log, log2

# log1p

Logarithm of x plus one.

### **Syntax**

```
y = log1p(x)
```

# **Description**

log1p(x) is log(1+x) with improved precision for small x.

## Example

```
log1p(1e-15)
1e-15
log(1 + 1e-15)
1.1102e-15
```

### See also

log, expm1

# log2

Base 2 logarithm.

## **Syntax**

```
y = log2(x)
```

# **Description**

log2(x) gives the base 2 logarithm of x, defined as log2(x)=log(x)/log(2). The result is complex if x is not real positive.

## **Example**

```
log2([1, 2, 1024, 2000, -5])
0 1 10 10.9658 2.3219+4.5324j
```

#### See also

log, log10

### mod

Modulo.

### **Syntax**

```
m = mod(x, y)
```

# **Description**

mod(x,y) gives the modulo of x divided by y, i.e. a number m between 0 and y such that x = q\*y+m with integer q. Imaginary parts, if they exist, are ignored.

## **Examples**

```
mod(10,7)

3

mod(-10,7)

4

mod(10,-7)

-4

mod(-10,-7)

-3
```

rem

#### nan

Not a Number.

## **Syntax**

```
x = nan
x = NaN
x = nan(n)
x = nan(n1,n2,...)
x = nan([n1,n2,...])
x = nan(..., type)
```

### **Description**

NaN (Not a Number) is the result of the primitive floating-point functions or operators called with invalid arguments. For example, 0/0, inf-inf and 0\*inf all result in NaN. When used in an expression, NaN propagates to the result. All comparisons involving NaN result in false, except for comparing NaN with any number for inequality, which results in true.

Contrary to built-in functions usually found in the underlying operating system, many functions which would result in NaNs give complex numbers when called with arguments in a certain range.

With integer non-negative arguments, nan creates arrays whose elements are NaN. Arguments are interpreted the same way as zeros and ones.

The last argument of nan can be a string to specify the type of the result: 'double' for double-precision (default), or 'single' for single-precision.

### **Examples**

```
nan
nan
0*nan
nan
nan==nan
false
nan==nan
true
log(-1)
0+3.1416j
```

#### See also

inf, isnan, zeros, ones

### nchoosek

Binomial coefficient.

## **Syntax**

```
b = nchoosek(n, k)
```

# **Description**

nchoosek(n,k) gives the number of combinations of n objects taken k at a time. Both n and k must be nonnegative integers with k < n.

## **Examples**

```
nchoosek(10,4)
210
nchoosek(10,6)
210
```

#### See also

factorial, gamma

#### nthroot

Real nth root.

## **Syntax**

```
y = nthroot(x,n)
```

# Description

nthroot(x,n) gives the real nth root of real number x. If x is positive, it is  $x.^(1./n)$ ; if x is negative, it is  $-abs(x).^(1./n)$  if n is an odd integer, or NaN otherwise.

#### **Example**

```
nthroot([-2,2], 3)
-1.2599 1.2599
[-2,2] .^ (1/3)
0.6300+1.0911i 1.2599
```

#### See also

```
operator .^, realsqrt, sqrt
```

# pdf

Probability density function.

### **Syntax**

```
y = pdf(distribution,x)
y = pdf(distribution,x,a1)
y = pdf(distribution,x,a1,a2)
```

## **Description**

pdf(distribution,x) gives the probability of a density function. The distribution is specified with the first argument, a string; case is ignored ('t' and 'T' are equivalent). Additional arguments must be provided for some distributions. See cdf for the list of distributions.

#### See also

cdf. random

## pi

Constant  $\pi$ .

## **Syntax**

```
x = pi
```

# Description

pi is the number  $\pi$ , up to the precision of its floating-point representation.

### **Example**

```
exp(1j * pi)
-1
```

#### See also

goldenratio, i, j, eps

# pol2cart

Polar to Cartesian coordinates transform.

## **Syntax**

```
(x, y) = pol2cart(phi, r)
(x, y, z) = pol2cart(phi, r, z)
```

## **Description**

(x,y)=pol2cart(phi,r) transforms polar coordinates phi and r to Cartesian coordinates x and y such that  $x=r\cos(\varphi)$  and  $x=r\sin(\varphi)$ . (x,y,z)=pol2cart(phi,r,z) transforms cylindrical coordinates to Cartesian coordinates, leaving z unchanged.

# **Example**

```
(x, y) = pol2cart(1, 2)

x =

1.0806

y =

1.6829
```

#### See also

cart2pol, cart2sph, sph2cart

#### random

Random generator for distribution function.

### **Syntax**

```
x = random(distribution)
x = random(distribution, a1)
x = random(distribution, a1, a2)
x = random(.... size)
```

### **Description**

random(distribution,a1,a2) calculates a pseudo-random number whose distribution function is specified by name distribution and parameters a1 and a2 (some distributions have a single parameter). The distributions are given in the table below. Unlike in functions pdf, cdf and icdf, parameters do not have default values and must be specified.

Additional input arguments specify the size of the result, either as a vector (or a single scalar for a square matrix) or as scalar values. The result is an array of the specified size where each value is an independent pseudo-random variable. The default size is 1 (scalar).

If the parameters are arrays, the result is an array of the same size and each element is an independent pseudo-random variable whose distribution has its parameters at the corresponding position. The size, if specified, must be the same.

Distribution	Name	Parameters
beta	beta	a and b
Cauchy	cauchy	a and b
<i>X</i> _	chi	deg. of freedom $ u$
$\chi^2$	chi2	deg. of freedom $ u$
γ	gamma	shape $lpha$ and $\lambda$
exponential	exp	mean
F	f	deg. of freedom $ u_1$ and $ u_2$
half-normal	half-normal	9
Laplace	laplace	mean and scale
lognormal	logn	mean and st. dev.
Nakagami	nakagami	$\mu$ and $\omega$
normal	norm	mean and st. dev.
Rayleigh	rayl	b
Student's T	t	deg. of freedom $ u$
uniform	unif	limits of the range
Weibull	weib	shape a and scale b

# Example

Array of 5 pseudo-random numbers whose distribution is  $\chi^2$  with 3 degrees of freedom:

```
random('chi2', 3, [1, 5])
1.6442 0.4164 2.0272 2.7962 4.5896
```

#### See also

```
pdf, cdf, icdf, rand, randn, rng
```

#### rat

Rational approximation.

### **Syntax**

```
(num, den) = rat(x)
(num, den) = rat(x, tol)
(num, den) = rat(x, tol=tol)
```

### **Description**

rat(x,tol) returns the numerator and the denominator of a rational approximation of real number x with the smallest integer numerator and denominator which fulfil absolute tolerance tol. If the input argument x is an array, output arguments are arrays of the same size. Negative numbers give a negative numerator. The tolerance can be passed as a named argument.

With one input argument, rat(x) uses tolerance tol=1e-6\*norm(x,1). With one output argument, rat(x) gives the rational approximation itself as a floating-point number.

With command format rat, all numeric results as displayed as rational aproximations with the default tolerance, including complex numbers.

## **Example**

```
(num,den) = rat(pi)
num =
   355
den =
   113
num/den
  3.141592920353982
```

#### See also

format

#### real

Real part of a complex number.

### **Syntax**

```
re = real(z)
```

## **Description**

real(z) is the real part of the complex number z, or z if z is real.

## **Examples**

```
real(1+2j)
1
real(1)
1
```

#### See also

imag, complex

# reallog

Real natural (base e) logarithm.

## **Syntax**

```
y = reallog(x)
```

# Description

reallog(x) gives the real natural logarithm of x. It is the inverse of exp for real numbers. The imaginary part of x is ignored. The result is NaN if x is negative.

# Example

```
reallog([-1,0,1,10,1+2j])
nan inf 0 2.3026 0
```

#### See also

log, realpow, realsqrt, exp

## realmax realmin

Largest and smallest real numbers.

### **Syntax**

```
x = realmax
x = realmax(n)
x = realmax(n1,n2,...)
x = realmax([n1,n2,...])
x = realmax(..., type)
x = realmin
x = realmin(...)
```

### **Description**

realmax gives the largest positive real number in double precision. realmin gives the smallest positive real number in double precision which can be represented in normalized form (i.e. with full mantissa precision).

With integer non-negative arguments, realmax and realmin create arrays whose elements are all set to the respective value. Arguments are interpreted the same way as zeros and ones.

The last argument of realmax and realmin can be a string to specify the type of the result: 'double' for double-precision (default), or 'single' for single-precision.

## **Examples**

```
realmin
2.2251e-308
realmin('single')
1.1755e-38
realmax
1.7977e308
realmax('single')
3.4028e38single
realmax + eps(realmax)
inf
```

#### See also

```
inf, ones, zeros, eps, flintmax
```

# realpow

Real power.

### **Syntax**

```
z = realpow(x, y)
```

## **Description**

realpow(x, y) gives the real value of x to the power y. The imaginary parts of x and y are ignored. The result is NaN if it is not defined for the input arguments. If the arguments are arrays, their size must match or one of them must be a scalar number; the power is performed element-wise.

#### See also

```
operator .^, reallog, realsqrt
```

# realsqrt

Real square root.

### **Syntax**

```
y = realsqrt(x)
```

## **Description**

realsqrt(x) gives the real square root of x. The imaginary part of x is ignored. The result is NaN if x is negative.

## Example

```
realsqrt([-1,0,1,10,1+2j])
nan 0 1 3.1623 1
```

#### See also

sgrt, reallog, realpow, nthroot

#### rem

Remainder of a real division.

#### Syntax

```
r = rem(x, y)
```

## **Description**

rem(x,y) gives the remainder of x divided by y, i.e. a number r between 0 and sign(x)\*abs(y) such that x = q\*y+r with integer q. Imaginary parts, if they exist, are ignored.

## **Examples**

```
rem(10,7)

3

rem(-10,7)

-3

rem(10,-7)

3

rem(-10,-7)
```

#### See also

mod

#### round

Rounding to the nearest integer.

# **Syntax**

```
y = round(x)
```

# **Description**

round(x) gives the integer nearest to x. If the argument is a complex number, the real and imaginary parts are handled separately.

# **Examples**

```
round(2.3)
2
round(2.6)
3
round(-2.3)
-2
```

#### See also

floor, ceil, fix, roundn

#### roundn

Rounding to a specified precision.

#### **Syntax**

```
y = roundn(x, n)
```

### **Description**

roundn(x,n) rounds x to the nearest multiple of  $10^n$ . If argument x is a complex number, the real and imaginary parts are handled separately. roundn(x,0) gives the same result as round(x).

Argument n must be a real integer. If x and/or n are arrays, rounding is performed separately on each element.

### **Examples**

```
roundn(pi, -2)
   3.1400
roundn(1000 * pi, 1)
   3140
roundn(pi, [-3, -1])
   3.1420   3.1000
```

#### See also

```
round, floor, ceil, fix
```

# sign

Sign of a real number or direction of a complex number.

# Syntax

```
s = sign(x)
z2 = sign(z1)
```

# **Description**

With a real argument, sign(x) is 1 if x>0, 0 if x==0, or -1 if x<0. With a complex argument, sign(z1) is a complex value with the same phase as z1 and whose magnitude is 1.

## **Examples**

```
sign(-2)
-1
sign(1+1j)
0.7071+0.7071j
sign([0, 5])
0 1
```

### See also

abs, angle

#### sec

Secant.

### **Syntax**

```
y = sec(x)
```

## **Description**

sec(x) gives the secant of x, which is complex if x is.

### See also

asec, sech, cos

### sech

Hyperbolic secant.

## **Syntax**

```
y = sech(x)
```

# **Description**

acot(x) gives the hyperbolic secant of x, which is complex if x is.

#### See also

asech, sec, cosh

#### sin

Sine.

## **Syntax**

```
y = \sin(x)
```

## **Description**

sin(x) gives the sine of x, which is complex if x is complex.

# **Example**

```
sin(2)
0.9093
```

#### See also

cos, asin, sinh

## sinc

Sinc.

## **Syntax**

```
y = sinc(x)
```

# **Description**

sinc(x) gives the sinc of x, i.e. sin(pi\*x)/(pi\*x) if x=0 or 1 if x=0. The result is complex if x is complex.

# **Example**

```
sinc(1.5)
-0.2122
```

#### See also

sin, sinh

# single

Conversion to single-precision numbers.

### **Syntax**

```
B = single(A)
```

## **Description**

single(A) converts number or array A to single precision. A can be any kind of numeric value (real, complex, or integer), or a character or logical array.

Single literal numbers can be entered as a floating-point number with the single suffix.

### **Examples**

```
single(pi)
  3.1416single
single('AB')
  1x2 single array
  65 66
3.7e4single
  37000single
```

#### See also

double, uint8 and related functions, operator +, setstr, char, logical

#### sinh

Hyperbolic sine.

# **Syntax**

```
y = sinh(x)
```

# **Description**

sinh(x) gives the hyperbolic sine of x, which is complex if x is complex.

#### **Example**

```
sinh(2)
3.6269
```

#### See also

cosh, asinh, sin

# sph2cart

Spherical to Cartesian coordinates transform.

### **Syntax**

```
(x, y, z) = sph2cart(phi, theta, r)
```

## **Description**

(x,y,z)=sph2cart(phi,theta,r) transforms polar coordinates phi, theta, and r to Cartesian coordinates x, y, and z such that  $x=r\cos(\varphi)\cos(\vartheta)$ ,  $y=r\sin(\varphi)\cos(\vartheta)$ , and  $z=r\sin(\vartheta)$ .

### **Example**

```
(x, y, z) = sph2cart(1, 2, 3)
x =
    -0.6745
y =
    -1.0505
z =
    2.7279
```

#### See also

cart2pol, cart2sph, pol2cart

# sqrt

Square root.

### Syntax

```
r = sqrt(z)
```

## **Description**

sqrt(z) gives the square root of z, which is complex if z is not real positive.

### **Examples**

```
sqrt(4)
  2
sqrt([1 4 -9 3+4j])
  1 2 3j 2+1j
```

#### See also

realsqrt, sqrtm, chol

# swapbytes

Conversion between big-endian and little-endian representation.

### **Syntax**

```
Y = swapbytes(X)
```

# **Description**

swapbytes (X) swaps the bytes representing number X. If X is an array, each number is swapped separately. The imaginary part, if any, is discarded. X can be of any numeric type. swapbytes is its own inverse for real numbers.

# **Example**

```
swapbytes(1uint32)
16777216uint32
```

#### See also

```
typecast, cast
```

#### tan

Tangent.

## **Syntax**

```
y = tan(x)
```

# **Description**

tan(x) gives the tangent of x, which is complex if x is complex.

## **Example**

```
tan(2)
-2.185
```

#### See also

atan, tanh

### tanh

Hyperbolic tangent.

# **Syntax**

```
y = tanh(x)
```

# **Description**

tanh(x) gives the hyperbolic tangent of x, which is complex if x is complex.

## **Example**

```
tanh(2)
0.964
```

### See also

atanh, tan

# typecast

Type conversion with same binary representation.

### **Syntax**

```
Y = typecast(X, type)
```

### **Description**

typecast(X, type) changes the numeric array X to the type given by string type, which can be 'double', 'single', 'int8' or any other signed or unsigned integer type, 'char', or 'logical'. The binary representation in memory is preserved. The imaginary part, if any, is discarded. Depending on the conversion, the number of elements is changed, so that the array size in bytes in preserved. The result is a row vector if X is a scalar or a row vector, or a column vector otherwise. The result depends on the computer architecture.

### **Example**

```
typecast(1uint32, 'uint8')
  1x4 uint8 array
    0    0    1
typecast(pi, 'uint8')
  1x8 uint8 array
  64    9    33    251    84    68    45    24
```

#### See also

swapbytes, bwrite, sread, cast

# 6.18 Linear Algebra

# addpol

Addition of two polynomials.

# Syntax

```
p = addpol(p1,p2)
```

# **Description**

addpol(p1,p2) adds two polynomials p1 and p2. Each polynomial is given as a vector of coefficients, with the highest power first; e.g.,  $x^2 + 2x - 3$  is represented by [1,2,-3]. Row vectors and column vectors are accepted, as well as matrices made of row vectors or column vectors, provided one matrix is not larger in one dimension and

smaller in the other one. addpol is equivalent to the plain addition when both arguments have the same size.

## **Examples**

```
addpol([1,2,3], [2,5])
1 4 8
addpol([1,2,3], -[2,5]) % subtraction
1 0 -2
addpol([1,2,3;4,5,6], [1;1])
1 2 4
4 5 7
```

#### See also

conv, deconv, operator +

### balance

Diagonal similarity transform for balancing a matrix.

#### **Syntax**

```
B = balance(A)
(T, B) = balance(A)
```

## **Description**

balance(A) applies a diagonal similarity transform to the square matrix A to make the rows and columns as close in norm as possible. Balancing may reduce the 1-norm of the matrix, and improves the accuracy of the computed eigenvalues and/or eigenvectors. To avoid round-off errors, balance scales A with powers of 2.

balance returns the balanced matrix B which has the same eigenvalues and singular values as A, and optionally the diagonal scaling matrix T such that  $T\A*T=B$ .

### **Example**

```
A = [1,2e6;3e-6,4];

(T,B) = balance(A)

T =

16384 0

0 3.125e-2

B =

1 3.8147

1.5729 4
```

#### See also

eig

#### care

Continuous-time algebraic Riccati equation.

### **Syntax**

```
(X, L, K) = care(A, B, Q)

(X, L, K) = care(A, B, Q, R)

(X, L, K) = care(A, B, Q, R, S)

(X, L) = care(A, S, Q, true)
```

### **Description**

care(A,B,Q) calculates the stable solution X of the following continuous-time algebraic Riccati equation:

$$A'X + XA - XBB'X + O = 0$$

All matrices are real; Q and X are symmetric.

With four input arguments, care(A,B,Q,R) (with R real symmetric) solves the following Riccati equation:

$$A'X + XA - XBR^{-1}B'X + Q = 0$$

With five input arguments, care(A,B,Q,R,S) solves the following equation:

$$A'X + XA - (S + XB)R^{-1}(S' + B'X) + Q = 0$$

With two or three output arguments, (X,L,K) = care(...) also returns the gain matrix K defined as

$$K = R^{-1}B'X$$

and the column vector of closed-loop eigenvalues

$$L = eig(A - BK)$$

care(A,S,Q,true) with up to two output arguments is equivalent to care(A,B,Q) or care(A,B,Q,false) with S=B\*B'.

#### **Example**

```
A = [-4,2;1,2];
B = [0;1];
C = [2, -1];
Q = C' * C;
R = 5;
(X, L, K) = care(A, B, Q, R)
      1.07
            3.5169
    3.5169
             23.2415
  L =
   -4.3488
   -2.2995
 K =
    0.7034 4.6483
A' * X + X * A - X * B / R * B' * X + 0
  1.7319e-14 1.1369e-13
  8.5265e-14 6.2528e-13
```

#### See also

dare

#### chol

Cholesky decomposition.

#### **Syntax**

```
M2 = chol(M1)
```

## Description

If a square matrix M1 is symmetric (or hermitian) and positive definite, it can be decomposed into the following product:

```
M_1=M_2^\prime M_2
```

where  $\tilde{\text{M2}}$  is an upper triangular matrix. The Cholesky decomposition can be seen as a kind of square root.

The part of M1 below the main diagonal is not used, because M1 is assumed to be symmetric or hermitian. An error occurs if M1 is not positive definite.

# **Example**

```
M = chol([5,3;3,8])
M =
```

```
2.2361 1.3416
0 2.4900
M'*M
5 3
3 8
```

#### See also

inv, sqrtm

#### cond

Condition number of a matrix.

### **Syntax**

```
x = cond(M)
```

## **Description**

cond (M) returns the condition number of matrix M, i.e. the ratio of its largest singular value divided by the smallest one, or infinity for singular matrices. The larger the condition number, the more ill-conditioned the inversion of the matrix.

# **Examples**

```
cond([1, 0; 0, 1])
1
cond([1, 1; 1, 1+1e-3])
4002.0008
```

#### See also

svd, rank

#### conv

Convolution or polynomial multiplication.

## **Syntax**

```
v = conv(v1,v2)
M = conv(M1,M2)
M = conv(M1,M2,dim)
M = conv(...,kind)
```

### **Description**

conv(v1,v2) convolves the vectors v1 and v2, giving a vector whose length is length(v1)+length(v2)-1, or an empty vector if v1 or v2 is empty. The result is a row vector if both arguments are row vectors, and a column vector if both arguments are column vectors. Otherwise, arguments are considered as matrices.

conv(M1,M2) convolves the matrices M1 and M2 column by columns. conv(M1,M2,dim) convolves along the dimension dim, 1 for columns and 2 for rows. If one of the matrices has only one column, or one row, it is repeated to match the size of the other argument.

Let n1 and n2 be the number of elements in M1 and M2, respectively, along the convolution dimension. By default, the result has n1+n2-1 elements along the convolution dimension. An additional string argument kind can specify a different number of elements in the result: with kind='same', the result has n1 elements (M has the same size as M1, i.e. M1 is filtered by the finite impulse response filter M2). With kind='valid', the result has n1-n2+1 elements, i.e. result elements impacted by boundaries are discarded. kind='full' produce the same result as if kind is missing.

### **Examples**

```
conv([1,2],[1,2,3])
  1 4 7 6
conv([1,2],[1,2,3;4,5,6],2)
  1 4 7 6
  4 13 16 12
conv([1,2,5,8,3],[1,2,1],'full')
  1 4 10 20 24 14 3
conv([1,2,5,8,3],[1,2,1],'same')
  4 10 20 24 14
conv([1,2,5,8,3],[1,2,1],'valid')
  10 20 24
```

#### See also

deconv, filter, addpol, conv2

#### conv2

Two-dimensions convolution of matrices.

### **Syntax**

```
M = conv2(M1,M2)
M = conv2(M1,M2,kind)
```

## **Description**

conv2(M1,M2) convolves the matrices M1 and M2 along both directions. The optional third argument specifies how to crop the result. Let (nr1,nc1)=size(M1) and (nr2,nc2)=size(M2). With kind='full' (default value), the result M has nr1+nr2-1 lines and nc1+nc2-1 columns. With kind='same', the result M has nr1 lines and nc1 columns; this options is very useful if M1 represents equidistant samples in a plane (e.g. pixels) to be filtered with the finite-impulse response 2-d filter M2. With kind='valid', the result M has nr1-nr2+1 lines and nc1-nc2+1 columns, or is the empty matrix []; if M1 represents data filtered by M2, the borders where the convolution sum is not totally included in M1 are removed.

## **Examples**

```
conv2([1,2,3;4,5,6;7,8,9],[1,1,1;1,1,1;1,1,1])
    1    3    6    5    3
    5    12    21    16    9
    12    27    45    33    18
    11    24    39    28    15
    7    15    24    17    9
conv2([1,2,3;4,5,6;7,8,9],[1,1,1;1,1,1;1,1,1],'same')
    12    21    16
    27    45    33
    24    39    28
conv2([1,2,3;4,5,6;7,8,9],[1,1,1;1,1,1;1,1,1],'valid')
    45
```

#### See also

conv

#### COV

Covariance.

### **Syntax**

```
M = cov(data)
M = cov(data, false)
M = cov(data, true)
```

# **Description**

cov(data) returns the best unbiased estimate m-by-m covariance matrix of the n-by-m matrix data for a normal distribution. Each row of

data is an observation where n quantities were measured. The covariance matrix is symmetric if data is real, and hermitian if data is complex (i.e. M==M'). The diagonal is the variance of each column of data.

```
cov(data, false) is the same as cov(data).
```

cov(data,true) returns the m-by-m covariance matrix of the n-by-m matrix data which contains the whole population.

#### **Example**

```
A = [1,2;2,4;3,5];
cov(A)
1 1.5
1.5 2.3333
```

The diagonal elements are the variance of the columns of A:

```
var(A)
1 2.3333
```

The covariance matrix can be computed as follows:

```
n = size(A, 1);
A1 = A - repmat(mean(A, 1), n, 1);
(A1' * A1) / (n - 1)
    1 1.5
    1.5 2.3333
```

#### See also

mean, var

#### cross

Cross product.

#### **Syntax**

```
v3 = cross(v1, v2)

v3 = cross(v1, v2, dim)
```

# Description

cross(v1,v2) gives the cross products of vectors v1 and v2. v1 and v2 must be row or columns vectors of three components, or arrays of the same size containing several such vectors. When there is ambiguity, a third argument dim may be used to specify the dimension of vectors: 1 for column vectors, 2 for row vectors, and so on.

#### **Examples**

```
cross([1; 2; 3], [0; 0; 1])
2
-1
0
cross([1, 2, 3; 7, 1, -3], [4, 0, 0; 0, 2, 0], 2)
0 12 -8
6 0 14
```

#### See also

dot, operator \*, det

#### cummax

Cumulative maximum.

### **Syntax**

```
M2 = cummax(M1)
M2 = cummax(M1,dim)
M2 = cummax(...,dir)
```

# **Description**

cummax(M1) returns a matrix M2 of the same size as M1, whose elements M2(i,j) are the maximum of all the elements M1(k,j) with k <= i. cummax(M1,dim) operates along the dimension dim (columnwise if dim is 1, row-wise if dim is 2).

An optional string argument dir specifies the processing direction. If it is 'reverse' or begins with 'r', cummax processes elements in reverse order, from the last one to the first one, along the processing dimension. If it is 'forward' or begins with 'f', it processes elements as if not specified, in the forward direction.

# **Examples**

```
cummax([1,2,3;5,1,4;2,8,7])
   1   2   3
   5   2   4
   5   8   7
cummax([1,2,3;5,1,4;2,8,7], 2)
   1   2   3
   5   5   5
   2   8   8
```

#### See also

max, cummin, cumsum, cumprod

#### cummin

Cumulative minimum.

#### **Syntax**

```
M2 = cummin(M1)
M2 = cummin(M1,dim)
M2 = cummin(...,dir)
```

### **Description**

cummin(M1) returns a matrix M2 of the same size as M1, whose elements M2(i,j) are the minimum of all the elements M1(k,j) with  $k \le i$ . cummin(M1,dim) operates along the dimension dim (columnwise if dim is 1, row-wise if dim is 2).

An optional string argument dir specifies the processing direction. If it is 'reverse' or begins with 'r', cummin processes elements in reverse order, from the last one to the first one, along the processing dimension. If it is 'forward' or begins with 'f', it processes elements as if not specified, in the forward direction.

#### See also

min, cummax, cumsum, cumprod

# cumprod

Cumulative products.

### **Syntax**

```
M2 = cumprod(M1)
M2 = cumprod(M1,dim)
M2 = cumprod(...,dir)
```

# **Description**

cumprod(M1) returns a matrix M2 of the same size as M1, whose elements M2(i,j) are the product of all the elements M1(k,j) with k <= i.

cumprod(M1,dim) operates along the dimension dim (column-wise if dim is 1. row-wise if dim is 2).

An optional string argument dir specifies the processing direction. If it is 'reverse' or begins with 'r', cumprod processes elements in reverse order, from the last one to the first one, along the processing dimension. If it is 'forward' or begins with 'f', it processes elements as if not specified, in the forward direction.

### **Examples**

```
cumprod([1,2,3;4,5,6])
    1    2    3
    4    10    18
cumprod([1,2,3;4,5,6],2)
    1    2    6
    4    20    120
```

#### See also

prod, cumsum, cummax, cummin

#### cumsum

Cumulative sums.

### **Syntax**

```
M2 = cumsum(M1)
M2 = cumsum(M1,dim)
M2 = cumsum(...,dir)
```

# **Description**

cumsum(M1) returns a matrix M2 of the same size as M1, whose elements M2(i,j) are the sum of all the elements M1(k,j) with k<=i. cumsum(M1,dim) operates along the dimension dim (column-wise if dim is 1. row-wise if dim is 2).

An optional string argument dir specifies the processing direction. If it is 'reverse' or begins with 'r', cumsum processes elements in reverse order, from the last one to the first one, along the processing dimension. If it is 'forward' or begins with 'f', it processes elements as if not specified, in the forward direction.

#### **Examples**

```
cumsum([1,2,3;4,5,6])
   1 2 3
   5 7 9
cumsum([1,2,3;4,5,6],2)
   1 3 6
   4 9 15
cumsum([1,2,3;4,5,6],2,'r')
   6  5 3
   15 11 6
```

#### See also

sum, diff, cumprod, cummax, cummin

### dare

Discrete-time algebraic Riccati equation.

### **Syntax**

$$(X, L, K) = dare(A, B, Q)$$
  
 $(X, L, K) = dare(A, B, Q, R)$ 

# Description

dare(A,B,Q) calculates the stable solution X of the following discretetime algebraic Riccati equation:

$$X = A'XA - A'XB(B'XB + I)^{-1}B'XA + O$$

All matrices are real; Q and X are symmetric.

With four input arguments, dare(A,B,Q,R) (with R real symmetric) solves the following Riccati equation:

$$X = A'XA - A'XB(B'XB + R)^{-1}B'XA + O$$

With two or three output arguments, (X,L,K) = dare(...) also returns the gain matrix K defined as

$$K = (B'XB + R)^{-1}B'XA$$

and the column vector of closed-loop eigenvalues

$$L = eig(A - BK)$$

#### **Example**

#### See also

care

#### deconv

Deconvolution or polynomial division.

## **Syntax**

```
q = deconv(a,b)
(q,r) = deconv(a,b)
```

# **Description**

(q,r)=deconv(a,b) divides the polynomial a by the polynomial b, resulting in the quotient q and the remainder r. All polynomials are given as vectors of coefficients, highest power first. The degree of the remainder is strictly smaller than the degree of b. deconv is the inverse of conv: a = addpol(conv(b,q),r).

# **Examples**

```
[q,r] = deconv([1,2,3,4,5],[1,3,2])
q =
   1 -1 4
r =
```

```
-6 -3
addpol(conv(q,[1,3,2]),r)
1 2 3 4 5
```

#### See also

```
conv, filter, addpol
```

#### det

Determinant of a square matrix.

## **Syntax**

```
d = det(M)
```

# **Description**

det(M) is the determinant of the square matrix M, which is 0 (up to the rounding errors) if M is singular. The function rank is a numerically more robust test for singularity.

## **Examples**

```
det([1,2;3,4])
-2
det([1,2;1,2])
```

#### See also

```
poly, rank
```

#### diff

Differences.

## **Syntax**

```
dm = diff(A)
dm = diff(A,n)
dm = diff(A,n,dim)
dm = diff(A,[],dim)
```

## **Description**

diff(A) calculates the differences between each elements of the columns of matrix A, or between each elements of A if it is a row vector.

diff(A,n) calculates the n:th order differences, i.e. it repeats n times the same operation. Up to a scalar factor, the result is an approximation of the n:th order derivative based on equidistant samples.

diff(A,n,dim) operates along dimension dim. If the second argument n is the empty matrix [], the default value of 1 is assumed.

## **Examples**

```
diff([1,3,5,4,8])
  2 2 -1 4
diff([1,3,5,4,8],2)
  0 -3 5
diff([1,3,5;4,8,2;3,9,8],1,2)
  2 2
  4 -6
  6 -1
```

#### See also

cumsum

# dlyap

Discrete-time Lyapunov equation.

# **Syntax**

$$X = dlyap(A, C)$$

# **Description**

dlyap(A,C) calculates the solution X of the following discrete-time Lyapunov equation:

$$AXA' - X + C = 0$$

All matrices are real.

#### **Example**

```
A = [3,1,2;1,3,5;6,2,1];

C = [7,1,2;4,3,5;1,2,1];

X = dlyap(A, C)

X =

-1.0505 3.2222 -1.2117

3.2317 -11.213 4.8234

-1.4199 5.184 -2.7424
```

#### See also

lyap, dare

#### dot

Scalar product.

#### **Syntax**

```
v3 = dot(v1, v2)

v3 = dot(v1, v2, dim)
```

## **Description**

dot(v1,v2) gives the scalar products of vectors v1 and v2. v1 and v2 must be row or columns vectors of same length, or arrays of the same size; then the scalar product is performed along the first dimension not equal to 1. A third argument dim may be used to specify the dimension the scalar product is performed along.

With complex values, complex conjugate values of the first array are multiplied with values of the second array.

## **Examples**

```
dot([1; 2; 3], [0; 0; 1])
3
dot([1, 2, 3; 7, 1, -3], [4, 0, 0; 0, 2, 0], 2)
4
2
dot([1; 2i], [3i; 5])
0 - 7i
```

#### See also

cross, operator \*, det

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# eig

Eigenvalues and eigenvectors of a matrix.

### **Syntax**

```
e = eig(M)
(V,D) = eig(M)
```

### **Description**

eig(M) returns the vector of eigenvalues of the square matrix M. (V,D) = eig(M) returns a diagonal matrix D of eigenvalues and a matrix V whose columns are the corresponding eigenvectors. They are such that M\*V = V\*D.

### **Examples**

Eigenvalues as a vector:

```
eig([1,2;3,4])
-0.3723
5.3723
```

Eigenvectors, and eigenvalues as a diagonal matrix:

```
(V,D) = eig([1,2;2,1])

V =

0.7071 0.7071

-0.7071 0.7071

D =

-1 0

0 3
```

Checking that the result is correct:

```
[1,2;2,1] * V
-0.7071 2.1213
0.7071 2.1213
V * D
-0.7071 2.1213
0.7071 2.1213
```

#### See also

schur, svd, det, roots

### expm

Exponential of a square matrix.

### **Syntax**

```
M2 = expm(M1)
```

### **Description**

expm(M) is the exponential of the square matrix M, which is usually different from the element-wise exponential of M given by exp.

## **Examples**

```
expm([1,1;1,1])
4.1945 3.1945
3.1945 4.1945
exp([1,1;1,1])
2.7183 2.7183
2.7183 2.7183
```

#### See also

```
logm, operator ^, exp
```

#### fft

Fast Fourier Transform.

## **Syntax**

```
F = fft(f)
F = fft(f,n)
F = fft(f,n,dim)
```

# **Description**

fft(f) returns the discrete Fourier transform (DFT) of the vector f, or the DFT's of each columns of the array f. With a second argument n, the n first values are used; if n is larger than the length of the data, zeros are added for padding. An optional argument dim gives the dimension along which the DFT is performed; it is 1 for calculating the DFT of the columns of f, 2 for its rows, and so on. fft(f,[],dim) specifies the dimension without resizing the array.

fft is based on a mixed-radix Fast Fourier Transform if the data length is non-prime. It can be very slow if the data length has large prime factors or is a prime number.

The coefficients of the DFT are given from the zero frequency to the largest frequency (one point less than the inverse of the sampling period). If the input f is real, its DFT has symmetries, and the first half contain all the relevant information.

#### **Examples**

```
fft(1:4)

10 -2+2j -2 -2-2j

fft(1:4, 3)

6 -1.5+0.866j -1.5-0.866j
```

#### See also

ifft

#### fft2

2-d Fast Fourier Transform.

## **Syntax**

```
F = fft2(f)
F = fft2(f, size)
F = fft2(f, nr, nc)
F = fft2(f, n)
```

# **Description**

fft2(f) returns the 2-d Discrete Fourier Transform (DFT along dimensions 1 and 2) of array f.

With two or three input arguments, fft2 resizes the two first dimensions by cropping or by padding with zeros. fft2(f,nr,nc) resizes first dimension to nr rows and second dimension to nc columns. In ft2(f,size), the new size is given as a two-element vector [nr,nc]. fft2(F,n) is equivalent to fft2(F,n,n).

If the first argument is an array with more than two dimensions, fft2 performs the 2-d DFT along dimensions 1 and 2 separately for each plane along remaining dimensions; fftn performs an DFT along each dimension.

#### See also

```
ifft2, fft, fftn
```

#### fftn

n-dimension Fast Fourier Transform.

## **Syntax**

```
F = fftn(f)
F = fftn(f, size)
```

## **Description**

fftn(f) returns the n-dimension Discrete Fourier Transform of array f (DFT along each dimension of f).

With two input arguments, fftn(f,size) resizes f by cropping or by padding f with zeros.

#### See also

```
ifftn, fft, fft2
```

#### filter

Digital filtering of data.

## **Syntax**

```
y = filter(b,a,u)
y = filter(b,a,u,x0)
y = filter(b,a,u,x0,dim)
(y, xf) = filter(...)
```

# **Description**

filter(b,a,u) filters vector u with the digital filter whose coefficients are given by polynomials b and a. The filtered data can also be an array, filtered along the first non-singleton dimension or along the dimension specified with a fifth input argument. The fourth argument, if provided and different than the empty matrix [], is a matrix whose columns contain the initial state of the filter and have  $\max(length(a), length(b))-1$  element. Each column correspond to a signal along the dimension of filtering. The result y, which has the

same size as the input, can be computed with the following code if u is a vector:

```
b = b / a(1);
a = a / a(1);
if length(a) > length(b)
  b = [b, zeros(1, length(a)-length(b))];
else
  a = [a, zeros(1, length(b)-length(a))];
end
n = length(x);
for i = 1:length(u)
  y(i) = b(1) * u(i) + x(1);
  for j = 1:n-1
     x(j) = b(j + 1) * u(i) + x(j + 1) - a(j + 1) * y(i);
  end
  x(n) = b(n + 1) * u(i) - a(n + 1) * y(i);
end
```

The optional second output argument is set to the final state of the filter.

## **Examples**

```
filter([1,2], [1,2,3], ones(1,10))
   1 1 -2 4 1 -11 22 -8 -47 121

u = [5,6,5,6,5,6,5];
p = 0.8;
filter(1-p, [1,-p], u, p*u(1))
   % low-pass with matching initial state
5 5.2 5.16 5.328 5.2624 5.4099 5.3279
```

#### See also

conv, deconv, conv2

#### funm

Matrix function.

```
Y = funm(X, fun)
(Y, err) = funm(X, fun)
```

funm(X, fun) returns the matrix function of square matrix X specified by function fun. fun takes a scalar input argument and gives a scalar output. It is either specified by its name or given as an anonymous or inline function or a function reference.

With a second output argument err, funm also returns an estimate of the relative error.

### **Examples**

```
funm([1,2;3,4], @sin)
  -0.4656   -0.1484
  -0.2226   -0.6882
X = [1,2;3,4];
funm(X, @(x) (1+x)/(2-x))
  -0.25   -0.75
  -1.125   -1.375
(eye(2)+X)/(2*eye(2)-X)
  -0.25   -0.75
  -1.125   -1.375
```

#### See also

expm, loam, sartm, schur

### householder

Householder transform.

### **Syntax**

```
(nu, beta) = householder(x)
```

## **Description**

The householder transform is an orthogonal matrix transform which sets all the elements of a column to zero, except the first one. It is the elementary step used by QR decomposition.

The matrix transform can be written as a product by an orthogonal square matrix P=I-beta\*nu\*nu', where I is the identity matrix, beta is a scalar, and nu is a column vector where nu(1) is 1. householder(x), where x is a real or complex non-empty column vector, gives nu and beta such that P\*x=[y;Z], where y is a scalar and Z a zero column vector.

## **Example**

```
x = [2; 5; 10];
(nu, beta) = householder(x)
  nu =
    1.0000
 0.3743
 0.7486
  beta =
    1.1761
P = eve(3) - beta * nu * nu'
  P =
    -0.1761 -0.4402 -0.8805
    -0.4402 0.8352 -0.3296
    -0.8805 -0.3296 0.3409
P * x
  ans =
   -11.3578
     0.0000
     0.0000
```

It is more efficient to avoid calculating P explicitly. Multiplication by P, either as P\*A (to set elements to zero) or B\*P' (to accumulate transforms), can be performed by passing nu and beta to householderapply:

```
householderapply(x, nu, beta)
ans =
-11.3578
0.0000
0.0000
```

#### See also

householderapply, gr

# householderapply

Apply Householder transform.

## **Syntax**

```
B = householderapply(A, nu, beta)
B = householderapply(A, nu, beta, 'r')
```

# **Description**

householderapply(A,nu,beta) apply the Householder transform defined by column vector nu (where nu(1) is 1) and real scalar

beta, as obtained by householder, to matrix A; i.e. it computes A-nu\*beta\*nu'\*A.

householderapply(A,nu,beta,'r') apply the inverse Householder transform to matrix A; i.e. it computes A-A\*nu\*beta\*nu'.

#### See also

householder

### ifft

Inverse Fast Fourier Transform.

## **Syntax**

```
f = ifft(F)
f = ifft(F, n)
f = ifft(F, n, dim)
```

## **Description**

ifft returns the inverse Discrete Fourier Transform (inverse DFT). Up to the sign and a scaling factor, the inverse DFT and the DFT are the same operation: for a vector, ifft(d) = conj(fft(d))/length(d). ifft has the same syntax as fft.

## **Examples**

```
F = fft([1,2,3,4])
F =
10 -2+2j -2 -2-2j
ifft(F)
1 2 3 4
```

#### See also

```
fft, ifft2, ifftn
```

### ifft2

Inverse 2-d Fast Fourier Transform.

## **Syntax**

```
f = ifft2(F)
f = ifft2(F, size)
f = ifft2(F, nr, nc)
f = ifft2(F, n)
```

## **Description**

ifft2 returns the inverse 2-d Discrete Fourier Transform (inverse DFT along dimensions 1 and 2).

With two or three input arguments, ifft2 resizes the two first dimensions by cropping or by padding with zeros. ifft2(F,nr,nc) resizes first dimension to nr rows and second dimension to nc columns. In ifft2(F,size), the new size is given as a two-element vector [nr,nc]. ifft2(F,n) is equivalent to ifft2(F,n,n).

If the first argument is an array with more than two dimensions, ifft2 performs the inverse 2-d DFT along dimensions 1 and 2 separately for each plane along remaining dimensions; ifftn performs an inverse DFT along each dimension.

Up to the sign and a scaling factor, the inverse 2-d DFT and the 2-d DFT are the same operation. ifft2 has the same syntax as fft2.

#### See also

```
fft2, ifft, ifftn
```

### ifftn

Inverse n-dimension Fast Fourier Transform.

## **Syntax**

```
f = ifftn(F)
f = ifftn(F, size)
```

## **Description**

ifftn(F) returns the inverse n-dimension Discrete Fourier Transform of array F (inverse DFT along each dimension of F).

With two input arguments, ifftn(F,size) resizes F by cropping or by padding F with zeros.

Up to the sign and a scaling factor, the inverse n-dimension DFT and the n-dimension DFT are the same operation. ifftn has the same syntax as fftn.

```
fftn, ifft, ifft2
```

### hess

Hessenberg reduction.

## **Syntax**

```
(P,H) = hess(A)
H = hess(A)
```

## **Description**

hess (A) reduces the square matrix A A to the upper Hessenberg form H using an orthogonal similarity transformation P\*H\*P'=A. The result H is zero below the first subdiagonal and has the same eigenvalues as A.

## **Example**

```
(P,H)=hess([1,2,3;4,5,6;7,8,9])
  1
          0
                  0
  0
         -0.4961 -0.8682
         -0.8682 0.4961
 0
 H =
         -3.597 -0.2481
  1
  -8.0623 14.0462 2.8308
         0.8308 -4.6154e-2
P*H*P'
ans =
 1
          2
                  3
  4
          5
                  6
  7
                  9
```

#### See also

lu, qr, schur

### inv

Inverse of a square matrix.

## **Syntax**

```
M2 = inv(M1)
```

## **Description**

inv(M1) returns the inverse M2 of the square matrix M1, i.e. a matrix of the same size such that M2\*M1 = M1\*M2 = eye(size(M1)). M1 must not be singular; otherwise, its elements are infinite.

To solve a set of linear of equations, the operator \ is more efficient.

## **Example**

```
inv([1,2;3,4])
-2 1
1.5 -0.5
```

#### See also

operator /, operator \, pinv, lu, rank, eye

### kron

Kronecker product.

### **Syntax**

```
M = kron(A, B)
```

## **Description**

kron(A,B) returns the Kronecker product of matrices A (size m1 by n1) and B (size m2 by n2), i.e. an m1\*m2-by-n1\*n2 matrix made of m1 by n1 submatrices which are the products of each element of A with B.

## **Example**

```
kron([1,2;3,4],ones(2))
1 1 2 2
1 1 2 2
3 3 4 4
3 3 4 4
```

repmat

### **kurtosis**

Kurtosis of a set of values.

## **Syntax**

```
k = kurtosis(A)
k = kurtosis(A, dim)
```

## **Description**

kurtosis(A) gives the kurtosis of the columns of array A or of the row vector A. The dimension along which kurtosis proceeds may be specified with a second argument.

The kurtosis measures how much values are far away from the mean. It is 3 for a normal distribution, and positive for a distribution which has more values far away from the mean.

## **Example**

```
kurtosis(rand(1, 10000))
   1.8055
```

#### See also

mean, var, skewness, moment

# linprog

Linear programming.

```
x = linprog(c, A, b)
x = linprog(c, A, b, xlb, xub)
```

linprog(c,A,b) solves the following linear programming problem:

$$min cx$$
  
s.t.  $Ax \le b$ 

The optimum x is either finite, infinite if there is no bounded solution, or not a number if there is no feasible solution.

Additional arguments may be used to constrain x between lower and upper bounds. linprog(c,A,b,xlb,xub) solves the following linear programming problem:

$$min c x$$
s.t.  $Ax \le b$ 

$$x \ge x_{lb}$$

$$x \le x_{ub}$$

If xub is missing, there is no upper bound. xlb and xub may have less elements than x, or contain -inf or +inf; corresponding elements have no lower and/or upper bounds.

## **Examples**

Maximize 3x + 2y subject to  $x + y \le 9$ ,  $3x + y \le 18$ ,  $x \le 7$ , and  $y \le 6$ :

```
c = [-3,-2];
A = [1,1; 3,1; 1,0; 0,1];
b = [9; 18; 7; 6];
x = linprog(c, A, b)
x =
    4.5
    4.5
```

A more efficient way to solve the problem, with bounds on variables:

```
c = [-3,-2];
A = [1,1; 3,1];
b = [9; 18];
xlb = [];
xub = [7; 6];
x = linprog(c, A, b, xlb, xub)
x =
4.5
4.5
```

Check that the solution is feasible and bounded:

```
all(isfinite(x))
  true
```

# logm

Matrix logarithm.

## **Syntax**

```
Y = logm(X)
(Y, err) = logm(X)
```

## **Description**

logm(X) returns the matrix logarithm of X, the inverse of the matrix exponential. X must be square. The matrix logarithm does not always exist.

With a second output argument err, logm also returns an estimate of the relative error norm(expm(logm(X))-X)/norm(X).

## **Example**

```
Y = logm([1,2;3,4])

Y =

-0.3504 + 2.3911j  0.9294 - 1.0938j

1.394 - 1.6406j  1.0436 + 0.7505j

expm(Y)

1 - 5.5511e-16j  2 -7.7716e-16j

3 - 8.3267e-16j  4
```

#### See also

expm, sqrtm, funm, schur, log

### lu

LU decomposition.

### **Syntax**

```
(L, U, P) = lu(A)
(L2, U) = lu(A)
Y = lu(A)
```

# **Description**

With three output arguments, lu(A) computes the LU decomposition of matrix A with partial pivoting, i.e. a lower triangular matrix L, an upper triangular matrix U, and a permutation matrix P such that P\*A=L\*U.

If A in an m-by-n mytrix, L is m-by-min(m,n), U is min(m,n)-by-n and P is m-by-m. A can be rank-deficient.

With two output arguments, lu(A) permutes the lower triangular matrix and gives L2=P'\*L, such that A=L2\*U.

With a single output argument, lu gives Y=L+U-eye(n).

### **Example**

```
X = [1,2,3;4,5,6;7,8,8];
(L,U,P) = lu(X)
I =
1
      0
            0
0.143 1
            0
0.571 0.5 1
U =
7
      8
     0.857 1.857
0
     0 0.5
P =
0 0 1
1 0 0
0 1 0
P*X-L*U
ans =
0 0 0
0 0 0
0 0 0
```

#### See also

inv, qr, svd

# lyap

Continuous-time Lyapunov equation.

## **Syntax**

```
X = lyap(A, B, C)

X = lyap(A, C)
```

# **Description**

lyap(A,B,C) calculates the solution X of the following continuous-time Lyapunov equation:

$$AX + XB + C = 0$$

All matrices are real.

With two input arguments, lyap(A,C) solves the following Lyapunov equation:

$$AX + XA' + C = 0$$

### **Example**

```
A = [3,1,2;1,3,5;6,2,1];

B = [2,7;8,3];

C = [2,1;4,5;8,9];

X = lyap(A, B, C)

X =

0.1635 -0.1244

-0.2628 0.1311

-0.7797 -0.7645
```

#### See also

dlyap, care

#### max

Maximum value of a vector or of two arguments.

## **Syntax**

```
x = max(v)
(v,ind) = max(v)
v = max(M,[],dim)
(v,ind) = max(M,[],dim)
M3 = max(M1,M2)
```

## **Description**

max(v) returns the largest number of vector v. NaN's are ignored. The optional second output argument is the index of the maximum in v; if several elements have the same maximum value, only the first one is obtained. The argument type can be double, single, or integer of any size.

max(M) operates on the columns of the matrix M and returns a row vector. max(M,[],dim) operates along dimension dim (1 for columns, 2 for rows).

max(M1,M2) returns a matrix whose elements are the maximum between the corresponding elements of the matrices M1 and M2. M1 and M2 must have the same size, or be a scalar which can be compared against any matrix.

## **Examples**

```
(mx,ix) = max([1,3,2,5,8,7])
  mx =
    8
  ix =
    5
max([1,3;5,nan], [], 2)
    3
    5
max([1,3;5,nan], 2)
    2 3
    5 2
```

### See also

min

#### mean

Arithmetic mean of a vector.

## **Syntax**

```
x = mean(v)
v = mean(M)
v = mean(M,dim)
```

# **Description**

mean(v) returns the arithmetic mean of the elements of vector v. mean(M) returns a row vector whose elements are the means of the corresponding columns of matrix M. mean(M,dim) returns the mean of matrix M along dimension dim; the result is a row vector if dim is 1, or a column vector if dim is 2.

# **Examples**

```
mean(1:5)
  7.5
mean((1:5)')
  7.5
mean([1,2,3;5,6,7])
  3 4 5
mean([1,2,3;5,6,7],1)
  3 4 5
mean([1,2,3;5,6,7],2)
```

2

#### See also

```
cov, std, var, median, sum, prod
```

### median

Median.

### **Syntax**

```
x = median(v)
v = median(M)
v = median(M, dim)
```

## **Description**

median(v) gives the median of vector v, i.e. the value x such that half of the elements of v are smaller and half of the elements are larger. The result is NaN if any value is NaN.

median(M) gives a row vector which contains the median of the columns of M. With a second argument, median(M, dim) operates along dimension dim.

## **Example**

```
median([1, 2, 5, 6, inf])
5
```

#### See also

mean, sort

### min

Minimum value of a vector or of two arguments.

```
x = min(v)
(v,ind) = min(v)
v = min(M,[],dim)
(v,ind) = min(M,[],dim)
M3 = min(M1,M2)
```

min(v) returns the largest number of vector v. NaN's are ignored. The optional second smallest argument is the index of the minimum in v; if several elements have the same minimum value, only the first one is obtained. The argument type can be double, single, or integer of any size.

min(M) operates on the columns of the matrix M and returns a row vector. min(M,[],dim) operates along dimension dim(1 for columns, 2 for rows).

min(M1,M2) returns a matrix whose elements are the minimum between the corresponding elements of the matrices M1 and M2. M1 and M2 must have the same size, or be a scalar which can be compared against any matrix.

## **Examples**

#### See also

max

### moment

Central moment of a set of values.

# Syntax

```
m = moment(A, order)
m = moment(A, order, dim)
```

# **Description**

moment(A,order) gives the central moment (moment about the mean) of the specified order of the columns of array A or of the row

vector A. The dimension along which moment proceeds may be specified with a third argument.

### **Example**

```
moment(randn(1, 10000), 3) 3.011
```

### See also

mean, var, skewness, kurtosis

#### norm

Norm of a vector or matrix.

### **Syntax**

```
x = norm(v)
x = norm(v,kind)
x = norm(M)
x = norm(M,kind)
```

# Description

With one argument, norm calculates the 2-norm of a vector or the induced 2-norm of a matrix. The optional second argument specifies the kind of norm.

Kind	Vector	Matrix
none or 2	<pre>sqrt(sum(abs(v).^2))</pre>	largest singular value
		(induced 2-norm)
1	sum(abs(V))	largest column sum of abs
<pre>inf or 'inf'</pre>	max(abs(v))	largest row sum of abs
-inf	min(abs(v))	invalid
p	sum(abs(V).^p)^(1/p)	invalid
'fro'	<pre>sqrt(sum(abs(v).^2))</pre>	sqrt(sum(diag(M'*M)))

## **Examples**

```
norm([3,4])
5
norm([2,5;9,3])
10.2194
norm([2,5;9,3],1)
11
```

```
abs, hypot, svd
```

### null

Null space.

### **Syntax**

```
Z = null(A)
Z = null(A, tol=tol)
```

## **Description**

null(A) returns a matrix Z whose columns are an orthonormal basis for the null space of m-by-n matrix A. Z has n-rank(A) columns, which are the last right singular values of A; that is, those corresponding to the singular values below a small tolerance. This tolerance can be specified with a named argument tol.

Without input argument, null gives the null value (the unique value of the special null type, not related to linear algebra).

## **Example**

```
null([1,2,3;1,2,4;1,2,5])
-0.8944
0.4472
8.0581e-17
```

#### See also

```
svd, orth, null (null value)
```

## orth

Orthogonalization.

```
Q = orth(A)
Q = orth(A, tol=tol)
```

orth(A) returns a matrix Q whose columns are an orthonormal basis for the range of those of matrix A. Q has rank(A) columns, which are the first left singular vectors of A (that is, those corresponding to the largest singular values).

Orthogonalization is based on the singular-value decomposition, where only the singular values larger than some small threshold are considered. This threshold can be specified with an optional named argument.

## **Example**

```
orth([1,2,3;1,2,4;1,2,5])
-0.4609 0.788
-0.5704 8.9369e-2
-0.6798 -0.6092
```

#### See also

svd. null

# pinv

Pseudo-inverse of a matrix.

## **Syntax**

```
M2 = pinv(M1)
M2 = pinv(M1, tol)
M2 = pinv(M1, tol=tol)
```

# **Description**

pinv(M1) returns the pseudo-inverse of matrix M. For a nonsingular square matrix, the pseudo-inverse is the same as the inverse. For an arbitrary matrix (possibly nonsquare), the pseudo-inverse M2 has the following properties: size(M2) = size(M1'), M1\*M2\*M1 = M1, M2\*M1\*M2 = M2, and the norm of M2 is minimum. The pseudo-inverse is based on the singular-value decomposition, where only the singular values larger than some small threshold are considered. This threshold can be specified with an optional second argument tol or as a named argument.

If M1 is a full-rank matrix with more rows than columns, pinv returns the least-square solution  $pinv(M1)*y = (M1'*M1)\M1'*y$  of the overdetermined system M1\*x = y.

### **Examples**

### See also

inv. svd

# poly

Characteristic polynomial of a square matrix or polynomial coefficients based on its roots.

## **Syntax**

```
pol = poly(M)
pol = poly(r)
```

# **Description**

With a matrix argument, poly(M) returns the characteristic polynomial det(x\*eye(size(M))-M) of the square matrix M. The roots of the characteristic polynomial are the eigenvalues of M.

With a vector argument, poly(r) returns the polynomial whose roots are the elements of the vector r. The first coefficient of the polynomial is 1. If the complex roots form conjugate pairs, the result is real.

## **Examples**

```
poly([1,2;3,4]
   1 -5 -2
roots(poly([1,2;3,4]))
   5.3723
   -0.3723
eig([1,2;3,4])
   -0.3723
```

```
5.3723
poly(1:3)
1 -6 11 -6
```

roots. det

# polyder

Derivative of a polynomial or a polynomial product or ratio.

## **Syntax**

```
A1 = polyder(A)
C1 = polyder(A, B)
(N1, D1) = polyder(N, D)
```

## **Description**

polyder (A) returns the polynomial which is the derivative of the polynomial A. Both polynomials are given as vectors of coefficients, highest power first. The result is a row vector.

With a single output argument, polyder(A,B) returns the derivative of the product of polynomials A and B. It is equivalent to polyder(conv(A,B)).

With two output arguments, (N1,D1)=polyder(N,D) returns the derivative of the polynomial ratio N/D as N1/D1. Input and output arguments are polynomial coefficients.

## **Examples**

```
Derivative of x^3 + 2x^2 + 5x + 2:

polyder([1, 2, 5, 2])
    3 4 5

Derivative of (x^3 + 2x^2 + 5x + 2)/(2x + 3):

(N, D) = polyder([1, 2, 5, 2], [2, 3])
    N =
    4 13 12 11
    D =
    4 12 9
```

polyint, polyval, poly, addpol, conv

# polyint

Integral of a polynomial.

## **Syntax**

```
pol2 = polyint(pol1)
pol2 = polyint(pol1, c)
```

## **Description**

polyint(pol1) returns the polynomial which is the integral of the polynomial pol1, whose zero-order coefficient is 0. Both polynomials are given as vectors of coefficients, highest power first. The result is a row vector. A second input argument can be used to specify the integration constant.

## **Example**

```
Y = polyint([1, 2, 3, 4, 5])
 Y =
   0.2
         0.5
               1
                2 5
                               0
y = polyder(Y)
 y =
   1
         2
               3
Y = polyint([1, 2, 3, 4, 5], 10)
              1
                    2
                          5
   0.2
         0.5
                                10
```

#### See also

polyder, polyval, poly, addpol, conv

# polyval

Numeric value of a polynomial evaluated at some point.

```
y = polyval(pol, x)
```

polyval(pol,x) evaluates the polynomial pol at x, which can be a scalar or a matrix of arbitrary size. The polynomial is given as a vector of coefficients, highest power first. The result has the same size as x.

### **Examples**

```
polyval([1,3,8], 2)
   18
polyval([1,2], 1:5)
   3 4 5 6 7
```

#### See also

```
polyder, polyint, poly, addpol, conv
```

## prod

Product of the elements of a vector.

## **Syntax**

```
x = prod(v)
v = prod(M)
v = prod(M,dim)
```

## **Description**

prod(v) returns the product of the elements of vector v. prod(M) returns a row vector whose elements are the products of the corresponding columns of matrix M. prod(M,dim) returns the product of matrix M along dimension dim; the result is a row vector if dim is 1, or a column vector if dim is 2.

## **Examples**

```
prod(1:5)
    120
prod((1:5)')
    120
prod([1,2,3;5,6,7])
    5 12 21
prod([1,2,3;5,6,7],1)
    5 12 21
prod([1,2,3;5,6,7],2)
```

```
6
210
```

```
sum, mean, operator *
```

### qr

QR decomposition.

## **Syntax**

```
(Q, R, E) = qr(A)
(Q, R) = qr(A)
R = qr(A)
(Qe, Re, e) = qr(A, false)
(Qe, Re) = qr(A, false)
Re = qr(A, false)
```

## **Description**

With three output arguments, qr(A) computes the QR decomposition of matrix A with column pivoting, i.e. a square unitary matrix Q and an upper triangular matrix R such that A\*E=Q\*R. With two output arguments, qr(A) computes the QR decomposition without pivoting, such that A=Q\*R. With a single output argument, qr gives R.

With a second input argument with the value false, if A has m rows and n columns with m>n, qr produces an m-by-n Q and an n-by-n R. Bottom rows of zeros of R, and the corresponding columns of Q, are discarded. With column pivoting, the third output argument e is a permutation vector: A(:,e)=Q\*R.

## **Example**

```
(Q,R) = qr([1,2;3,4;5,6])
0 =
             0.8971
                      0.4082
   -0.169
             0.276
                     -0.8165
   -0.5071
   -0.8452
           -0.345
                     0.4082
R =
   -5.9161
           -7.4374
             0.8281
        0
        0
                  0
(Q,R) = qr([1,2;3,4;5,6],false)
   0.169
             0.8971
```

```
0.5071 0.276
0.8452 -0.345
R =
5.9161 7.4374
0 0.8281
```

lu, schur, hess, svd

### rank

Rank of a matrix.

## **Syntax**

```
x = rank(M)
x = rank(M, tol)
x = rank(M, tol=tol)
```

## **Description**

rank(M) returns the rank of matrix M, i.e. the number of lines or columns linearly independent. To obtain it, the singular values are computed and the number of values significantly larger than 0 is counted. The value below which they are considered to be 0 can be specified with the optional second argument or named argument.

## **Examples**

```
rank([1,1;0,0])
1
rank([1,1;0,1j])
2
```

#### See also

svd, cond, pinv, det

## roots

Roots of a polynomial.

## **Syntax**

```
r = roots(pol)
r = roots(M)
r = roots(M,dim)
```

## **Description**

roots(pol) calculates the roots of the polynomial pol. The polynomial is given by the vector of its coefficients, highest power first, while the result is a column vector.

With a matrix as argument, roots (M) calculates the roots of the polynomials corresponding to each column of M. An optional second argument is used to specify in which dimension roots operates (1 for columns, 2 for rows). The roots of the i:th polynomial are in the i:th column of the result, whatever the value of dim is.

## **Examples**

```
roots([1, 0, -1])
1
-1
roots([1, 0, -1]')
1
-1
roots([1, 1; 0, 5; -1, 6])
1 -2
-1 -3
roots([1, 0, -1]', 2)
[]
```

#### See also

```
poly, eig
```

#### schur

Schur factorization.

```
(U,T) = schur(A)
T = schur(A)
(U,T) = schur(A, 'c')
T = schur(A, 'c')
```

schur(A) computes the Schur factorization of square matrix A, i.e. a unitary matrix U and a square matrix T (the *Schur matrix*) such that A=U\*T\*U'. If A is complex, the Schur matrix is upper triangular, and its diagonal contains the eigenvalues of A; if A is real, the Schur matrix is real upper triangular, except that there may be 2-by-2 blocks on the main diagonal which correspond to the complex eigenvalues of A. To force a complex Schur factorization with an upper triangular matrix T, schur is given a second input argument 'c' or 'complex'.

### **Examples**

Schur factorization:

```
A = [1,2;3,4];
(U,T) = schur(A)
U =
-0.8246 -0.5658
0.5658 -0.8246
T =
-0.3723 -1
0 5.3723
```

Since T is upper triangular, its diagonal contains the eigenvalues of A:

```
eig(A)
ans =
-0.3723
5.3723
```

For a matrix with complex eigenvalues, the real Schur factorization has 2x2 blocks on its diagonal:

```
T = schur([1,0,0;0,1,2;0,-3,1])
  T =
                   0
     1
            0
            1
                   2
     0
           - 3
                   1
T = schur([1,0,0;0,1,2;0,-3,1],'c')
  T =
     1
                     0
                                     0
     0
                     1 + 2.4495j
                                     1
     0
                     0
                                     1 - 2.4495i
```

#### See also

lu, hess, qr, eig

### skewness

Skewness of a set of values.

## **Syntax**

```
s = skewness(A)
s = skewness(A, dim)
```

## **Description**

skewness (A) gives the skewness of the columns of array A or of the row vector A. The dimension along which skewness proceeds may be specified with a second argument.

The skewness measures how asymmetric a distribution is. It is 0 for a symmetric distribution, and positive for a distribution which has more values much larger than the mean.

## **Example**

```
skewness(randn(1, 10000).^2)
2.6833
```

#### See also

mean, var, kurtosis, moment

# sqrtm

Matrix square root.

## **Syntax**

```
Y = sqrtm(X)
(Y, err) = sqrtm(X)
```

# **Description**

sqrtm(X) returns the matrix square root of X, such that  $sqrtm(X)^2=X$ . X must be square. The matrix square root does not always exist.

With a second output argument err, sqrtm also returns an estimate of the relative error  $norm(sqrtm(X)^2-X)/norm(X)$ .

### **Example**

```
Y = sqrtm([1,2;3,4])

Y =

0.5537 + 0.4644j  0.807 - 0.2124j

1.2104 - 0.3186j  1.7641 + 0.1458j

Y^2

1  2

3  4
```

#### See also

expm, logm, funm, schur, chol, sgrt

#### std

Standard deviation.

## **Syntax**

```
x = std(v)
x = std(v, p)
v = std(M)
v = std(M, p)
v = std(M, p, dim)
```

## **Description**

std(v) gives the standard deviation of vector v, normalized by length(v)-1. With a second argument, std(v,p) normalizes by length(v)-1 if p is false, or by length(v) if p is true.

std(M) gives a row vector which contains the standard deviation of the columns of M. With a third argument, std(M,p,dim) operates along dimension dim.

## **Example**

```
std([1, 2, 5, 6, 10, 12])
4.3359
```

### See also

mean, var, cov

#### sum

Sum of the elements of a vector.

## **Syntax**

```
x = sum(v)
v = sum(M)
v = sum(M,dim)
```

## **Description**

sum(v) returns the sum of the elements of vector v. sum(M) returns a row vector whose elements are the sums of the corresponding columns of matrix M. sum(M,dim) returns the sum of matrix M along dimension dim; the result is a row vector if dim is 1, or a column vector if dim is 2.

## **Examples**

```
sum(1:5)
  15
sum((1:5)')
  15
sum([1,2,3;5,6,7])
  6 8 10
sum([1,2,3;5,6,7],1)
  6 8 10
sum([1,2,3;5,6,7],2)
  6
  18
```

#### See also

```
prod, mean, operator +
```

#### svd

Singular value decomposition.

```
s = svd(M)
(U,S,V) = svd(M)
(U,S,V) = svd(M,false)
```

The singular value decomposition (U,S,V) = svd(M) decomposes the m-by-n matrix M such that M = U\*S\*V', where S is an m-by-n diagonal matrix with decreasing positive diagonal elements (the singular values of M), U is an m-by-m unitary matrix, and V is an n-by-n unitary matrix. The number of non-zero diagonal elements of S (up to rounding errors) gives the rank of M.

When M is rectangular, in expression U\*S\*V', some columns of U or V are multiplied by rows or columns of zeros in S, respectively. (U,S,V) = svd(M,false) produces U, S and V where these columns or rows are discarded (relationship M = U\*S\*V' still holds):

```
Size of A Size of U Size of S Size of V m \text{ by } n, m <= n  m \text{ by } m m \text{ by } m n \text{ by } m n \text{ by } m n \text{ by } n
```

svd(M,true) produces the same result as svd(M).

With one output argument, s = svd(M) returns the vector of singular values s=diag(S).

The singular values of M can also be computed with s = sqrt(eig(M'\*M)), but svd is faster and more robust.

## **Examples**

```
(U,S,V)=svd([1,2;3,4])

U =

0.4046 0.9145

0.9145 -0.4046

S =

5.465 0

0 0.366

V =

0.576 -0.8174

0.8174 0.576

U*S*V'

1 2

3 4

svd([1,2;1,2])

3.1623

3.4697e-19
```

#### See also

eig, pinv, rank, cond, norm

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#### trace

Trace of a matrix.

## **Syntax**

```
tr = trace(M)
```

## **Description**

trace(M) returns the trace of the matrix M, i.e. the sum of its diagonal elements.

## **Example**

```
trace([1,2;3,4])
5
```

#### See also

norm, diag

#### var

Variance of a set of values.

## **Syntax**

```
s2 = var(A)
s2 = var(A, p)
s2 = var(A, p, dim)
```

## **Description**

var(A) gives the variance of the columns of array A or of the row vector A. The variance is normalized with the number of observations minus 1, or by the number of observations if a second argument is true. The dimension along which var proceeds may be specified with a third argument.

#### See also

mean, std, cov, kurtosis, skewness, moment

# **6.19 Array Functions**

# arrayfun

Function evaluation for each element of an array.

## **Syntax**

```
(B1, \ldots) = arrayfun(fun, A1, \ldots)
```

## **Description**

arrayfun(fun,A) evaluates function fun for each element of numeric array A. Each evaluation must give a scalar result of numeric (or logical or char) type; results are returned as a numeric array the same size as A. First argument is a function reference, an inline function, or the name of a function as a string.

With more than two input arguments, arrayfun calls function fun as feval(fun,A1(i),A2(i),...). All array arguments must have the same size, but their type can be different.

With two output arguments or more, arrayfun evaluates function fun with the same number of output arguments and builds a separate array for each output. Without output argument, arrayfun evaluates fun without output argument.

arrayfun differs from cellfun: all input arguments of arrayfun are arrays of any type (not necessarily cell arrays), and corresponding elements are provided provided to fun. With map, input arguments as well as output arguments are cell arrays.

## **Examples**

```
arrayfun(@isempty, {1, ''; {}, ones(5)})
  F T
  T F
map(@isempty, {1, ''; {}, ones(5)})
  2x2 cell array
(m, n) = arrayfun(@size, {1, ''; {}, ones(2, 5)})
  m =
    1  0
    0  2
  n =
    1  0
    0  5
```

#### See also

cellfun, map, fevalx

### cat

Array concatenation.

## **Syntax**

```
cat(dim, A1, A2, ...)
```

## **Description**

cat(dim,A1,A2,...) concatenates arrays A1, A2, etc. along dimension dim. Other dimensions must match. cat is a generalization of the comma and the semicolon inside brackets.

## **Examples**

```
cat(2, [1,2;3,4], [5,6;7,8])
1  2  5  6
3  4  7  8
cat(3, [1,2;3,4], [5,6;7,8])
2x2x2 array
  (:,:,1) =
    1  2
    3  4
  (:,:,2) =
    5  6
    7  8
```

### See also

```
operator [], operator;, operator,
```

### cell

Cell array of empty arrays.

```
C = cell(n)
C = cell(n1,n2,...)
C = cell([n1,n2,...])
```

cell builds a cell array whose elements are empty arrays []. The size of the cell array is specified by one integer for a square array, or several integers (either as separate arguments or in a vector) for a cell array of any size.

### **Example**

```
cell(2, 3)
2x3 cell array
```

#### See also

```
zeros, operator {}, iscell
```

### cellfun

Function evaluation for each cell of a cell array.

## **Syntax**

```
A = cellfun(fun, C)
A = cellfun(fun, C, ...)
A = cellfun(fun, S)
A = cellfun(fun, S, ...)
```

# Description

cellfun(fun,C) evaluates function fun for each cell of cell array C. Each evaluation must give a scalar result of numeric, logical, or character type; results are returned as a non-cell array the same size as C. First argument is a function reference, an inline function, or the name of a function as a string.

With more than two input arguments, cellfun calls function fun as feval(fun,  $C\{i\}$ , other), where  $C\{i\}$  is each cell of C in turn, and other stands for the remaining arguments of cellfun.

The second argument can be a structure array S instead of a cell array. In that case, fun is called with S(i).

cellfun differs from map in two ways: the result is a non-cell array, and remaining arguments of cellfun are provided directly to fun.

## **Examples**

```
cellfun(@isempty, {1, ''; {}, ones(5)})
  F T
  T F
map(@isempty, {1, ''; {}, ones(5)})
  2x2 cell array
cellfun(@size, {1, ''; {}, ones(5)}, 2)
  1 0
  0 5
```

### See also

map, arrayfun

# diag

Creation of a diagonal matrix or extraction of the diagonal elements of a matrix.

## **Syntax**

```
M = diag(v)
M = diag(v,k)
v = diag(M)
v = diag(M,k)
```

# **Description**

With a vector input argument, diag(v) creates a square diagonal matrix whose main diagonal is given by v. With a second argument, the diagonal is moved by that amount in the upper right direction for positive values, and in the lower left direction for negative values.

With a matrix input argument, the main diagonal is extracted and returned as a column vector. A second argument can be used to specify another diagonal.

## **Examples**

```
diag(1:3)
    1 0 0
    0 2 0
    0 3
diag(1:3,1)
    0 1 0 0
    0 2 0
    0 0 3
```

```
0 0 0 0

M = magic(3)

M =

8 1 6

3 5 7

4 9 2

diag(M)

8

5

2

diag(M,1)

1
```

tril, triu, eye, trace

## eye

Identity matrix.

## **Syntax**

```
M = eye(n)
M = eye(m,n)
M = eye([m,n])
M = eye(..., type)
```

## **Description**

eye builds a matrix whose diagonal elements are 1 and other elements 0. The size of the matrix is specified by one integer for a square matrix, or two integers (either as two arguments or in a vector of two elements) for a rectangular matrix.

An additional input argument can be used to specify the type of the result. It must be the string 'double', 'single', 'int8', 'int16', 'int32', 'int64', 'uint8', 'uint16', 'uint32', or 'uint64' (64-bit arrays are not supported on all platforms).

## **Examples**

```
eye(3)

1 0 0

0 1 0

0 0 1

eye(2, 3)
```

```
1 0 0
0 1 0
eye(2, 'int8')
2x2 int8 array
1 0
0 1
```

ones, zeros, diag

## fevalx

Function evaluation with array expansion.

## **Syntax**

```
(Y1,...) = fevalx(fun,X1,...)
```

## **Description**

(Y1, Y2,...)=fevalx(fun, X1, X2,...) evaluates function fun with input arguments X1, X2, etc. Arguments must be arrays, which are expanded if necessary along singleton dimensions so that all dimensions match. For instance, three arguments of size 3x1x2, 1x5 and 1x1 are replicated into arrays of size 3x5x2. Output arguments are assigned to Y1, Y2, etc. Function fun is specified either by its name as a string, by a function reference, or by an inline or anonymous function.

## **Example**

```
fevalx(@plus, 1:5, (10:10:30)')
   11
         12
               13
                    14
                          15
   21
         22
               23
                    24
                          25
         32
             33
   31
                   34
                          35
```

### See also

feval, meshgrid, repmat, inline, operator @

#### find

Find the indices of the non-null elements of an array.

## **Syntax**

```
ix = find(v)
[s1,s2] = find(M)
[s1,s2,x] = find(M)
... = find(..., n)
... = find(..., n, dir)
```

## **Description**

With one output argument, find(v) returns a vector containing the indices of the nonzero elements of v. v can be an array of any dimension; the indices correspond to the internal storage ordering and can be used to access the elements with a single subscript.

With two output arguments, find(M) returns two vectors containing the subscripts (row in the first output argument, column in the second output argument) of the nonzero elements of 2-dim array M. To obtain subscripts for an array of higher dimension, you can convert the single output argument of find to subscripts with ind2sub.

With three output arguments, find(M) returns in addition the nonzero values themselves in the third output argument.

With a second input argument n, find limits the maximum number of elements found. It searches forward by default; with a third input argument dir, find gives the n first nonzero values if dir is 'first' or 'f', and the n last nonzero values if dir is 'last' or 'l'.

### **Examples**

```
ix = find([1.2,0;0,3.6])
ix =
  1
[s1,s2] = find([1.2,0;0,3.6])
s1 =
  1
s2 =
  1
  2
[s1,s2,x] = find([1.2,0;0,3.6])
  1
  2
s2 =
  1
  2
x =
  1.2
```

```
3.6

A = rand(3)

A =

0.5599 0.3074 0.5275

0.3309 0.8077 0.3666

0.7981 0.6424 0.6023

find(A > 0.7, 2, 'last')

7
```

nnz, sort

# flipdim

Flip an array along any dimension.

## **Syntax**

```
B = flipdim(A, dim)
```

# **Description**

flipdim(A,dim) gives an array which has the same size as A, but where indices of dimension dim are reversed.

# **Examples**

```
flipdim(cat(3, [1,2;3,4], [5,6;7,8]), 3)
    2x2x2 array
    (:,:,1) =
        5  6
        7  8
    (:,:,2) =
        1  2
        3  4
```

#### See also

fliplr, flipud, rot90, reshape

# fliplr

Flip an array or a list around its vertical axis.

## **Syntax**

```
A2 = fliplr(A1)
list2 = fliplr(list1)
```

## **Description**

fliplr(A1) gives an array A2 which has the same size as A1, but where all columns are placed in reverse order.

fliplr(list1) gives a list list2 which has the same length as list1, but where all top-level elements are placed in reverse order. Elements themselves are left unchanged.

## **Examples**

```
fliplr([1,2;3,4])
2 1
4 3
fliplr({1, 'x', {1,2,3}})
{{1,2,3}, 'x', 1}
```

#### See also

flipud, flipdim, rot90, reshape

# flipud

Flip an array upside-down.

# **Syntax**

```
A2 = flipud(A1)
```

# **Description**

flipud(A1) gives an array A2 which has the same size as A1, but where all lines are placed in reverse order.

## **Example**

```
flipud([1,2;3,4])
3 4
1 2
```

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### See also

fliplr, flipdim, rot90, reshape

### ind2sub

Conversion from single index to row/column subscripts.

## **Syntax**

```
(i, j, ...) = ind2sub(size, ind)
```

## **Description**

ind2sub(size,ind) gives the subscripts of the element which would be retrieved from an array whose size is specified by size by the single index ind. size must be either a scalar for square matrices or a vector of two elements or more for arrays. ind can be an array; the result is calculated separately for each element and has the same size.

## **Example**

```
M = [3, 6; 8, 9];
M(3)
8
(i, j) = ind2sub(size(M), 3)
i =
   2
j =
   1
M(i, j)
8
```

#### See also

sub2ind, size

# interp1

1D interpolation.

## **Syntax**

```
yi = interp1(x, y, xi)
yi = interp1(x, y, xi, method)
yi = interp1(y, xi)
yi = interp1(y, xi, method)
yi = interp1(..., method, extraval)
```

## **Description**

interp1(x,y,xi) interpolates data along one dimension. Input data are defined by vector y, where element y(i) corresponds to coordinates x(i). Interpolation is performed at points defined in vector xi; the result is a vector of the same size as xi.

If y is an array, interpolation is performed along dimension 1 (i.e. along its columns), and size(y,1) must be equal to length(x). Then if xi is a vector, interpolation is performed at the same points for each remaining dimensions of y, and the result is an array of size [length(xi), size(y)(2:end)]; if xi is an array, all sizes must match y except for the first one.

If x is missing, it defaults to 1:size(y,1).

The default interpolation method is piecewise linear. An additional input argument can be provided to specify it with a string (only the first character is considered):

Argument	Meaning
'0' or 'nearest'	nearest value
'<'	lower coordinate
'>'	higher coordinate
'1' or 'linear'	piecewise linear
'3' or 'cubic'	piecewise cubic
'p' or 'pchip'	pchip

Cubic interpolation gives continuous values and first derivatives, and null second derivatives at end points. Pchip (piecewise cubic Hermite interpolation) also gives continuous values and first derivatives, but guarantees that the interpolant stays within the limits of the data in each interval (in particular monotonicity is preserved) at the cost of larger second derivatives.

With vectors, interp1 produces the same result as interpn; vector orientations do not have to match, though.

When the method is followed by a scalar number extraval, that value is assigned to all values outside the range defined by x (i.e. extrapolated values). The default is NaN.

# **Examples**

One-dimension interpolation:

```
interp1([1, 2, 5, 8], [0.1, 0.2, 0.5, 1], [0, 2, 3, 7])
  nan  0.2000  0.3000  0.8333
interp1([1, 2, 5, 8], [0.1, 0.2, 0.5, 1], [0, 2, 3, 7], '0')
  nan  0.2000  0.2000  1.0000
```

Interpolation of multiple values:

```
t = 0:10;
y = [sin(t'), cos(t')];
tnew = 0:0.4:8;
ynew = interp1(t, y, tnew)
    ynew =
        0.0000     1.0000
        0.3366     0.8161
        ...
        0.8564     0.2143
        0.9894     -0.1455
```

### See also

interpn

# interpn

Multidimensional interpolation.

## **Syntax**

```
Vi = interpn(x1, ..., xn, V, xi1, ..., xin)
Vi = interpn(x1, ..., xn, V, xi1, ..., xin, method)
Vi = interpn(..., method, extraval)
```

# **Description**

interpn(x1,...,xn,V,xi1,...,xin) interpolates data in a space of n dimensions. Input data are defined by array V, where element V(i,j,...) corresponds to coordinates x1(i), x2(j), etc. Interpolation is performed for each coordinates defined by arrays xi1, xi2, etc., which must all have the same size; the result is an array of the same size.

Length of vectors x1, x2, ... must match the size of V along the corresponding dimension. Vectors x1, x2, ... must be sorted (monotonically increasing or decreasing), but they do not have to be spaced uniformly. Interpolated points outside the input volume are set to nan. Input and output data can be complex. Imaginary parts of coordinates are ignored.

The default interpolation method is multilinear. An additional input argument can be provided to specify it with a string (only the first character is considered):

Argument	Meaning
'0' or 'nearest'	nearest value
'<'	lower coordinates
'>'	higher coordinates
'1' or 'linear'	multilinear

Method '<' takes the sample where each coordinate has its index as large as possible, lower or equal to the interpolated value, and smaller than the last coordinate. Method '>' takes the sample where each coordinate has its index greater or equal to the interpolated value.

When the method is followed by a scalar number extraval, that value is assigned to all values outside the input volume (i.e. extrapolated values). The default is NaN.

### **Examples**

One-dimension interpolation:

```
interpn([1, 2, 5, 8], [0.1, 0.2, 0.5, 1], [0, 2, 3, 7])
  nan   0.2000   0.3000   0.8333
interpn([1, 2, 5, 8], [0.1, 0.2, 0.5, 1], [0, 2, 3, 7], '0')
  nan   0.2000   0.2000   1.0000
```

Three-dimension interpolation:

```
D = cat(3,[0,1;2,3],[4,5;6,7]);
interpn([0,1], [0,1], [0,1], D, 0.2, 0.7, 0.5)
3.1000
```

Image rotation (we define original coordinates between -0.5 and 0.5 in vector c and arrays X and Y, and the image as a linear gradient between 0 and 1):

```
c = -0.5:0.01:0.5;
X = repmat(c, 101, 1);
Y = X';
phi = 0.2;
Xi = cos(phi) * X - sin(phi) * Y;
Yi = sin(phi) * X + cos(phi) * Y;
D = 0.5 + X;
E = interpn(c, c, D, Xi, Yi);
E(isnan(E)) = 0.5;
```

#### See also

interp1

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### intersect

Set intersection.

## **Syntax**

```
c = intersect(a, b)
(c, ia, ib) = intersect(a, b)
```

## **Description**

intersect(a,b) gives the intersection of sets a and b, i.e. it gives the set of members of both sets a and b. Sets are any type of numeric, character or logical arrays, or lists or cell arrays of character strings. Multiple elements of input arguments are considered as single members; the result is always sorted and has unique elements.

The optional second and third output arguments are vectors of indices such that if (c,ia,ib)=intersect(a,b), then c is a(ia) as well as b(ib).

## **Example**

Set exclusive or can also be computed as the union of a and b minus the intersection of a and b:

```
setdiff(union(a, b), intersect(a, b))
{'a','aa','de','z'}
```

### See also

unique, union, setdiff, setxor, ismember

## inthist

Histogram of an integer array.

## **Syntax**

```
h = inthist(A, n)
```

## **Description**

inthist(A,n) computes the histogram of the elements of integer array A between 0 and n-1. A must have an integer type (int8/16/32/64 or uint8/16/32/64). The result is a row vector h of length n, where h(i) is the number of elements in A with value i-1.

## **Example**

```
A = map2int(rand(100), 0, 1, 'uint8');
h = inthist(A, 10)
h =
    37 31 34 34 32 35 38 36 36 32
```

#### See also

hist

# ipermute

Inverse permutation of the dimensions of an array.

# **Syntax**

```
B = ipermute(A, perm)
```

# **Description**

ipermute(A, perm) returns an array with the same elements as A, but where dimensions are permuted according to the vector of dimensions perm. It performs the inverse permutation of permute. perm must contain integers from 1 to n; dimension i in A becomes dimension perm(i) in the result.

## **Example**

```
size(ipermute(rand(3,4,5), [2,3,1]))
5 3 4
```

permute, ndims, squeeze

# isempty

Test for empty array, list or struct.

## **Syntax**

```
b = isempty(A)
b = isempty(list)
b = isempty(S)
```

## **Description**

isempty(obj) gives true if obj is the empty array [] of any type (numeric, char, logical or cell array) or the empty struct, and false otherwise.

## **Examples**

```
isempty([])
  true
isempty(0)
  false
isempty('')
  true
isempty({{}})
  true
isempty({{}})
  false
isempty(struct)
  true
```

#### See also

size, length

### iscell

Test for cell arrays.

```
b = iscell(X)
```

iscell(X) gives true if X is a cell array or a list, and false otherwise.

## **Examples**

```
iscell({1;2})
    true
iscell({1,2})
    true
islist({1;2})
    false
```

#### See also

islist

## ismember

Test for set membership.

# **Syntax**

```
b = ismember(m, s)
(b, ix) = ismember(m, s)
```

# **Description**

ismember (m, s) tests if elements of array m are members of set s. The result is a logical array the same size as m; each element is true if the corresponding element of m is a member of s, or false otherwise. m must be a numeric array or a cell array, matching type of set s.

With a second output argument ix, ismember also gives the index of the corresponding element of m in s, or 0 if the element is not a member of s.

## **Example**

```
s = {'a','bc','bbb','de'};
m = {'d','a','x';'de','a','z'};
(b, ix) = ismember(m, s)
b =
F T F
T T F
ix =
0 1 0
4 1 0
```

intersect, union, setdiff, setxor

# length

Length of a vector or a list.

## **Syntax**

```
n = length(v)
n = length(list)
```

## **Description**

length(v) gives the length of vector v. length(A) gives the number of elements along the largest dimension of array A. length(list) gives the number of elements in a list.

## **Examples**

```
length(1:5)
5
length((1:5)')
5
length(ones(2,3))
3
length({1, 1:6, 'abc'})
3
length({{}})
1
```

### See also

```
size, numel, end
```

# linspace

Sequence of linearly-spaced elements.

```
v = linspace(x1, x2)
v = linspace(x1, x2, n)
```

linspace(x1,x2) produces a row vector of 50 values spaced linearly from x1 to x2 inclusive. With a third argument, linspace(x1,x2,n) produces a row vector of n values.

## **Examples**

```
linspace(1,10)
  1.0000 1.1837 1.3673 ... 9.8163 10.0000
linspace(1,2,6)
  1.0 1.2 1.4 1.6 1.8 2.0
```

#### See also

logspace, operator:

# logspace

Sequence of logarithmically-spaced elements.

## **Syntax**

```
v = logspace(x1, x2)
v = logspace(x1, x2, n)
```

# Description

logspace(x1,x2) produces a row vector of 50 values spaced logarithmically from  $10^x1$  to  $10^x2$  inclusive. With a third argument, logspace(x1,x2,n) produces a row vector of n values.

# **Example**

```
logspace(0,1)
1.0000 1.0481 1.0985 ... 9.1030 9.5410 10.0000
```

#### See also

linspace, operator:

# magic

Magic square.

## **Syntax**

```
M = magic(n)
```

## **Description**

A magic square is a square array of size n-by-n which contains each integer between 1 and  $n^2$ , and whose sum of each column and of each line is equal. magic(n) returns magic square of size n-by-n.

There is no 2-by-2 magic square. If the size is 2, the matrix [1,3;4,2] is returned instead.

## **Example**

```
magic(3)
8 1 6
3 5 7
4 9 2
```

#### See also

zeros, ones, eye, rand, randn

# meshgrid

Arrays of X-Y coordinates.

# **Syntax**

```
(X, Y) = meshgrid(x, y)
(X, Y) = meshgrid(x)
```

# **Description**

meshgrid(x,y) produces two arrays of x and y coordinates suitable for the evaluation of a function of two variables. The input argument x is copied to the rows of the first output argument, and the input argument y is copied to the columns of the second output argument, so that both arrays have the same size. meshgrid(x) is equivalent to meshgrid(x,x).

### **Example**

```
(X, Y) = meshgrid(1:5, 2:4)
 X =
   1 2
        3 4
               5
     2 3 4
               5
   1
              5
   1 2 3 4
 Y =
     2 2 2
   2
              2
   3 3 3 3 3
   4
      4
        4
           4
              4
Z = atan2(X, Y)
 7 =
   0.4636
             0.7854
                      0.9828
                                1.1071
                                         1.1903
   0.3218
             0.5880
                      0.7854
                                0.9273
                                         1.0304
   0.2450
             0.4636
                      0.6435
                                0.7854
                                         0.8961
```

#### See also

ndgrid, repmat

# ndgrid

Arrays of N-dimension coordinates.

## **Syntax**

```
(X1, \ldots, Xn) = ndgrid(x1, \ldots, xn)
(X1, \ldots, Xn) = ndgrid(x)
```

# **Description**

ndgrid(x1,...,xn) produces n arrays of n dimensions. Array i is obtained by reshaping input argument i as a vector along dimension i and replicating it along all other dimensions to match the length of other input vectors. All output arguments have the same size.

With one input argument, ndgrid reuses it to match the number of output arguments.

```
(Y,X)=ndgrid(y,x) is equivalent to (X,Y)=meshgrid(x,y).
```

## **Example**

```
(X1, X2) = ndgrid(1:3)

X1 =

1    1    1

2    2    2

3    3    3
```

```
X2 =

1 2 3

1 2 3

1 2 3
```

meshgrid, repmat

### ndims

Number of dimensions of an array.

# **Syntax**

```
n = ndims(A)
```

## **Description**

ndims(A) returns the number of dimensions of array A, which is at least 2. Scalars, row and column vectors, and matrices have 2 dimensions.

# **Examples**

```
ndims(magic(3))
2
ndims(rand(3,4,5))
3
```

### See also

size, squeeze, permute, ipermute

### nnz

Number of nonzero elements.

```
n = nnz(A)
```

nnz(A) returns the number of nonzero elements of array A. Argument A must be a numeric, char or logical array.

### **Examples**

```
nnz(-2:2)
4
nnz(magic(3) > 3)
6
```

#### See also

find

### num2cell

Conversion from numeric array to cell array.

## **Syntax**

```
C = num2cell(A)
C = num2cell(A, dims)
```

# **Description**

num2cell(A) creates a cell array the same size as numeric array A. The value of each cell is the corresponding elements of A.

num2cell(A,dims) cuts array A along the dimensions *not* in dims and creates a cell array with the result. Dimensions of cell array are the same as dimensions of A for dimensions not in dims, and 1 for dimensions in dims; dimensions of cells are the same as dimensions of A for dimensions in dims. and 1 for dimensions not in dims.

Argument A can be a numeric array of any dimension and class, a logical array, or a char array.

## **Examples**

```
num2cell([1, 2; 3, 4])
    {1, 2; 3, 4}
num2cell([1, 2; 3, 4], 1)
    {[1; 3], [2; 4]}
num2cell([1, 2; 3, 4], 2)
    {[1, 2]; [3, 4]}
```

num2list, permute

### numel

Number of elements of an array.

# **Syntax**

```
n = numel(A)
```

## **Description**

numel(A) gives the number of elements of array A. It is equivalent to prod(size(A)).

# **Examples**

```
numel(1:5)
   5
numel(ones(2, 3))
   6
numel({1, 1:6; 'abc', []})
   4
numel({2, 'vwxyz'})
   7
```

#### See also

size, length

#### ones

Array of ones.

```
A = ones(n)
A = ones(n1, n2, ...)
A = ones([n1, n2, ...])
A = ones(..., type)
```

ones builds an array whose elements are 1. The size of the array is specified by one integer for a square matrix, or several integers (either as separate arguments or in a vector) for an array of any size.

An additional input argument can be used to specify the type of the result. It must be the string 'double', 'single', 'int8', 'int16', 'int32', 'int64', 'uint8', 'uint16', 'uint32', or 'uint64' (64-bit arrays are not supported on all platforms).

## **Examples**

```
ones(2,3)
   1 1 1
   1 1 1
ones(2, 'int32')
   2x2 int32 array
   1 1
   1 1
```

#### See also

zeros, eye, rand, randn, repmat

# permute

Permutation of the dimensions of an array.

## **Syntax**

```
B = permute(A, perm)
```

# Description

permute(A,perm) returns an array with the same elements as A, but where dimensions are permuted according to the vector of dimensions perm. It is a generalization of the matrix transpose operator. perm must contain integers from 1 to n; dimension perm(i) in A becomes dimension i in the result.

## **Example**

```
size(permute(rand(3,4,5), [2,3,1]))
4 5 3
```

ndims, squeeze, ipermute, num2cell

#### rand

Uniformly-distributed random number.

## **Syntax**

```
x = rand
A = rand(n)
A = rand(n1, n2, ...)
A = rand([n1, n2, ...])
A = rand(..., type)
rand('seed', s);
```

## **Description**

rand builds a scalar pseudo-random number uniformly distributed between 0 and 1. The lower bound 0 may be reached, but the upper bound 1 is never. The default generator is based on a scalar 64-bit seed, which theoretically has a period of 2^64-2^32 numbers. This seed can be set with the arguments rand('seed',s), where s is a scalar. rand('seed',s) returns the empty array [] as output argument. To discard it, the statement should be followed by a semicolon. The generator can be changed with rng.

rand(n), rand(n1,n2,...) and rand([n1,n2,...]) return an n-by-n square array or an array of arbitrary size whose elements are pseudo-random numbers uniformly distributed between 0 and 1.

An additional input argument can be used to specify the type of the result, 'double' (default) or 'single'. With the special value 'raw', rand returns an unscaled integer result of type double which corresponds to the uniform output of the random generator before it is mapped to the range between 0 and 1. The scaling factor can be retrieved in the field rawmax of the structure returned by rng.

## **Examples**

```
rand
   0.2361
rand(1, 3)
   0.6679 0.8195 0.2786
rand('seed',0);
rand
   0.2361
```

randn, randi, rng

### randi

Uniformly-distributed integer random number.

## **Syntax**

```
x = randi(nmax)
x = randi(range)
M = randi(..., n)
M = randi(..., n1, n2, ...)
M = randi(..., [n1, n2, ...])
M = randi(..., class)
```

## **Description**

randi(nmax) produces a scalar pseudo-random integer number uniformly distributed between 1 and nmax. randi(range), where range is a two-element vector [nmin,nmax], produces a scalar pseudo-random integer number uniformly distributed between nmin and nmax.

With more numeric input arguments, randi produces arrays of pseudo-random integer numbers. randi(range,n) produces an n-by-n square array, and randi(range,[n1,n2,...]) or randi(range,n1,n2,...) produces an array of the specified size.

The number class of the result can be specified with a final string argument. The default is 'double'.

#### **Examples**

```
randi(10)
   3
randi(10, [1, 5])
   3 4 6 8 1
randi([10,15], [1, 5])
   12 14 13 10 13
randi(8, [1, 5], 'uint8')
   1x5 uint8 array
   3 4 5 7 2
```

#### See also

rand, randn, rng

### randn

Normally-distributed random number

## **Syntax**

```
x = randn
A = randn(n)
A = randn(n1, n2, ...)
A = randn([n1, n2, ...])
A = randn(..., type)
randn('seed', s);
```

## **Description**

randn builds a scalar pseudo-random number chosen from a normal distribution with zero mean and unit variance. The default generator is based on a scalar 64-bit seed, which theoretically has a period of 2^64-2^32 numbers. This seed can be set with the arguments randn('seed',s), where s is a scalar. The seed is the same as the seed of rand and rng. randn('seed',s) returns the empty array [] as output argument. To discard it, the statement should be followed by a semicolon. The generator can be changed with rng.

randn(n), randn(n1,n2,...) and randn([n1,n2,...]) return an n-by-n square array or an array of arbitrary size whose elements are pseudo-random numbers chosen from a normal distribution.

An additional input argument can be used to specify the type of the result. It must be the string 'double' (default) or 'single'.

## **Examples**

```
randn

1.5927

randn(1, 3)

0.7856 0.6489 -0.8141

randn('seed',0);

randn

1.5927
```

#### See also

```
rand, randi, rng
```

# repmat

Replicate an array.

## **Syntax**

```
B = repmat(A, n)
B = repmat(A, m, n)
B = repmat(A, [n1,...])
```

## **Description**

repmat creates an array with multiple copies of its first argument. It can be seen as an extended version of ones, where 1 is replaced by an arbitrary array.

With 3 input arguments, repmat(A,m,n) replicates array A m times vertically and n times horizontally. The type of the first argument (number, character, logical, cell, or structure array) is preserved.

With two input arguments, repmat (A,n) produces the same result as repmat (A,n,n).

With a vector as second argument, the array can be replicated along more than two dimensions; repmat(A,m,n) produces the same result as repmat(A,[m,n]).

## **Examples**

```
repmat([1,2;3,4], 1, 2)
    1 2 1 2
    3 4 3 4
repmat('abc', 3)
    abcabcabc
    abcabcabc
    abcabcabc
    abcabcabc
```

#### See also

```
zeros, ones, operator:, kron, replist
```

# reshape

Rearrange the elements of an array to change its shape.

```
A2 = reshape(A1)
A2 = reshape(A1, n1, n2, ...)
A2 = reshape(A1, [n1, n2, ...])
```

reshape(A1) gives a column vector with all the elements of array A1. If A1 is a variable, reshape(A1) is the same as A1(:).

reshape(A1,n1,n2,...) or reshape(A1,[n1,n2,...]) changes the dimensions of array A1 so that the result has m rows and n columns. A1 must have n1\*n2\*... elements; read row-wise, both A1 and the result have the same elements.

When dimensions are given as separate elements, one of them can be replaced with the empty array []; it is replaced by the value such that the number of elements of the result matches the size of input array.

**Remark:** code should not rely on the internal data layout. Array elements are currently stored row-wise, but this may change in the future. reshape will remain consistant with indexing, though; reshape (A,s)(i)==A(i) for any compatible size s.

## **Example**

```
reshape([1,2,3;10,20,30], 3, 2)
1 2
3 10
20 30
reshape(1:12, 3, [])
1 2 3 4
5 6 7 8
9 10 11 12
```

#### See also

```
operator ()
```

## rng

State of random number generator.

```
rng(type)
rng(seed)
rng(seed, type)
rng(state)
state = rng
```

Random (actually pseudo-random) number generators produce sequences of numbers whose statistics make them difficult to distinguish from true random numbers. They are used by functions rand, randi, randn and random. They are characterized by a type string and a state.

With a numeric input argument, rng(seed) sets the state based on a seed. The state is usually an array of unsigned 32-bit integer numbers. rng uses the seed to produce an internal state which is valid for the type of random number generator. The default seed is 0.

With a string input argument, rng(type) sets the type of the random number generator and resets the state to its initial value (default seed). The following types are recognized:

```
'original' Original generator used until LME 6.
```

'kiss' **or** 'default' Combination of mwc, a 3-shift register, and a congruential generator. The period is about 2^123.

With two input arguments, rng(seed, type) sets both the seed and the type of the random number generator.

With an output argument, state=rng gets the current state, which can be restored later by calling rng(state). The state is a structure.

### **Examples**

```
rng(123);
R = rand(1,2)
R =
    0.2838    0.4196
s = rng
s =
    type: 'original'
    state: real 2x1
    rawmax: 4294967296
R = rand
R =
    0.5788
rng(s)
R = rand
R =
    0.5788
```

<sup>&#</sup>x27;mcg16807' Multiplicative congruential generator. The state is defined by s(i+1)=mod(a\*s(i),m) with  $a=7^5$  and  $m=2^31-1$ , and the generated value is s(i)/m.

<sup>&#</sup>x27;mwc' Concatenation of two 16-bit multiply-with-carry generators. The period is about 2<sup>60</sup>.

#### Reference

The MWC and KISS generators are described in George Marsaglia, *Random numbers for C: The END?*, Usenet, sci.stat.math, 20 Jan 1999.

#### See also

rand, randn, randi

## rot90

Array rotation.

## **Syntax**

```
A2 = rot90(A1)

A2 = rot90(A1, k)
```

## **Description**

rot90(A1) rotates array A1 90 degrees counter-clockwise; the top left element of A1 becomes the bottom left element of A2. If A1 is an array with more than two dimensions, each plane corresponding to the first two dimensions is rotated.

In rot90(A1,k), the second argument is the number of times the array is rotated 90 degrees counter-clockwise. With k=2, the array is rotated by 180 degrees; with k=3 or k=-1, the array is rotated by 90 degrees clockwise.

## **Examples**

```
rot90([1,2,3;4,5,6])
   3 6
   2 5
   1 4
rot90([1,2,3;4,5,6], -1)
   4 1
   5 2
   6 3
rot90([1,2,3;4,5,6], -1)
   6 5 4
   3 2 1
fliplr(flipud([1,2,3;4,5,6]))
   6 5 4
   3 2 1
```

fliplr, flipud, reshape

### setdiff

Set difference.

## **Syntax**

```
c = setdiff(a, b)
(c, ia) = setdiff(a, b)
```

## **Description**

setdiff(a,b) gives the difference between sets a and b, i.e. the set of members of set a which do not belong to b. Sets are any type of numeric, character or logical arrays, or lists or cell arrays of character strings. Multiple elements of input arguments are considered as single members; the result is always sorted and has unique elements.

The optional second output argument is a vector of indices such that if (c,ia)=setdiff(a,b), then c is a(ia).

## **Example**

```
a = {'a','bc','bbb','de'};
b = {'z','bc','aa','bbb'};
(c, ia) = setdiff(a, b)
    c =
        {'a','de'}
    ia =
        1 4
a(ia)
    {'a','de'}
```

#### See also

unique, union, intersect, setxor, ismember

#### setxor

Set exclusive or.

## **Syntax**

```
c = setxor(a, b)
(c, ia, ib) = setxor(a, b)
```

## **Description**

setxor(a,b) performs an exclusive or operation between sets a and b, i.e. it gives the set of members of sets a and b which are not members of the intersection of a and b. Sets are any type of numeric, character or logical arrays, or lists or cell arrays of character strings. Multiple elements of input arguments are considered as single members; the result is always sorted and has unique elements.

The optional second and third output arguments are vectors of indices such that if (c,ia,ib)=setxor(a,b), then c is the union of a(ia) and b(ib).

## **Example**

Set exclusive or can also be computed as the union of a and b minus the intersection of a and b:

```
setdiff(union(a, b), intersect(a, b))
{'a','aa','de','z'}
```

#### See also

unique, union, intersect, setdiff, ismember

#### size

Size of an array.

## **Syntax**

```
v = size(A)
(m, n) = size(A)
m = size(A, i)
```

## **Description**

size(A) returns the number of rows and the number of elements along each dimension of array A, either in a row vector or as scalars if there are two output arguments or more.

size(A,i) gives the number of elements in array A along dimension i: size(A,1) gives the number of rows and size(A,2) the number of columns.

## **Examples**

```
M = ones(3, 5);
size(M)
   3 5
(m, n) = size(M)
   m =
     3
   n =
     5
size(M, 1)
   3
size(M, 2)
   5
```

#### See also

length, numel, ndims, end

### sort

Array sort.

```
(A_sorted, ix) = sort(A)
(A_sorted, ix) = sort(A, dim)
(A_sorted, ix) = sort(A, dir)
(A_sorted, ix) = sort(A, dim, dir)
(list_sorted, ix) = sort(list)
(list_sorted, ix) = sort(list, dir)
```

sort(A) sorts separately the elements of each column of array A, or the elements of A if it is a row vector. The result has the same size as A. Elements are sorted in ascending order, with NaNs at the end. For complex arrays, numbers are sorted by magnitude.

The optional second output argument gives the permutation array which transforms A into the sorted array. It can be used to reorder elements in another array or to sort the rows of a matrix with respect to one of its columns, as shown in the last example below. Order of consecutive identical elements is preserved.

If a second numeric argument dim is provided, the sort is performed along dimension dim (columns if dim is 1, rows if 2, etc.)

An additional argument can specify the ordering direction. It must be the string 'ascending' (or 'a') for ascending order, or 'descending' (or 'd') for descending order. In both cases, NaNs are moved to the end.

sort(list) sorts the elements of a list, which must be strings. Cell arrays are sorted like lists, not column-wise like numeric arrays. The second output argument is a row vector. The direction can be specified with a second input argument.

## **Examples**

```
sort([3,6,2,3,9,1,2])
  1 2 2 3 3 6 9
sort([2,5,3;nan,4,2;6,1,1])
 2
      1
          2
 6
     4
 nan 5
          3
sort([2,5,3;nan,4,2;6,1,1], 'd')
 6
    5
         3
  2
      4
 nan 1
         1
sort({'def', 'abcd', 'abc'})
  {'abc', 'abcd', 'def'}
```

To sort the rows of an array after the first column, one can obtain the permutation vector by sorting the first column, and use it as subscripts on the array rows:

```
M = [2,4; 5,1; 3,9; 4,0]
2 4
5 1
3 9
4 0
(Ms, ix) = sort(M(:,1));
M(ix,:)
```

- 2 4
- 3 9
- 4 0
- 5 1

## **Algorithm**

Shell sort.

#### See also

unique

## squeeze

Suppression of singleton dimensions of an array.

## **Syntax**

```
B = squeeze(A)
```

## **Description**

squeeze(A) returns an array with the same elements as A, but where dimensions equal to 1 are removed. The result has at least 2 dimensions; row and column vectors keep their dimensions.

# **Examples**

```
size(squeeze(rand(1,2,3,1,4)))
2 3 4
size(squeeze(1:5))
1 5
```

#### See also

permute, ndims

### sub2ind

Conversion from row/column subscripts to single index.

```
ind = sub2ind(size, i, j)
ind = sub2ind(size, i, j, k, ...)
```

sub2ind(size,i,j) gives the single index which can be used to retrieve the element corresponding to the i:th row and the j:th column of an array whose size is specified by size. size must be either a scalar for square matrices or a vector of two elements or more for other arrays. If i and j are arrays, they must have the same size: the result is calculated separately for each element and has the same size.

sub2ind also accepts sizes and subscripts for arrays with more than 2 dimensions. The number of indices must match the length of size.

### **Example**

```
M = [3, 6; 8, 9];
M(2, 1)
    8
sub2ind(size(M), 2, 1)
    7
M(3)
    8
```

#### See also

ind2sub, size

#### tril

Extraction of the lower triangular part of a matrix.

### **Syntax**

```
L = tril(M)
L = tril(M.k)
```

# Description

tril(M) extracts the lower triangular part of a matrix; the result is a matrix of the same size where all the elements above the main diagonal are set to zero. A second argument can be used to specify another diagonal: 0 is the main diagonal, positive values are above and negative values below.

## **Examples**

```
M = magic(3)
M =
    8 1 6
    3 5 7
    4 9 2
tril(M)
    8 0 0
    3 5 0
    4 9 2
tril(M,1)
    8 1 0
    3 5 7
    4 9 2
```

#### See also

triu, diag

### triu

Extraction of the upper triangular part of a matrix.

## **Syntax**

```
U = triu(M)
U = triu(M,k)
```

# Description

tril(M) extracts the upper triangular part of a matrix; the result is a matrix of the same size where all the elements below the main diagonal are set to zero. A second argument can be used to specify another diagonal; 0 is the main diagonal, positive values are above and negative values below.

## **Examples**

```
M = magic(3)
M =
    8 1 6
    3 5 7
    4 9 2
triu(M)
    8 1 6
    0 5 7
```

```
0 0 2
triu(M,1)
0 1 6
0 0 7
0 0 0
```

tril, diag

#### union

Set union.

## **Syntax**

```
c = union(a, b)
(c, ia, ib) = union(a, b)
```

## **Description**

union(a,b) gives the union of sets a and b, i.e. it gives the set of members of sets a or b or both. Sets are any type of numeric, character or logical arrays, or lists or cell arrays of character strings. Multiple elements of input arguments are considered as single members; the result is always sorted and has unique elements.

The optional second and third output arguments are vectors of indices such that if (c,ia,ib)=union(a,b), then elements of c are the elements of a(ia) or b(ib); the intersection of a(ia) and b(ib) is empty.

## **Example**

Set exclusive or can also be computed as the union of a and b minus the intersection of a and b:

```
setdiff(union(a, b), intersect(a, b))
{'a','aa','de','z'}
```

#### See also

unique, intersect, setdiff, setxor, ismember

# unique

Keep unique elements.

## **Syntax**

```
v2 = unique(v1)
list2 = unique(list1)
(b, ia, ib) = unique(a)
```

## **Description**

With an array argument, unique(v1) sorts its elements and removes duplicate elements. Unless v1 is a row vector, v1 is considered as a column vector.

With an argument which is a list of strings, unique(list) sorts its elements and removes duplicate elements.

The optional second output argument is set to a vector of indices such that if (b,ia)=unique(a), then b is a(ia).

The optional third output argument is set to a vector of indices such that if (b,ia,ib)=unique(a), then a is b(ib).

# **Examples**

```
(b,ia,ib) = unique([4,7,3,8,7,1,3])
b =
    1 3 4 7 8
ia =
    6 3 1 2 4
ib =
    3 4 2 5 4 1 2
unique({'def', 'ab', 'def', 'abc'})
    {'ab', 'abc', 'def'}
```

#### See also

sort, union, intersect, setdiff, setxor, ismember

## unwrap

Unwrap angle sequence.

## **Syntax**

```
a2 = unwrap(a1)
a2 = unwrap(a1, tol)
A2 = unwrap(A1, tol, dim)
```

## **Description**

unwrap(a1), where a1 is a vector of angles in radians, returns a vector a2 of the same length, with the same values modulo  $2\pi$ , starting with the same value, and where differences between consecutive values do not exceed  $\pi$ . It is useful for interpolation in a discrete set of angles and for plotting.

With two input arguments, unwrap(a1, tol) reduces the difference between two consecutive values only if it is larger (in absolute value) than tol. If tol is smaller than  $\pi$ , or the empty array [], the default value of  $\pi$  is used.

With three input arguments, unwrap(A1,tol,dim) operates along dimension dim. The result is an array of the same size as A1. The default dimension for arrays is 1.

# **Example**

```
unwrap([0, 2, 4, 6, 0, 2])
0.00 2.00 4.00 6.00 6.28 8.28
```

#### See also

mod, rem

#### zeros

Null array.

## **Syntax**

```
A = zeros(n)
A = zeros(n1,n2,...)
A = zeros([n1,n2,...])
A = zeros(..., type)
```

## **Description**

zeros builds an array whose elements are 0. The size of the array is specified by one integer for a square matrix, or several integers (either as separate arguments or in a vector) for an array of any size.

An additional input argument can be used to specify the type of the result. It must be the string 'double', 'single', 'int8', 'int16', 'int32', 'int64', 'uint8', 'uint16', 'uint32', or 'uint64' (64-bit arrays are not supported on all platforms).

# **Examples**

```
zeros([2,3])

0 0 0

0 0 0

zeros(2)

0 0

0 0

zeros(1, 5, 'uint16')

1x5 uint16 array

0 0 0 0
```

#### See also

ones, cell, eye, rand, randn, repmat

# 6.20 Triangulation Functions

# delaunay

2-d Delaunay triangulation.

# **Syntax**

```
t = delaunay(x, y)

(t, e) = delaunay(x, y)
```

# Description

delaunay (x,y) calculates the Delaunay triangulation of 2-d points given by arrays x and y. Both arrays must have the same number of values, m. The result is an array of three columns. Each row corresponds to a triangle; values are indices in x and y.

The second output argument, if requested, is a logical vector of size m-by-1; elements are true if the corresponding point in x and y belongs to the convex hull of the set of points.

The Delaunay triangulation is a net of triangles which link all the starting points in such a way that no point is included in the circumscribed circle of any other triangle. Triangles are "as equilateral" as possible.

### **Example**

Delaunay triangulation of 20 random points:

```
x = rand(20, 1);
y = rand(20, 1);
(t, e) = delaunay(x, y);
```

With Sysquake graphical functions, points belonging to the convex hull are displayed as crosses and interior points as circles:

```
clf;
scale equal;
plot(x(e), y(e), 'x');
plot(x(~e), y(~e), 'o');
```

Array of vertex indices is modified to have closed triangles:

```
t = [t, t(:, 1)];
```

Triangles are displayed:

```
plot(x(t), y(t));
```

#### See also

delaunayn, voronoi

# delaunayn

N-d Delaunay triangulation.

# **Syntax**

```
t = delaunayn(x)
(t, e) = delaunayn(x)
```

## **Description**

delaunayn(x) calculates the Delaunay triangulation of points given by the rows of array x in a space of dimension size(x,2). The result is an array with one more column. Each row corresponds to a simplex; values are row indices in x and give the vertices of each polyhedron.

The second output argument, if requested, is a logical vector with as many elements as rows in x; elements are true if the corresponding point in x belongs to the convex hull of the set of points.

#### See also

delaunay, tsearchn, voronoin

# griddata

Data interpolation in 2-d plane.

## **Syntax**

```
vi = griddata(x, y, v, xi, yi)
vi = griddata(x, y, v, xi, yi, method)
```

# **Description**

griddata(x,y,v,xi,yi) interpolates values at coordinates given by the corresponding elements of arrays xi and yi in a 2-dimension plane. Original data are defined by corresponding elements of arrays x, y, and v (which must have the same size), such that the value at coordinate [x(i);y(i)] is v(i). The result is an array with the same size as xi and yi where vi(j) is the value interpolated at [xi(j);yi(j)].

All coordinates are real (imaginary components are ignored). Values v and vi can be real or complex. The result for coordinates outside the convex hull defined by x and y is NaN.

griddata is based on Delaunay triangulation. The interpolation method used in each triangle is linear by default, or can be specified with an additional input argument, a string:

Argument	Meaning
'0' or 'nearest'	nearest value
'1' or 'linear'	linear

## **Example**

Nearest value interpolation in 2D plane of a few values v(x,y). The plane is sampled with a regular grid with meshgrid.

```
x = [0.2; 1.8; 0.7; 0.9; 1.6];
y = [0.2; 0.7; 1.8; 1.1; 1.7];
v = [0.1; 0.3; 0.9; 0.5; 0.4];
(xi, yi) = meshgrid(0:0.01:2);
vi = griddata(x, y, v, xi, yi, '0');
```

In Sysquake, the result can be displayed as a contour plot. For locations where the values cannot be interpolated, i.e. outside the convex hull defined by x and y, values are set to 0.

```
vi(isnan(vi)) = 0;
contour(vi, [], 20);
```

#### See also

delaunay, tsearch, griddatan, interpn

# griddatan

Data interpolation in N-d space.

# **Syntax**

```
vi = griddatan(x, v, xi)
vi = griddatan(x, v, xi, method)
```

# **Description**

griddatan(x,v,xi) interpolates values at coordinates given by the p rows of p-by-n array xi in an n-dimension space. Original data are defined by m-by-n array x and m-by-1 column vector v, such that the value at coordinate x(i,:)' is v(i). The result is a p-by-1 column vector vi where vi(j) is the value interpolated at xi(j,:)'.

Coordinates x and xi are real (imaginary components are ignored). Values v and vi can be real or complex. The result for coordinates outside the convex hull defined by x is NaN.

griddatan is based on Delaunay triangulation. The interpolation method used in each simplex is linear by default, or can be specified with an additional input argument, a string:

```
Argument Meaning
'0' or 'nearest' nearest value
'1' or 'linear' linear
```

### **Example**

Linear interpolation in 2D plane of a few values v(x,y). The plane is sampled with a regular grid with meshgrid. Since griddatan interpolates a 1-dim array of points, the result is reshaped to match x and y (compare with the example of griddata).

```
x = [0.2; 1.8; 0.7; 0.9; 1.6];
y = [0.2; 0.7; 1.8; 1.1; 1.7];
v = [0.1; 0.3; 0.9; 0.5; 0.4];
(xi, yi) = meshgrid(0:0.01:2);
vi = griddatan([x,y], v, [xi(:),yi(:)], '1');
vi = reshape(vi, size(xi));
```

In Sysquake, the result can be displayed as a contour plot. For locations where the values cannot be interpolated, i.e. outside the convex hull defined by x and y, values are set to 0.

```
vi(isnan(vi)) = 0;
contour(vi, [], 20);
```

#### See also

delaunayn, tsearchn, griddata, interpn

### tsearch

Search of points in triangles.

# Syntax

```
ix = tsearch(x, y, t, xi, yi)
```

# Description

tsearch(x,y,t,xi,yi) searches in which triangle is located each point given by the corresponding elements of arrays xi and yi. Corresponding elements of arrays x and y represent the vertices of the triangles, and rows of array t represent their indices in x and y; array t is usually the result of delaunay. Dimensions of x and y, and of xi and yi, must be equal. The result is an array with the same size as xi and yi where each element is the row index in t of the first triangle which contains the point, or NaN if the point is outside all triangles (i.e. outside the convex hull of points defined by x and y if t is a proper triangulation such as the one computed with delaunay).

### **Example**

Search for triangles containing points [0,0] and [0,1] corresponding to Delaunay triangulation of 20 random points:

```
x = randn(20, 1);
y = randn(20, 1);
t = delaunay(x, y);
xi = [0, 0];
yi = [0, 1];
ix = tsearch(x, y, t, xi, yi);
```

#### See also

tsearchn, delaunay, voronoi, griddata

### tsearchn

Search of points in triangulation simplices.

# **Syntax**

```
ix = tsearchn(x, t, xi)
```

# **Description**

tsearchn(x,t,xi) searches in which simplex each point given by the rows of array xi is located. Rows of array x represent the vertices of the simplices, and rows of array t represent their indices in x; array t is usually the result of delaunayn. Dimensions must match: in a space of n dimensions, x and xi have n columns, and t has n+1 columns. The result is a column vector with one element for each row of xi, which is the row index in t of the first simplex which contains the point, or NaN if the point is outside all simplices (i.e. outside the convex hull of points x if t is a proper triangulation of x such as the one computed with delaunayn).

# **Example**

Search for simplices containing points [0,0] and [0,1] corresponding to Delaunay triangulation of 20 random points:

```
x = randn(20, 2);
t = delaunayn(x);
xi = [0, 0; 0, 1];
ix = tsearchn(x, t, xi);
```

tsearch, delaunayn, voronoin, griddatan

#### voronoi

2-d Voronoi tessalation.

### **Syntax**

```
(v, p) = voronoi(x, y)
```

# **Description**

voronoi(x,y) calculates the Voronoi tessalation of the set of 2-d points given by arrays x and y. Both arrays must have the same number of values, m. The first output argument v is an array of two columns which contains the coordinates of the vertices of the Voronoi cells, one row per vertex. The first row contains infinity and is used as a marker for unbounded Voronoi cells. The second output argument p is a list of vectors of row indices in v; each element describes the Voronoi cell corresponding to a point in x. In each cell, vertices are sorted counterclockwise.

Voronoi tessalation is a tessalation (a partition of the plane) such that each region is the set of points closer to one of the initial point than to any other one. Two regions are in contact if and only if their initial points are linked in the corresponding Delaunay triangulation.

# **Example**

Voronoi tessalation of 20 random points:

```
x = rand(20, 1);
y = rand(20, 1);
(v, p) = voronoi(x, y);
```

These points are displayed as crosses with Sysquake graphical functions. The scale is fixed, because Voronoi polygons can have vertices which are far away from the points.

```
clf;
scale('equal', [0,1,0,1]);
plot(x, y, 'x');
```

Voronoi polygons are displayed in a loop, skipping unbounded polygons. The first vertex is repeated to have closed polygons. Since plot expects row vectors, vertex coordinates are transposed.

```
for p1 = p
  if ~any(p1 == 1)
    p1 = [p1, p1(1)];
    plot(v(p1,1)', v(p1,2)');
  end
end
```

voronoin, delaunay

### voronoin

N-d Voronoi tessalation.

## **Syntax**

```
(v, p) = voronoin(x)
```

# **Description**

voronoin(x) calculates the Voronoi tessalation of the set of points given by the rows of arrays x in a space of dimension n=size(x,2). The first output argument v is an array of n columns which contains the coordinates of the vertices of the Voronoi cells, one row per vertex. The first row contains infinity and is used as a marker for unbounded Voronoi cells. The second output argument p is a list of vectors of row indices in v; each element describes the Voronoi cell corresponding to a point in x. In each cell, vertices are sorted by index.

#### See also

voronoi, delaunayn

# 6.21 Integer Functions

# uint8 uint16 uint32 uint64 int8 int16 int32 int64

Conversion to integer types.

## **Syntax**

```
B = uint8(A)

B = uint16(A)

B = uint32(A)

B = uint64(A)

B = int8(A)

B = int16(A)

B = int32(A)

B = int64(A)
```

## Description

The functions convert a number or an array to unsigned or signed integers. The name contains the size of the integer in bits.

To avoid a conversion from double to integer, constant literal numbers should be written with a type suffix, such as 12int32. This is the only way to specify large 64-bit numbers, because double-precision floating-point numbers have a mantissa of 56 bits.

Constant arrays of uint8 can also be encoded in a compact way using base64 inline data.

uint64 and int64 are not supported on platforms with tight memory constraints.

# **Examples**

```
uint8(3)
  3uint8
3uint8
  3uint8
uint8([50:50:400])
  1x8 uint8 array
    50 100 150 200 250 44 94 144
@/base64 MmSWyPosXpA=
   50
  100
  . . .
  144
int8([50:50:400])
  1x8 int8 array
    50 100 - 106 - 56 - 6
                             44
                                   94 - 112
```

The base64 data above is obtained with the following expression:

```
base64encode(uint8([50:50:400]))
```

#### See also

double, single, char, logical, map2int

### intmax

Largest integer.

## **Syntax**

```
i = intmax
i = intmax(type)
```

# **Description**

Without input argument, intmax gives the largest signed 32-bit integer. intmax(type) gives the largest integer of the type specified by string type, which can be 'uint8', 'uint16', 'uint32', 'uint64', 'int8', 'int16', 'int32', or 'int64' (64-bit integers are not supported on all platforms). The result has the corresponding integer type.

## **Examples**

```
intmax
  2147483647int32
intmax('uint16')
  65535uint16
```

#### See also

intmin, realmax, flintmax, uint8 and related functions, map2int

### intmin

Smallest integer.

# Syntax

```
i = intmin
i = intmin(type)
```

# **Description**

Without input argument, intmin gives the smallest signed 32-bit integer. intmin(type) gives the largest integer of the type specified by string type, which can be 'uint8', 'uint16', 'uint32', 'uint64', 'int8', 'int16', 'int32', or 'int64' (64-bit integers are not supported on all platforms). The result has the corresponding integer type.

### **Examples**

```
intmin
    -2147483648int32
intmin('uint16')
    0uint16
```

#### See also

intmax, realmin, uint8 and related functions, map2int

# map2int

Mapping of a real interval to an integer type.

# **Syntax**

```
B = map2int(A)
B = map2int(A, vmin, vmax)
B = map2int(A, vmin, vmax, type)
```

# Description

map2int(A,vmin,vmax) converts number or array A to 8-bit unsigned integers. Values between vmin and vmax in A are mapped linearly to values 0 to 255. With a single input argument, the default input interval is 0 to 1.

map2int(A,vmin,vmax,type) converts A to the specified type, which can be any integer type given as a string: 'uint8', 'uint16', 'uint32', 'uint64', 'int8', 'int16', 'int32', or 'int64' (64-bit integers are not supported on all platforms). The input interval is mapped to its full range.

In all cases, input values outside the interval are clipped to the minimum or maximum values.

# **Examples**

```
map2int(-0.2:0.2:1.2)
  1x5 uint8 array
    0    0   51 102 153 204 255 255
map2int([1,3,7], 0, 10, 'uint16')
  1x3 uint16 array
    6553 19660 45875
map2int([1,3,7], 0, 10, 'int16')
  1x3 int16 array
    -26214 -13107 13107
```

uint8 and related functions.

# 6.22 Non-Linear Numerical Functions

### **fminbnd**

Minimum of a function.

## **Syntax**

```
(x, y) = fminbnd(fun, x0)
(x, y) = fminbnd(fun, [xlow,xhigh])
(x, y) = fminbnd(..., options)
(x, y) = fminbnd(..., options, ...)
(x, y, didConverge) = fminbnd(...)
```

## **Description**

fminbnd(fun,...) finds numerically a local minimum of function fun. fun is either specified by its name or given as an anonymous or inline function or a function reference. It has at least one input argument x, and it returns one output argument, also a real number. fminbnd finds the value x such that fun(x) is minimized.

Second argument tells where to search; it can be either a starting point or a pair of values which must bracket the minimum.

The optional third argument may contain options. It is either the empty array [] for default options, or the result of optimset.

Remaining input arguments of fminbnd, if any, are given as additional input arguments to function fun. They permit to parameterize the function. For example fminbnd('fun',x0,[],2,5) calls fun as fun(x,2,5) and minimizes its value with respect to x.

The first output argument of fminbnd is the value of x at optimum. The second output argument, if it exists, is the value of fun(x) at optimum. The third output argument, if it exists, is set to true if fminbnd has converged to an optimum, or to false if it has not; in that case, other output arguments are set to the best value obtained. With one or two output arguments, fminbnd throws an error if it does not converge.

# **Examples**

Minimum of a sine near 2, displayed with 15 digits:

```
fprintf('%.15g\n', fminbnd(@sin, 2));
  4.712389014989218
```

To find the minimum of  $ce^x - \sin x$  between -1 and 10 with c = 0.1, the expression is written as an inline function stored in variable fun:

```
fun = inline('c*exp(x)-sin(x)', 'x', 'c');
```

Then fminbnd is used, with the value of c passed as an additional argument:

```
x = fminbnd(fun,[-1,10],[],0.1)
x =
    1.2239
```

With an anonymous function, this becomes

```
c = 0.1;
fun = @(x) c*exp(x)-sin(x);
x = fminbnd(fun,[-1,10])
x =
1.2239
```

Attempt to find the minimum of an unbounded function:

```
(x,y,didConverge) = fminbnd(@exp,10)
x =
    -inf
y =
    0
didConverge =
    false
```

#### See also

optimset, fminsearch, fzero, inline, operator@

### **fminsearch**

Minimum of a function in R<sup>n</sup>.

### **Syntax**

```
x = fminsearch(fun, x0)
x = fminsearch(..., options)
x = fminsearch(..., options, ...)
(x, y, didConverge) = fminsearch(...)
```

## **Description**

fminsearch(fun,x0,...) finds numerically a local minimum of function fun. fun is either specified by its name or given as an anonymous or inline function or a function reference. It has at least one input argument x, a real scalar, vector or array, and it returns one output argument, a scalar real number. fminsearch finds the value x such that fun(x) is minimized, starting from point x0.

The optional third input argument may contain options. It is either the empty array [] for default options, or the result of optimset.

Remaining input arguments of fminsearch, if any, are given as additional input arguments to function fun. They permit to parameterize the function. For example fminsearch('fun',x0,[],2,5) calls fun as fun(x,2,5) and minimizes its value with respect to x.

The first output argument of fminsearch is the value of x at optimum. The second output argument, if it exists, is the value of fun(x) at optimum. The third output argument, if it exists, is set to true if fminsearch has converged to an optimum, or to false if it has not; in that case, other output arguments are set to the best value obtained. With one or two output arguments, fminsearch throws an error if it does not converge.

# **Algorithm**

fminsearch implements the Nelder-Mead simplex method. It starts from a polyhedron centered around x0 (the "simplex"). Then at each iteration, either vertex  $x_i$  with the maximum value fun( $x_i$ ) is moved to decrease it with a reflexion-expansion, a reflexion, or a contraction; or the simplex is shrinked around the vertex with minimum value. Iterations stop when the simplex is smaller than the tolerance, or when the maximum number of iterations or function evaluations is reached (then an error is thrown).

# **Examples**

Minimum of a sine near 2, displayed with 15 digits:

```
fprintf('%.15g\n', fminsearch(@sin, 2));
  4.712388977408411
```

Maximum of  $xe^{-x^2y^2}xy - 0.1x^2$  The function is defined as an anonymous function stored in variable fun:

```
fun = @(x,y) x.*exp(-(x.*y).^2).*x.*y-0.1*x.^2;
```

In Sysquake, the contour plot can be displayed with the following commands:

```
[X,Y] = meshgrid(0:0.02:3, 0:0.02:3);
contour(feval(fun, X, Y), [0,3,0,3], 0.1:0.05:0.5);
```

The maximum is obtained by minimizing the opposite of the function, rewritten to use as input a single variable in R<sup>2</sup>:

Here is another way to find this maximum, by calling fun from an intermediate anonymous function:

```
fminsearch(@(X) -fun(X(1),X(2)), [1, 2]) 2.1444 \quad 0.3297
```

For the same function with a constraint x < 1, the objective function can be modified to return  $+\infty$  for inputs outside the feasible region (note that we can start on the constraint boundary, but starting from the infeasible region would probably fail):

```
fminsearch(@(X) X(1) < 1? -fun(X(1),X(2)) : inf, [1, 2]) 1 0.7071
```

#### See also

optimset, fminbnd, lsgnonlin, fsolve, inline, operator @

### fsolve

Solve a system of nonlinear equations.

# **Syntax**

```
x = fsolve(fun, x0)
x = fsolve(..., options)
x = fsolve(..., options, ...)
(x, y, didConverge) = fsolve(...)
```

# **Description**

fsolve(fun,  $x0, \ldots$ ) finds numerically a zero of function fun. fun is either specified by its name or given as an anonymous or inline function or a function reference. It has at least one input argument x, a real scalar, vector or array, and it returns one output argument y whose size should match x. fsolve attempts to find the value y such that fun(y) is zero, starting from point y0. Depending on the

existence of any solution and on the choice of x0, fsolve may fail to find a zero.

The optional third input argument may contain options. It is either the empty array [] for default options, or the result of optimset.

Remaining input arguments of fsolve, if any, are given as additional input arguments to function fun. They permit to parameterize the function. For example fsolve(@fun,x0,[],2,5) finds the value of x such that the result of fun(x,2,5) is zero.

The first output argument of fsolve is the value of x at zero. The second output argument, if it exists, is the value of fun(x) at zero; it should be a vector or array whose elements are zero, up to the tolerance, unless fsolve cannot find it. The third output argument, if it exists, is set to true if fsolve has converged to a solution, or to false if it has not; in that case, other output arguments are set to the best value obtained. With one or two output arguments, fsolve throws an error if it does not converge.

## **Algorithm**

fsolve minimizes the sum of squares of the vector elements returned by fun using the Nelder-Mead simplex method of fminsearch.

# **Example**

One of the zeros of  $x1^2+x2^2=10$ , x2=exp(x1):

```
[x, y, didConverge] = fsolve(@(x) [x(1)^2+x(2)^2-10; x(2)-exp(x(1))], [0; 0]
x =
   -3.1620
    0.0423
y =
   -0.0000
   -0.0000
didConverge =
   true
```

#### See also

optimset, fminsearch, fzero, inline, operator@

#### fzero

Zero of a function.

## **Syntax**

```
x = fzero(fun,x0)
x = fzero(fun,[xlow,xhigh])
x = fzero(...,options)
x = fzero(...,options,...)
```

## **Description**

fzero (fun,...) finds numerically a zero of function fun. fun is either specified by its name or given as an anonymous or inline function or a function reference. It has at least one input argument x, and it returns one output argument, also a real number. fzero finds the value x such that fun(x)==0, up to some tolerance.

Second argument tells where to search; it can be either a starting point or a pair of values xlow and xhigh which must bracket the zero, such that fun(xlow) and fun(xhigh) have opposite sign.

The optional third argument may contain options. It is either the empty array [] for the default options, or the result of optimset.

Additional input arguments of fzero are given as additional input arguments to the function specified by fun. They permit to parameterize the function.

## **Examples**

Zero of a sine near 3, displayed with 15 digits:

```
fprintf('%.15g\n', fzero(@sin, 3));
  3.141592653589793
```

To find the solution of  $e^x = c + \sqrt{x}$  between 0 and 100 with c = 10, a function f whose zero gives the desired solution is written:

```
function y = f(x,c)

y = exp(x) - c - sqrt(x);
```

Then fsolve is used, with the value of c passed as an additional argument:

```
x = fzero(@f,[0,100],[],10)
x =
    2.4479
f(x,10)
    1.9984e-15
```

An anonymous function can be used to avoid passing 10 as an additional argument, which can be error-prone since a dummy empty option arguments has to be inserted.

```
x = fzero(@(x) f(x,10), [0,100])

x = 2.4479
```

optimset, fminsearch, inline, operator @, roots

# integral

Numerical integration.

## **Syntax**

```
y = integral(fun, a, b)
y = integral(fun, a, b, options)
```

# **Description**

integral (fun,a,b) integrates numerically function fun between a and b. fun is either specified by its name or given as an anonymous or inline function or a function reference. It has a single input argument and a single output argument, both scalar real or complex.

Options can be provided with named arguments. The following options are accepted:

Name	Default	Meaning
AbsTol	1e-6	maximum absolute error
RelTol	1e-3	maximum relative error
Display	false	statistics display

# **Example**

$$\int_0^2 te^{-t}\mathrm{d}t$$
 integral(@(t) t\*exp(-t), 0, 2, AbsTol=1e-9) 0.5940

#### See also

sum, ode45, inline, operator @

# **Isqcurvefit**

Least-square curve fitting.

## **Syntax**

```
param = lsqcurvefit(fun, param0, x, y)
param = lsqcurvefit(..., options)
param = lsqcurvefit(..., options, ...)
(param, r, didConverge) = lsqcurvefit(...)
```

## **Description**

lsqcurvefit(fun,p0,x,y,...) finds numerically the parameters of function fun such that it provides the best fit for the curve defined by x and y in a least-square sense. fun is either specified by its name or given as an anonymous or inline function or a function reference. It has at least two input arguments: p, the parameter vector, and x, a vector or array of input data; it returns one output argument, a vector or array the same size as x and y. Its header could be

```
function y = f(param, x)
```

lsqcurvefit finds the value p which minimizes  $sum((fun(p,x)-y).^2)$ , starting from parameters p0. All values are real.

The optional fifth input argument may contain options. It is either the empty array [] for default options, or the result of optimset.

Remaining input arguments of lsqcurvefit, if any, are given as additional input arguments to function fun. They permit to parameterize the function. For example lsqcurvefit('fun',p0,x,y,[],2,5) calls fun as fun(p,x,2,5) and find the (local) least-square solution with respect to p.

The first output argument of lsqcurvefit is the value of p at optimum. The second output argument, if it exists, is the value of the cost function at optimum. The third output argument, if it exists, is set to true if lsqcurvefit has converged to an optimum, or to false if it has not; in that case, other output arguments are set to the best value obtained. With one or two output arguments, lsqcurvefit throws an error if it does not converge.

# **Algorithm**

Like Isqnonlin, Isqcurvefit is based on the Nelder-Mead simplex method.

# Example

Find the best curve fit of y=a\*sin(b\*x+c) with respect to parameters a, b and c, where x and y are given (see the example of lsqnonlin for another way to solve the same problem).

```
% assume nominal parameter values a0=2, b0=3, c0=1
 a0 = 2: b0 = 3: c0 = 1:
 % reset the seed of rand and randn for reproducible results
 rand('s', 0); randn('s', 0);
 % create x and y, with noise
 x0 = rand(1, 100);
 x = x0 + 0.05 * randn(1, 100);
 y = a0 * sin(b0 * x0 + c0) + 0.05 * randn(1, 100);
 % find least-square curve fit, starting from 1, 1, 1
 p0 = [1; 1; 1];
 p_{s} = lsqcurvefit(@(p, x) p(1) * sin(p(2) * x + p(3)), p0, x, y)
   p_ls =
     2.0060
     2.8504
     1.0836
In Sysquake, the solution can be displayed with
 fplot(@(x) a0 * sin(b0 * x + c0), [0,1], 'r');
```

 $fplot(@(x) p_ls(1)*sin(p_ls(2)*x+p_ls(3)), [min(x), max(x)]);$ 

#### See also

plot(x, y, 'o');

optimset, lsgnonlin, inline, operator @

legend('Nominal\nSamples\nLS fit', 'r\_kok\_');

# Isqnonlin

Nonlinear least-square solver.

# **Syntax**

```
x = lsqnonlin(fun, x0)
x = lsqnonlin(..., options)
x = lsqnonlin(..., options, ...)
(x, y, didConverge) = lsqnonlin(...)
```

# **Description**

lsqnonlin(fun,x0,...) finds numerically the value such that the sum of squares of the output vector produced by fun is a local minimum. fun is either specified by its name or given as an anonymous or inline function or a function reference. It has at least one input argument x, a real scalar, vector or array, and it returns one output argument, a real vector or array. Its header could be

```
function y = f(x)
```

lsqnonlin finds the value x such that  $sum(fun(x(:)).^2)$  is minimized, starting from point x0.

The optional third input argument may contain options. It is either the empty array [] for default options, or the result of optimset.

Remaining input arguments of lsqnonlin, if any, are given as additional input arguments to function fun. They permit to parameterize the function. For example lsqnonlin('fun',x0,[],2,5) calls fun as fun(x,2,5) and find the (local) least-square solution with respect to x.

The first output argument of lsqnonlin is the value of x at optimum. The second output argument, if it exists, is the value of fun(x) at optimum. The third output argument, if it exists, is set to true if lsqnonlin has converged to an optimum, or to false if it has not; in that case, other output arguments are set to the best value obtained. With one or two output arguments, lsqnonlin throws an error if it does not converge.

### **Algorithm**

Like fminsearch, lsqnonlin is based on the Nelder-Mead simplex method.

# **Example**

Find the least-square solution of a\*sin(b\*x+c)-y with respect to parameters a, b and c, where x and y are given (see the example of lsqcurvefit for another way to solve the same problem).

```
% assume nominal parameter values a0=2, b0=3, c0=1 a0 = 2; b0 = 3; c0 = 1; % reset the seed of rand and randn for reproducible results rand('s', 0); randn('s', 0); % create x and y, with noise x0 = \text{rand}(1, 100); x = x0 + 0.05 * \text{randn}(1, 100); y = a0 * \sin(b0 * x0 + c0) + 0.05 * \text{randn}(1, 100); % find least-square solution, starting from 1, 1, 1 p0 = [1; 1; 1]; p_ls = lsqnonlin(@(p) p(1) * <math>\sin(p(2) * x + p(3)) - y, p0) p_ls = 2.0060 2.8504 1.0836
```

In Sysquake, the solution can be displayed with

```
fplot(@(x) a0 * sin(b0 * x + c0), [0,1], 'r');
plot(x, y, 'o');
```

```
fplot(@(x) p_ls(1)*sin(p_ls(2)*x+p_ls(3)), [min(x), max(x)]); legend('Nominal\nSamples\nLS fit', 'r_kok_');
```

optimset, fminsearch, lsqcurvefit, inline, operator@

# ode23 ode45

Ordinary differential equation integration.

## **Syntax**

```
(t,y) = ode23(fun,[t0,tend],y0)
(t,y) = ode23(fun,[t0,tend],y0,options)
(t,y) = ode23(fun,[t0,tend],y0,options,...)
(t,y,te,ye,ie) = ode23(...)
(t,y) = ode45(fun,[t0,tend],y0)
(t,y) = ode45(fun,[t0,tend],y0,options)
(t,y) = ode45(fun,[t0,tend],y0,options,...)
(t,y,te,ye,ie) = ode45(...)
```

# **Description**

ode23(fun,[t0,tend],y0) and ode45(fun,[t0,tend],y0) integrate numerically an ordinary differential equation (ODE). Both functions are based on a Runge-Kutta algorithm with adaptive time step; ode23 is low-order and ode45 high-order. In most cases for non-stiff equations, ode45 is the best method. The function to be integrated is either specified by its name or given as an anonymous or inline function or a function reference. It should have at least two input arguments and exactly one output argument:

```
function yp = f(t,y)
```

The function calculates the derivative yp of the state vector y at time t

Integration is performed over the time range specified by the second argument [t0,tend], starting from the initial state y0. It may stop before reaching tend if the integration step cannot be reduced enough to obtain the required tolerance. If the function is continuous, you can try to reduce MinStep in the options argument (see below).

The optional fourth argument may contain options. It is either the empty array [] for the default options, or the result of odeset (the use of a vector of option values is deprecated.)

Events generated by options Events or EventTime can be obtained by three additional output arguments: (t,y,te,ye,ie)=... returns

event times in te, the corresponding states in ye and the corresponding event identifiers in ie.

Additional input arguments of ode45 are given as additional input arguments to the function specified by fun. They permit to parameterize the ODE.

### **Examples**

Let us integrate the following ordinary differential equation (Van Der Pol equation), parameterized by  $\mu$ :

$$x'' = \mu(1-x^2)x' - x$$

Let  $y_1 = x$  and  $y_2 = x'$ ; their derivatives are

$$y'_1 = y_2$$
  
 $y'_2 = \mu(1-y_1^2)y_2-y_1$ 

and can be computed by the following function:

```
function yp = f(t, y, mu)

yp = [y(2); mu*(1-y(1)^2)*y(2)-y(1)];
```

The following ode45 call integrates the Van Der Pol equation from 0 to 10 with the default options, starting from x(0) = 2 and x'(0) = 0, with  $\mu = 1$  (see Fig. 6.1):

```
(t, y) = ode45(@f, [0,10], [2;0], [], 1);
```

The same result can be obtained with an anonymous function:

```
mu=1;

(t, y) = ode45(@(t,y) [y(2); mu*(1-y(1)^2)*y(2)-y(1)],

[0,10], [2;0]);
```

The plot command expects traces along the second dimension; consequently, the result of ode45 should be transposed.

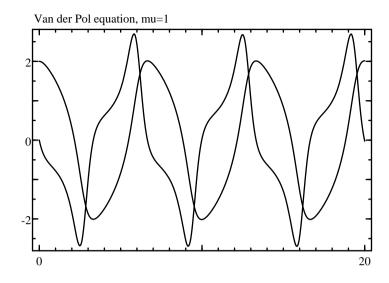
```
plot(t', y');
```

#### See also

odeset, integral, inline, operator @, expm

### odeset

Options for ordinary differential equation integration.



**Figure 6.1** Van der Pol equation with  $\mu = 1$  integrated with ode45

## **Syntax**

```
options = odeset
options = odeset(name1=value1, ...)
options = odeset(name1, value1, ...)
options = odeset(options0, name1=value1, ...)
options = odeset(options0, name1, value1, ...)
```

# **Description**

odeset(name1,value1,...) creates the option argument used by ode23 and ode45. Options are specified with name/value pairs, where the name is a string which must match exactly the names in the table below. Case is significant. Alternatively, options can be given with named arguments. Options which are not specified have a default value. The result is a structure whose fields correspond to each option. Without any input argument, odeset creates a structure with all the default options. Note that ode23 and ode45 also interpret the lack of an option argument, or the empty array [], as a request to use the default values. Options can also be passed directly to ode23 or ode45 as named arguments.

When its first input argument is a structure, odeset adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options (empty arrays mean "auto-

#### matic"):

Name	Default	Meaning
AbsTol	1e-6	maximum absolute error
Events	[] (none)	state-based event function
EventTime	[] (none)	time-based event function
InitialStep	[](10*MinStep)	initial time step
MaxStep	[] (time range/10)	maximum time step
MinStep	[] (time range/1e6)	minimum time step
NormControl	false	error control on state norm
OnEvent	[] (none)	event function
OutputFcn	[] (none)	output function
Past	false	provide past times and states
PreArg	{}	list of prepended input arguments
Refine	[] (1, 4 for ode45)	refinement factor
RelTol	1e-3	maximum relative error
Stats	false	statistics display

### Time steps and output

Several options control how the time step is tuned during the numeric integration. Error is calculated separately on each element of y if NormControl is false, or on norm(y) if it is true; time steps are chosen so that it remains under AbsTol or RelTol times the state, whichever is larger. If this cannot be achieved, for instance if the system is stiff and requires an integration step smaller than MinStep, integration is aborted.

'Refine' specifies how many points are added to the result for each integration step. When it is larger than 1, additional points are interpolated, which is much faster than reducing MaxStep.

The output function OutputFcn, if defined, is called after each step. It is a function name in a string, a function reference, or an anonymous or inline function, which can be defined as

```
function stop = outfun(tn, yn)
```

where tn is the time of the new samples, yn their values, and stop a logical value which is false to continue integrating or true to stop. The number of new samples is given by the value of Refine; when multiple values are provided, tn is a row vector and yn is a matrix whose columns are the corresponding states. The output function can be used for incremental plots, for animations, or for managing large amounts of output data without storing them in variables.

#### **Events**

Events are additional time steps at controlled time, to change instantaneously the states, and to base the termination condition on the states. Time instants where events occur are either given explicitly by EventTime, or implicitly by Events. There can be multiple streams of events, which are checked independently and are identified by a positive integer for Events, or a negative integer for EventTime. For instance, for a ball which bounces between several walls, the intersection between each wall and the ball trajectory would be a different event stream.

For events which occur at regular times, EventTime is an n-by-two matrix: for each row, the first column gives the time step ts, and the second column gives the offset to. Non-repeating events are specified with an infinite time step ts. Events occur at time t=to+k\*ts, where k is an integer.

When event time is varying, EventTime is a function which can be defined as

```
function eventTime = eventtimefun(t, y, ...)
```

where t is the current time, y the current state, and the ellipsis stand for additional arguments passed to ode\*. The function returns a (column) vector whose elements are the times where the next event occurs. In both cases, each row corresponds to a different event stream.

For events which are based on the state, the value of a function which depends on the time and the states is checked; the event occurs when its sign changes. Events is a function which can be defined as

Input arguments are the same as for EventTime. Output arguments are (column) vectors where each element i corresponds to an event stream. An event occurs when value(i) crosses zero, in either direction if direction(i)==0, from negative to nonnegative if direction(i)>0, or from positive to nonpositive if direction(i)<0. The event terminates integration if isterminal(i) is true. The Events function is evaluated for each state obtained by integration; intermediate time steps obtained by interpolation when Refine is larger than 1 are not considered. When an event occurs, the integration time step is reset to the initial value, and new events are disabled during the next integration step to avoid shattering. MaxStep should be used if events are missed when the result of Events is not monotonous between events.

When an event occurs, function 0nEvent is called if it exists. It can be defined as

```
function yn = onevent(t, y, i, ...)
```

where i identifies the event stream (positive for events produced by Events or negative for events produced by EventTime); and the output yn is the new value of the state, immediately after the event.

The primary goal of ode\* functions is to integrate states. However, there are systems where some states are constant between events, and are changed only when an event occurs. For instance, in a relay with hysteresis, the output is constant except when the input overshoots some value. In the general case, ni states are integrated and n-ni states are kept constant between events. The total number of states n is given by the length of the initial state vector y0, and the number of integrated states ni is given by the size of the output of the integrated function. Function OnEvent can produce a vector of size n to replace all the states, of size n-ni to replace the non-integrated states, or empty to replace no state (this can be used to display results or to store them in a file, for instance).

Event times are computed after an integration step has been accepted. If an event occurs before the end of the integration step, the step is shortened; event information is stored in the output arguments of ode\* te, ie and ye; and the OnEvent function is called. The output arguments t and y of ode\* contain two rows with the same time and the state right before the event and right after it. The time step used for integration is not modified by events.

# Additional arguments

Past is a logical value which, if it is true, specifies that the time and state values computed until now (what will eventually be the result of ode23 or ode45) are passed as additional input arguments to functions called during intergration. This is especially useful for delay differential equations (DDE), where the state at some time point in the past can be interpolated from the integration results accumulated until now with interp1. Assuming no additional parameters or PreArg (see below), functions must be defined as

PreArg is a list of additional input arguments for all functions called during integration; they are placed before normal arguments. For example, if its value is  $\{1,'abc'\}$ , the integrated function is called with fun(1,'abc',t,y), the output function as outfun(1,'abc',tn,yn), and so on.

## **Examples**

### **Default options**

```
odeset
  AbsTol: 1e-6
  Events: []
  EventTime: []
  InitialStep: []
  MaxStep: []
  MinStep: []
  NormControl: false
  OnEvent: []
  OutputFcn: []
  PreArg: {}
  Refine: []
  RelTol: 1e-3
  Stats: false
```

## Option 'refine'

ode45 is typically able to use large time steps to achieve the requested tolerance. When plotting the output, however, interpolating it with straight lines produces visual artifacts. This is why ode45 inserts 3 interpolated points for each calculated point, based on the fifth-order approximation calculated for the integration (Refine is 4 by default). In the following code, curves with and without interpolation are compared (see Fig. 6.2). Note that the numbers of evaluations of the function being integrated are the same.

```
mu = 1:
fun = @(t,y) [y(2); mu*(1-y(1)^2)*y(2)-y(1)];
(t, v) = ode45(fun, [0,5], [2;0], ...
               odeset('Refine',1,'Stats',true));
  Number of function evaluations: 289
 Successful steps: 42
 Failed steps (error too large): 6
size(v)
 43 2
(ti, yi) = ode45(fun, [0,5], [2;0], ...
                 odeset('Stats',true));
 Number of function evaluations: 289
 Successful steps: 42
  Failed steps (error too large): 6
size(yi)
  169 2
plot(ti', yi', 'g');
plot(t', y');
```

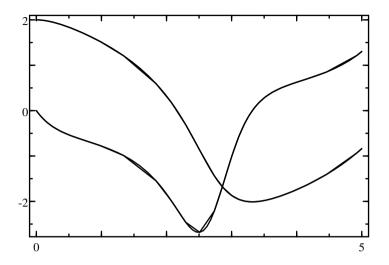


Figure 6.2 Van der Pol equation with Refine set to 1 and 4

#### State-based events

For simulating a ball bouncing on the ground, an event is generated every time the ball hits the ground, and its speed is changed instantaneously. Let y(1) be the height of the ball above the ground, and y(2) its speed (SI units are used). The state-space model is

```
y' = [y(2); -9.81];
```

An event occurs when the ball hits the ground:

```
value = y(1);
isterminal = false;
direction = -1;
```

When the event occurs, a new state is computed:

```
yn = [0; -damping*y(2)];
```

To integrate this, the following functions are defined:

```
function yp = ballfun(t, y, damping)
  yp = [y(2); -9.81];
function (v, te, d) = ballevents(t, y, damping)
  v = y(1);     // event when the height becomes negative
  te = false;     // do not terminate
  d = -1;     // only for negative speeds
function yn = ballonevent(t, y, i, damping)
  yn = [0; -damping*y(2)];
```

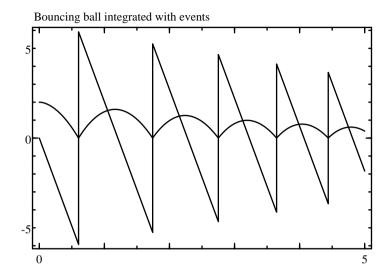


Figure 6.3 Bouncing ball integrated with events

Ball state is integrated during 5 s (see Fig. 6.3) with

#### Time events with discontinuous function

If the function being integrated has discontinuities at known time instants, option EventTime can be used to insure an accurate switching time. Consider a first-order filter with input u(t), where u(t)=0 for t<1 and u(t)=1 for  $t\geq 1$ . The following function is defined for the state derivative:

```
function yp = filterfun(t, y)
   yp = -y + (t <= 1 ? 0 : 1);

A single time event is generated at t = 1:
   opt = odeset('EventTime', [inf, 1]);
   (t, y) = ode45(@filterfun, [0, 5], 0, opt);
   plot(t', y');</pre>
```

Function filterfun is integrated in the normal way until t=1 inclusive, with u=0. This is why the conditional expression in filterfun is *less than or equal to* and not *less than*. Then the event occurs, and integration continues from  $t=1+\epsilon$  with u=0.

### Non-integrated state

For the last example, we will consider a system made of an integrator and a relay with hysteresis in a loop. Let y(1) be the output of the integrator and y(2) the output of the relay. Only y(1) is integrated:

```
yi' = y(2);
```

An event occurs when the integrator is larger or smaller than the hysteresis:

```
value = y(1) - y(2);
isTerminal = false;
direction = sign(y(2));
```

When the event occurs, a new value is computed for the 2nd state:

```
yn = -y(2);
```

To integrate this, the following functions are defined:

```
function yp = relayfun(t, y)
  yp = y(2);
function (v, te, d) = relayevents(t, y)
  v = y(1) - y(2);
  te = false;
  d = sign(y(2));
function yn = relayonevent(t, y, i)
  yn = -y(2);
```

The initial state is [0;1]; 0 for the integrator, and 1 for the output of the relay. State is integrated during 5 s (see Fig. 6.4) with

```
(t, y) = ode45(@relayfun, [0, 5], [0; 1], ...
odeset('Events', @relayevents, 'OnEvent', @relayonevent));
plot(t', y');
```

# Delay differential equation

A system whose Laplace transform is  $Y(s)/U(s) = e^{-ds}/(s^2 + s)$  (first order + integrator + delay d) is simulated with unit negative feedback. The reference signal is 1 for t > 0. First, the open-loop system is converted from transfer function to state-space, such that x'(t) = Ax(t) + Bu(t) and y(t) = Cx(t-d). The closed-loop state-space model is obtained by setting u(t) = 1 - y(t), which gives x'(t) = Ax(t) + BCx(t-d).

Delayed state is interpolated from past results with interp1. Note that values for t < 0 (extrapolated) are set to 0, and that values more recent than the last result are interpolated with the state passed to f for current t.

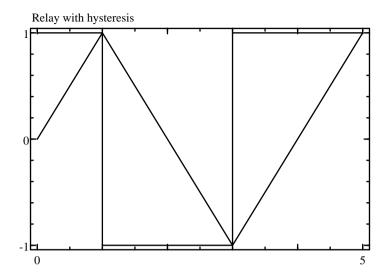


Figure 6.4 Relay with hysteresis integrated with events

ode23, ode45, optimset, interp1

# optimset

Options for minimization and zero finding.

# **Syntax**

```
options = optimset
options = optimset(name1=value1, ...)
options = optimset(name1, value1, ...)
options = optimset(options0, name1=value1, ...)
options = optimset(options0, name1, value1, ...)
```

# **Description**

optimset(name1,value1,...) creates the option argument used by fminbnd, fminsearch, fzero, fsolve, and other optimization functions. Options are specified with name/value pairs, where the name is a string which must match exactly the names in the table below. Case is significant. Alternatively, options can be given with named arguments. Options which are not specified have a default value. The result is a structure whose fields correspond to each option. Without any input argument, optimset creates a structure with all the default options. Note that fminbnd, fminsearch, and fzero also interpret the lack of an option argument, or the empty array [], as a request to use the default values. Options can also be passed directly to fminbnd and other similar functions as named arguments.

When its first input argument is a structure, optimset adds or changes fields which correspond to the name/value pairs which follow

Here is the list of permissible options (empty arrays mean "automatic"):

Name	Default	Meaning
Display	false	detailed display
MaxFunEvals	1000	maximum number of evaluations
MaxIter	500	maximum number of iterations
TolX	[]	maximum relative error

The default value of TolX is eps for fzero and sqrt(eps) for fminbnd and fminsearch.

#### **Examples**

Default options:

```
optimset
Display: false
MaxFunEvals: 1000
MaxIter: 500
TolX: []
```

Display of the steps performed to find the zero of  $\cos x$  between 1 and 2:

```
fzero(@cos, [1,2], optimset('Display',true))
  Checking lower bound
  Checking upper bound
  Inverse quadratic interpolation 2,1.5649,1
  Inverse quadratic interpolation 1.5649,1.571,2
  Inverse quadratic interpolation 1.571,1.5708,1.5649
  Inverse quadratic interpolation 1.5708,1.5708,1.571
```

```
Inverse quadratic interpolation 1.5708,1.5708,1.571
ans =
   1.5708
```

fzero, fminbnd, fminsearch, lsqnonlin, lsqcurvefit

# quad

Numerical integration.

## **Syntax**

```
y = quad(fun, a, b)
y = quad(fun, a, b, tol)
y = quad(fun, a, b, tol, trace)
y = quad(fun, a, b, tol, trace, ...)
```

# **Description**

quad(fun,a,b) integrates numerically real function fun between a and b. fun is either specified by its name or given as an anonymous or inline function or a function reference.

The optional fourth argument is the requested relative tolerance of the result. It is either a positive real scalar number or the empty matrix (or missing argument) for the default value, which is sqrt(eps). The optional fifth argument, if true or nonzero, makes quad displays information at each step.

Additional input arguments of quad are given as additional input arguments to function fun. They permit to parameterize the function.

## **Example**

$$\int_{0}^{2} t e^{-t} dt$$
 quad(@(t) t\*exp(-t), 0, 2) 0.5940

### Remark

Function quad is obsolete and should be replaced with integral, which supports named options and complex numbers.

integral, operator @

# 6.23 String Functions

## base32decode

Decode base32-encoded data.

## **Syntax**

strb = base32decode(strt)

# **Description**

base32decode(strt) decodes the contents of string strt which represents data encoded with base32. Characters which are not 'A'-'Z', '2'-'7', or '=' are ignored. Decoding stops at the end of the string or when '=' is reached.

#### See also

base32encode, base64decode

### base32encode

Encode data using base32.

### **Syntax**

strt = base32encode(strb)

# **Description**

base32encode(strb) encodes the contents of string strb which represents binary data. The result contains only characters 'A'-'Z' and '2'-'7', and linefeed every 56 characters. It is suitable for transmission or storage on media which accept only uppercase letters and digits, without '0' or '1' easy to misinterpret as letters.

Each character of encoded data represents 5 bits of binary data; i.e. one needs eight characters for five bytes. The five bits represent 32 different values, encoded with the characters 'A' to 'Z' and '2' to '7' in this order. When the binary data have a length which is not a

multiple of 5, encoded data are padded with 2, 3, 5 or 6 characters '=' to have a multiple of 8.

Base32 encoding is an Internet standard described in RFC 4648.

# **Example**

```
s = base32encode(char(0:10))
s =
    AAAQEAYEAUDAOCAJBI======
d = double(base32decode(s))
d =
    0 1 2 3 4 5 6 7 8 9 10
```

### See also

base32decode, base64encode

### base64decode

Decode base64-encoded data.

# **Syntax**

```
strb = base64decode(strt)
```

# **Description**

base64decode(strt) decodes the contents of string strt which represents data encoded with base64. Characters which are not 'A'-'Z', 'a'-'z', '0'-'9', '+', '/', or '=' are ignored. Decoding stops at the end of the string or when '=' is reached.

### See also

base64encode, base32decode

# base64encode

Encode data using base64.

# Syntax

```
strt = base64encode(strb)
```

# **Description**

base64encode(strb) encodes the contents of string strb which represents binary data. The result contains only characters 'A'-'Z', 'a'-'z', '0'-'9', '+', '/', and '='; and linefeed every 60 characters. It is suitable for transmission or storage on media which accept only text.

Each character of encoded data represents 6 bits of binary data; i.e. one needs four characters for three bytes. The six bits represent 64 different values, encoded with the characters 'A' to 'Z', 'a' to 'z', '0' to '9', '+', and '/' in this order. When the binary data have a length which is not a multiple of 3, encoded data are padded with one or two characters '=' to have a multiple of 4.

Base64 encoding is an Internet standard described in RFC 2045.

### **Example**

```
s = base64encode(char(0:10))
s =
    AAECAwQFBgcICQo=
double(base64decode(s))
0 1 2 3 4 5 6 7 8 9 10
```

### See also

base64decode. base32encode

### char

Convert an array to a character array (string).

# **Syntax**

```
s = char(A)
S = char(s1, s2, ...)
```

# **Description**

char(A) converts the elements of matrix A to characters, resulting in a string of the same size. Characters are stored in unsigned 16-bit words. The shape of A is preserved. Even if most functions ignore the string shape, you can force a row vector with char(A(:).').

char(s1,s2,...) concatenates vertically the arrays given as arguments to produce a string matrix. If the strings do not have the same number of columns, blanks are added to the right.

# **Examples**

```
char(65:70)
  ABCDEF
char([65, 66; 67, 68](:).')
  ABCD
char('ab','cde')
  ab
  cde
char('abc',['de';'fg'])
  abc
  de
  fg
```

### See also

setstr, uint16, operator :, operator .', ischar, logical, double,
single

### deblank

Remove trailing blank characters from a string.

# **Syntax**

```
s2 = deblank(s1)
```

# **Description**

deblank(s1) removes the trailing blank characters from string s1. Blank characters are spaces (code 32), tabulators (code 9), carriage returns (code 13), line feeds (code 10), and null characters (code 0).

# **Example**

```
double(' \tAB CD\r\n\0')
  32  9 65 66 32 32 67 68 13 10 0
double(deblank(' \tAB CD\n\r\0')))
  32  9 65 66 32 32 67 68
```

### See also

strtrim

### hmac

HMAC authentication hash.

# **Syntax**

```
hash = hmac(hashtype, key, data)
hash = hmac(hashtype, key, data, type=t)
```

# Description

hmac (hashtype, key, data) calculates the authentication hash of data with secret key key and the method specified by hashtype: 'md5', 'sha1', 'sha224', 'sha256', 'sha384', or 'sha512'. Both arguments data and key can be strings (char arrays) which are converted to UTF-8, or int8 or uint8 arrays. The key can be up to 64 bytes; longer keys are truncated. The result is a string of hexadecimal digits whose length depends on the hash method, from 32 for HMAC-MD5 to 128 for HMAC-SHA512.

Named argument type can change the output type. It can be 'uint8' for an uint8 array of 16 or 20 bytes (raw HMAC-MD5 or HMAC-SHA1 hash result), 'hex' for its representation as a string of 32 or 40 hexadecimal digits (default), or base64 for its conversion to Base64 in a string of 24 or 28 characters.

HMAC is an Internet standard described in RFC 2104.

# **Examples**

HMAC-MD5 of 'Authenticated message' using secret key 'secret':

```
hmac('md5', 'secret', 'Authenticated message')
4f557b1f67bc4790e6e9568e2f458cf0
```

Same result computed explicitly, with the notations of RFC 2104: B is the block length, L is the hash length (16 for HMAC-MD5 or 20 for HMAC-SHA1), K is the key padded with zeros to have size B, and H is the hash function, defined here to produce a uint8 hash instead of an hexadecimal string like the LME functions md5 or sha1.

```
B = 64;
L = 16;
H = @(a) uint8(sscanf(md5(a), '%2x')');
key = uint8('secret');
data = uint8('Authenticated message');
K = [key, zeros(1, B - length(key), 'uint8')];
hash = H([bitxor(K, 0x5cuint8), H([bitxor(K, 0x36uint8), data])]);
sprintf('%.2x', hash)
```

Simple implementation of the HOTP and TOTP password algorithms (RFC 4226 and 6238) often used for two-factor authentication, with their default parameter values. The password is assumed to be base32-encoded.

```
function n = hotp(pass, cnt)
  k = uint8(base32decode(pass));
  c = bwrite(cnt, 'uint64;b');
  // or c=bwrite([floor(c/2^32),mod(c,2^32)],'uint32;b');
  hs = hmac('sha1', k, c, type='uint8');
  ob = mod(hs(20), 16);
  dt = mod(sread(hs(ob + (1:4)), [], 'uint32;b'), 2^31);
  n = mod(dt, 1e6);
function n = totp(pass)
  t = floor(posixtime / 30);
  n = hotp(pass, t);
```

Simple implementation of the PBKDF2 key stretching algorithm (RFC 2898):

```
function dk = pbkdf2_hmac(hashtype, p, salt, c, dkLen)
  hLen = length(hmac(hashtype, '', '')) / 2;
  dk = uint8([]);
  for i = 1:ceil(dkLen / hLen)
    u = hmac(hashtype, p, [salt, bwrite(i, 'uint32;b')], type='uint8');
    f = u;
    for j = 2:c
        u = hmac(hashtype, p, u, type='uint8');
        f = bitxor(f, u);
    end
        dk = [dk, f];
  end
    dk = dk(1:dkLen);
```

Test of PBKDF2-HMAC-SHA1 with values provided in RFC 6070 (output format is switched to hexadecimal for easier comparison):

```
format int x
pbkdf2_hmac_sha1('sha1', 'password', 'salt', 4096, 20)
    0x4b    0x0    0x79    0x1    0xb7    0x65    0x48    0x9a    0xbe    0xad
    0x49    0xd9    0x26    0xf7    0x21    0xd0    0x65    0xa4    0x29    0xc1
format
```

### See also

md5, sha1

### ischar

Test for a string object.

### **Syntax**

```
b = ischar(obj)
```

# **Description**

ischar(obj) is true if the object obj is a character string, false otherwise. Strings can have more than one line.

# **Examples**

```
ischar('abc')
  true
ischar(0)
  false
ischar([])
  false
ischar('')
  true
ischar(['abc';'def'])
  true
```

### See also

isletter, isspace, isnumeric, islogical, isinteger, islist,
isstruct, setstr, char

# isdigit

Test for decimal digit characters.

# **Syntax**

```
b = isdigit(s)
```

# **Description**

For each character of string s, isdigit(s) is true if it is a digit ('0' to '9') and false otherwise. The result is a logical array with the same size as the input argument.

# **Examples**

```
isdigit('a123bAB12*')
FTTTFFFTTFF
```

### See also

isletter, isspace, lower, upper, ischar

### isletter

Test for letter characters.

# **Syntax**

```
b = isletter(s)
```

# **Description**

For each character of string s, isletter(s) is true if it is an ASCII letter (a-z or A-Z) and false otherwise. The result is a logical array with the same size as the input argument.

isletter gives false for letters outside the 7-bit ASCII range; unicodeclass should be used for Unicode-aware tests.

# **Examples**

```
isletter('abAB12*')
  T T T F F F F
```

### See also

isdigit, isspace, lower, upper, ischar, unicodeclass

# isspace

Test for space characters.

# **Syntax**

```
b = isspace(s)
```

# **Description**

For each character of string s, isspace(s) is true if it is a space, a tabulator, a carriage return or a line feed, and false otherwise. The result is a logical array with the same size as the input argument.

# **Example**

```
isspace('a\tb c\nd')
  F T F T F T F
```

### See also

isletter, isdigit, ischar

### latex2mathml

Convert LaTeX equation to MathML.

### **Syntax**

```
str = latex2mathml(tex)
str = latex2mathml(tex, mml1, mml2, ...)
str = latex2mathml(..., displaymath=b)
```

# **Description**

latex2mathml(tex) converts LaTeX equation in string tex to MathML. LaTeX equations may be enclosed between dollars or double-dollars, but this is not mandatory. In string literals, backslash and tick characters must be escaped as \\ and \' respectively.

With additional arguments, which must be strings containing MathML, parameters #1, #2, ... in argument tex are converted to argument i+1.

The following LaTeX features are supported:

- variables (each letter is a separate variable)
- numbers (sequences of digit and dot characters)
- superscripts and subscripts, prime (single or multiple)
- braces used to group subexpressions or specify arguments with more than one token
- operators (+, -, comma, semicolon, etc.)
- control sequences for character definitions, with greek characters in lower case (\alpha, ..., \omega, \varepsilon, \vartheta, \varphi) and upper case (\Alpha, ..., \Omega), arrows (\leftarrow or \gets, \rightarrow or \to, \uparrow, \downarrow, \leftrightarrow, \updownarrow, \Leftarrow, \Rightarrow, \Uparrow, \Downarrow, \Leftrightarrow, \Updownarrow, nwarrow, nearrow, searrow, swarrow, mapsto, hookleftarrow. hookrightarrow. Longleftrightarrow. longmapsto), and symbols (\|, \ell, \partial, \infty, \emptyset, \nabla, \perp, \angle, \triangle, \backslash,
  \forall, \exists, \flat, \natural, \sharp, \pm, \mp, \cdot, \times, \star, \diamond, \cap, \cup, etc.)

- \not followed by comparison operator, such as \not< or \not\approx
- control sequences for function definitions (\arccos, \arcsin, \arctan, \arg, \cos, \cosh, \cot, \coth, \csc, \deg, \det, \dim, \exp, \gcd, \hom, \inf, \injlim, \ker, \lg, \liminf, \limsup, \ln, \log, \max, \min, \Pr, \projlim, \sec, \sin, \sinh, \sup, \tan, \tanh)
- accents (\hat, \check, \tilde, \acute, grave, \dot, \ddot, 
  \dddot, breve, \bar, \vec, \overline, \widehat, \widetilde, 
  \underline)
- \left and \right
- fractions with \frac or \over
- roots with \sqrt (without optional radix) or \root...\of...
- \atop
- large operators (\bigcap, \bigcup, \bigodot, \bigoplus, \bigotimes, \bigsqcup, \biguplus, \bigvee, \bigwedge, \coprod, \prod, and \sum with implicit \limits for limits below and above the symbol; and \int, \iint, \iiint, \iiint, \iiint, \oint, and \oiint with implicit \nolimits for limits to the right of the symbol)
- \limits and \nolimits for functions and large operators
- matrices with \matrix, \pmatrix, \bmatrix, \bmatrix, \vmatrix, \vmatrix, \begin{array}{...}.../\end{array};
  values are separated with & and rows with \cr or \\
- font selection with \rm for roman, \bf for bold face, and \mit for math italic
- color with \color{c} where c is black, red, green, blue, cyan, magenta, yellow, white, orange, violet, purple, brown, darkgray, gray, or lightgray
- hidden element with \phantom
- text with \hbox{...} (brace contents is taken verbatim)
- horizontal spaces with \, \: \: \; \quad \qquad and \!

LaTeX features not enumerated above, such as definitions and nested text and equations, are not supported.

latex2mathml has also features which are missing in LaTeX. Unicode is used for both LaTeX input and MathML output. Some semantics is recognized to build subexpressions which are revealed in the

resulting MathML. For instance, in x+(y+z)w, (y+z) is a subpexpressions; so is (y+z)w with an implicit multiplication (resulting in the <mo>&it;<mo> MathML operator), used as the second operand of the addition. LaTeX code (like mathematical notation) is sometimes ambiguous and is not always converted to the expected MathML (e.g. a(b+c) is converted to a function call while the same notation could mean the product of a and b+c), but this should not have any visible effect when the MathML is typeset.

Operators can be used as freely as in LaTeX. Missing operands result in <none/>, as if there were an empty pair of braces {}. Consecutive terms are joined with implicit multiplications.

Named argument displaymath specifies whether the vertical space is tight, like in inline equations surrounded by text (false), or unconstrained, as rendered in separate lines (true). It affects the position of some limits. The default is true.

# **Examples**

```
 \begin{array}{l} latex2mathml('xy^2') \\ <mrow><mi>x</mi><mo>&it;</mo><msup><mi>y</mi><mn>2</mn></mrow></mrow><mml = latex2mathml('\\frac\{x_3+5\}\{x_1+x_2\}');\\ mml = latex2mathml('\\root n \\of x$');\\ mml = latex2mathml('\\pmatrix\{x & \\sqrt y \\cr \\sin\\phi & \\hat\\ell\}');\\ mml = latex2mathml('\\dot x = #1', mathml([1,2;3,0], false));\\ mml = latex2mathml('\\lim_{x \\rightarrow 0} f(x)', displaymath=true)\\ mml = latex2mathml('\\lim_{x \\rightarrow 0} f(x)', displaymath=false)\\ \end{array}
```

### See also

mathml

### lower

Convert all uppercase letters to lowercase.

# **Syntax**

```
s2 = lower(s1)
```

# **Description**

lower(s1) converts all the uppercase letters of string s1 to lowercase, according to the Unicode Character Database.

# **Example**

```
lower('abcABC123')
  abcabc123
```

### See also

```
upper, isletter
```

### mathml

Conversion to MathML.

# **Syntax**

```
str = mathml(x)
str = mathml(x, false)
str = mathml(..., Format=f, NPrec=n)
```

# **Description**

mathml(x) converts its argument x to MathML presentation, returned as a string.

By default, the MathML top-level element is <math>. If the result is to be used as a MathML subelement of a larger equation, a second input argument equal to the logical value false can be specified to suppress <math>.

By default, mathml converts numbers like format '%g' of sprintf. Named arguments can override them: format is a single letter format recognized by sprintf and NPrec is the precision (number of decimals).

# **Example**

### See also

mathmlpoly, latex2mathml, sprintf

# mathmlpoly

Conversion of a polynomial to MathML.

# **Syntax**

```
str = mathmlpoly(pol)
str = mathmlpoly(pol, var)
str = mathmlpoly(..., power)
str = mathmlpoly(..., false)
str = mathmlpoly(..., Format=f, NPrec=n)
```

# **Description**

mathmlpoly(coef) converts polynomial coefficients pol to MathML presentation, returned as a string. The polynomial is given as a vector of coefficients, with the highest power first; e.g.,  $x^2 + 2x - 3$  is represented by [1,2,-3].

By default, the name of the variable is x. An optional second argument can specify another name as a string, such as 'y', or a MathML fragment beginning with a less-than character, such as '<mn>3</mn>'.

Powers can be specified explicitly with an additional argument, a vector which must have the same length as the polynomial coefficients. Negative and fractional numbers are allowed; the imaginary part, if any, is ignored.

By default, the MathML top-level element is <math>. If the result is to be used as a MathML subelement of a larger equation, an additional input argument (the last unnamed argument) equal to the logical value false can be specified to suppress <math>.

Named arguments format and NPrec have the same effect as with mathml.

# **Examples**

Simple third-order polynomial:

```
mathmlpoly([1,2,5,3])
```

Polynomial with negative powers of variable q:

```
c = [1, 2.3, 4.5, -2];
mathmlpoly(c, 'q', -(0:numel(c)-1))
```

### Rational fraction:

```
str = sprintf('<mfrac>%s%s</mfrac>',
  mathmlpoly(num, false),
  mathmlpoly(den, false));
```

### See also

mathml

# md5

Calculate MD5 digest.

# **Syntax**

```
digest = md5(strb)
digest = md5(fd)
digest = md5(..., type=t)
```

# **Description**

md5(strb) calculates the MD5 digest of strb which represents binary data. strb can be a string (only the least-significant byte of each character is considered) or an array of bytes of class uint8 or int8. The result is a string of 32 hexadecimal digits. It is believed to be hard to create the input to get a given digest, or to create two inputs with the same digest.

md5(fd) calculates the MD5 digest of the bytes read from file descriptor fd until the end of the file. The file is left open.

Named argument type can change the output type. It can be 'uint8' for an uint8 array of 16 bytes (raw MD5 hash result), 'hex' for its representation as a string of 32 hexadecimal digits (default), or base64 for its conversion to Base64 in a string of 24 characters.

MD5 digest is an Internet standard described in RFC 1321.

# **Examples**

```
MD5 of the three characters 'a', 'b', and 'c':

md5('abc')
900150983cd24fb0d6963f7d28e17f72
```

This can be compared to the result of the command tool md5 found on many unix systems:

```
$ echo -n abc | md5
900150983cd24fb0d6963f7d28e17f72
```

The following statements calculate the digest of the file 'somefile':

```
fd = fopen('somefile');
digest = md5(fd);
fclose(fd);
```

### See also

shal. hmac

# regexp regexpi

Regular expression match.

# **Syntax**

```
(startIx, endIx, length, grExt) = regexp(str, re)
(startIx, endIx, grExt) = regexpi(str, re)
```

# **Description**

regexp(str,re) matches regular expression re in string str. A regular expression is a string which contains meta-characters to match classes of characters, repetitions and alternatives, as described below.

Once a match is found, the remaining part of str is parsed from the end of the previous match to find more matches. The result of regexp is an array of start indices in str and an array of corresponding end indices. Empty matches have a length endIx-startIx-1=0.

The third output argument, if present, is set to a list whose items correspond to matches. Items are arrays of size 2-by-ng. Each row corresponds to a group, i.e. a subexpression in parentheses in the regular expression; the first column contains the index of the first character in str and the second column contains the index of the last character.

regexpi is similar to regexp, except that letter case is ignored. The following regular expression elements are recognized:

# **Any character other than those described below** Literal match.

```
. (dot) Any character.
\0 Nul (0).
\t Tab (9).
\n Newline (10).
\v Vertical tab (11).
\f Form feed (12).
\r Carriage return (13).
\P where P is one of \()[]{}?*+/ P
```

- \xNN Character whose code is NN in hexadecimal.
- \uNNNN Character whose code is NNNN in hexadecimal.
- [...] Any of the characters in brackets. Characters can be enumarated (e.g. [ax2] to match a, x or 2), provided as ranges with a hyphen (e.g. [a-c] to match a, b or c) or any combination. Caret ^ must not appear first; closing bracket ] must appear first; and hyphen must not be used in a way which could be interpreted as a range.
- [^...] Any character not enumated in brackets (e.g. [^a-z] for any character except for lowercase letters).
- AB Catenation of A and B.
- A|B One of A or B. | has the lowest priority: ab|c matches ab or c.
- A? A (if possible) or nothing.
- A\* As many repetitions of A as possible, including none.
- A+ As many repetitions of A as possible, at least one.
- A{n} Exactly n repetitions of A.
- $A\{n_i\}$  At least n repetitions of A (as many as possible).
- $A\{n,m\}$  Between n and m repetitions of A (as many as possible).
- A?? Nothing (if possible) or A.
- A\*? As few repetitions of A as possible, including none.
- A+? As few repetitions of A as possible, at least one.
- $A\{n,\}$ ? At least n repetitions of A (as few as possible).
- A $\{n,m\}$ ? Between n and m repetitions of A (as few as possible).
- A?+, A\*?, A++, A{...}+ Possessive repetitions: as many as possible, but once the maximum number has been found, does not try less repetitions should the remaining part of the regular expression fail to match anything.
- (A) Group; matches subexpression A, which is captured for further reference as \N.
- (?:A) Group without capture; just matches subexpression A.
- \N where N is a digit from 1 to 9 Character substring which was matched by the N:th group delimited by parentheses.
- ^ Matches beginning of string.

- \$ Matches end of string.
- \b Beginning or end of word.
- (?=A) Positive lookahead: succeeds if what follows matches A without consuming A.
- (?!A) Negative lookahead: succeeds if what follows does not match A without consuming A.
- (?# comment (ignored).
- \d Digit (can be used inside or outside brackets).
- \D Not a digit (can be used inside or outside brackets).
- \s White space (can be used inside or outside brackets).
- \S Not white space (can be used inside or outside brackets).
- \w Alphanumeric (can be used inside or outside brackets).
- \W Not alphanumeric (can be used inside or outside brackets).
- [:alnum:] Same as A-Za-z0-9 (must be used inside brackets, e.g.
  [[:alnum:]])
- [:alpha:] Same as A-Za-z (must be used inside brackets, e.g.
   [[:alpha:]])
- [:blank:] Same as \x20\x09, i.e. space or tab (must be used inside brackets, e.g. [[:blank:]])
- [:cntrl:] Same as \0-\x1f (must be used inside brackets, e.g.
   [[:cntrl:]])
- [:digit:] Same as 0-9 (must be used inside brackets, e.g.
   [[:digit:]])
- [:graph:] Same as \x21-\x7e, i.e. ASCII characters without space and control characters (must be used inside brackets, e.g. [[:graph:]])
- [:lower:] Same as a-z (must be used inside brackets, e.g.
   [^[:lower:][:digit:]] which is equivalent to [^a-z0-9])
- [:print:] Same as \x20-\x7e, i.e. ASCII characters without control characters (must be used inside brackets, e.g. [[:print:]])
- [:punct:] Same as !"#\$%&'()\*+,-./:;<=>?@[\]^\_'{|}~ (must be used inside brackets, e.g. [[:punct:]])
- [:space:] Same as \x20\x09\x0a\x0c\x0d or \s (must be used inside brackets, e.g. [[:space:]])

```
[:upper:]
            Same as A-Z (must be used inside brackets, e.g.
 [[:upper:]])
[:word:] Same as [:alnum:]_ (must be used inside brackets, e.g.
 [[:word:]])
[:xdigit:] Same as 0-9A-Fa-f (must be used inside brackets.
 e.g. [[:xdigit:]])
```

Quantifiers ?, \* and +, and their lazy and possessive versions (suffixed with ? or + respectively) have the highest priority. Priority can be changed with parentheses, e.g. (abc)\* or (albc)d.

# **Examples**

startIx = 1 4 endIx =3 11

Simple match without metacharacter:

```
(startIx, endIx) = regexp('Some random string', 'om')
   startIx =
      2 10
   endTx =
      3 11
Dot to match any character:
 regexp('Some random string', 'S..e')
Anchor to end of string:
 regexp('Some random string', '..$')
   17
Repetition:
 regexp('Some random string', 'r.*m')
   6
By default, repetitions are greedy (as many as possible):
 (startIx, endIx) = regexp('Some random string', '.*m')
   startIx =
      1
   endIx =
     11
Lazy repetition (as few as possible):
 (startIx, endIx) = regexp('Some random string', '.*?m')
```

Possessive repetitions keep the largest number of repetitions which provides a match regardless of subsequent failures:

Since backslash is an escape character in LME strings, it must be escaped itself:

```
(startIx, endIx) = regexp('Some random string', '\\b\\w.+?\\b')
startIx =
    1  6  13
endIx =
    4  11  18
```

Reference to a captured group:

```
(startIx, endIx) = regexp('xx-ab-ab', '(.+)-\\1')
startIx =
   4
endIx =
   8
```

Positive lookahead to find words followed by a colon without picking the colon itself:

```
(startIx, endIx) = regexp('mailto:foo@example.com', '\b\\w+(?=:)')
    startIx =
        1
    endIx =
        6
```

Group (the extent of the whole match is ignored using placeholder output arguments ~):

```
(~, ~, grExt) = regexp('Regexp are fun', '\\b(\\w+)\\s+(\\w+)\\b');
grExt{1}
    1    6
    8    10
    12    14
```

Match ignoring case:

```
regexpi('Some random string', 'some')
1
```

Case-explicit character classes are still case-significant, but character enumerations or ranges are not:

```
regexpi('Some random string', '^[[:lower:]]')
[]
regexpi('Some random string', '^[a-z]')
1
```

### See also

strfind. strtok

### setstr

Conversion of an array to a string.

# **Syntax**

```
str = setstr(A)
```

# **Description**

setstr(A) converts the elements of array A to characters, resulting in a string of the same size. Characters are stored in unsigned 16-bit words.

# **Example**

```
setstr(65:75)
ABCDEFGHIJK
```

### See also

char, uint16, logical, double

# sha1 sha2

Calculate SHA-1 or SHA-2 digest.

### Syntax

```
digest = sha1(strb)
digest = sha1(fd)
digest = sha1(..., type=t)
digest = sha2(...)
digest = sha2(..., variant=v)
```

# **Description**

shal(strb) calculates the SHA-1 digest of strb which represents binary data. strb can be a string (only the least-significant byte of each character is considered) or an array of bytes of class uint8 or int8. The result is a string of 40 hexadecimal digits. It is believed to be hard to create the input to get a given digest, or to create two inputs with the same digest.

shal(fd) calculates the SHA-1 digest of the bytes read from file descriptor fd until the end of the file. The file is left open.

Named argument type can change the output type. It can be 'uint8' for an uint8 array of 20 bytes (raw SHA-1 hash result), 'hex' for its representation as a string of 40 hexadecimal digits (default), or base64 for its conversion to Base64 in a string of 28 characters.

SHA-1 digest is an Internet standard described in RFC 3174.

sha2 calculates the SHA-256 digest, a 256-bit variant of the SHA-2 hash algorithm. Its arguments are the same as those of sha1. In addition, named argument variant can specify one of the supported SHA-2 variants: 224, 256 (default), 384, or 512.

# **Example**

```
SHA-1 digest of the three characters 'a', 'b', and 'c':

sha1('abc')
    a9993e364706816aba3e25717850c26c9cd0d89d

SHA-224 digest of the empty message '':

sha2('', variant=224)
    d14a028c2a3a2bc9476102bb288234c415a2b01f828ea62ac5b3e42f
```

### See also

md5. hmac

# split

Split a string.

# **Syntax**

```
L = split(string, separator)
```

# **Description**

split(string, separator) finds substrings of string separated by separator and return them as a list. Empty substring are discarded. sepatator is a string of one or more characters.

# **Examples**

```
split('abc;de;f', ';')
   {'abc', 'de', 'f'}
split('++a+++b+++','++')
   {'a', '+b', '+'}
```

### See also

strfind

# strcmp

String comparison.

# **Syntax**

```
b = strcmp(s1, s2)
b = strcmp(s1, s2, n)
```

# **Description**

strcmp(s1, s2) is true if the strings s1 and s2 are equal (i.e. same length and corresponding characters are equal). strcmp(s1, s2, n) compares the strings up to the n:th character. Note that this function does not return the same result as the strcmp function of the standard C library.

# **Examples**

```
strcmp('abc','abc')
  true
strcmp('abc','def')
  false
strcmp('abc','abd',2)
  true
```

```
strcmp('abc','abd',5)
false
```

### See also

strcmpi, operator ===, operator ==, strfind, strmatch

# strcmpi

String comparison with ignoring letter case.

# **Syntax**

```
b = strcmpi(s1, s2)
b = strcmpi(s1, s2, n)
```

# **Description**

strcmpi compares strings for equality, ignoring letter case. In every other respect, it behaves like strcmp.

# **Examples**

```
strcmpi('abc','aBc')
  true
strcmpi('Abc','abd',2)
  true
```

### See also

strcmp, operator ===, operator ==, strfind, strmatch

### strfind

Find a substring in a string.

# **Syntax**

```
pos = strfind(str, sub)
```

# **Description**

strfind(str,sub) finds occurrences of string sub in string str and returns a vector of the positions of all occurrences, or the empty vector [] if there is none. Occurrences may overlap.

### **Examples**

```
strfind('ababcdbaaab','ab')
  1 3 10
strfind('ababcdbaaab','ac')
  []
strfind('aaaaaaa','aaa')
  1 2 3
```

### See also

find, strcmp, strrep, split, strmatch, strtok

### strmatch

String match.

# **Syntax**

```
i = strmatch(str, strMatrix)
i = strmatch(str, strList)
i = strmatch(..., 'exact')
```

# **Description**

strmatch(str,strMatrix) compares string str with each row of the character matrix strMatrix; it returns the index of the first row whose beginning is equal to str, or 0 if no match is found. Case is significant.

strmatch(str,strList) compares string str with each element of list or cell array strList, which must be strings.

With a third argument, which must be the string 'exact', str must match the complete row or element of the second argument, not only the beginning.

# **Examples**

```
strmatch('abc',['axyz';'uabc';'abcd';'efgh'])
3
strmatch('abc',['axyz';'uabc';'abcd';'efgh'],'exact')
0
strmatch('abc',{'ABC','axyz','abcdefg','ab','abcd'})
3
```

### See also

```
strcmp, strfind
```

# strrep

Replace a substring in a string.

# **Syntax**

```
newstr = strrep(str, sub, repl)
```

# Description

strrep(str, sub, repl) replaces all occurrences of string sub in string str with string repl.

# **Examples**

```
strrep('ababcdbaaab','ab','X')
  'XXcdbaaX'
strrep('aaaaaaaa','aaa','12345')
  '1234512345a'
```

### See also

strfind, strcmp, strmatch, strtok

### strtok

Token search in string.

# **Syntax**

```
(token, remainder) = strtok(str)
(token, remainder) = strtok(str, separators)
```

# **Description**

strtok(str) gives the first token in string str. A token is defined as a substring delimited by separators or by the beginning or end of the string; by default, separators are spaces, tabulators, carriage returns and line feeds. If no token is found (i.e. if str is empty or contains only separator characters), the result is the empty string.

The optional second output is set to what follows immediately the token, including separators. If no token is found, it is the same as str.

An optional second input argument contains the separators in a string.

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# **Examples**

Strings are displayed with quotes to show clearly the separators.

```
strtok(' ab cde ')
   'ab'
(t, r) = strtok(' ab cde ')
   t =
        'ab'
   r =
        ' cde '
(t, r) = strtok('2, 5, 3')
   t =
        '2'
   r =
        ', 5, 3'
```

### See also

strmatch, strfind, strtrim

### strtrim

Remove leading and trailing blank characters from a string.

# **Syntax**

```
s2 = strtrim(s1)
```

# **Description**

strtrim(s1) removes the leading and trailing blank characters from string s1. Blank characters are spaces (code 32), tabulators (code 9), carriage returns (code 13), line feeds (code 10), and null characters (code 0).

# Example

```
double(' \tAB CD\r\n\0')
  32  9 65 66 32 32 67 68 13 10 0
double(strtrim(' \tAB CD\n\r\0')))
  65 66 32 32 67 68
```

### See also

deblank, strtok

### unicodeclass

Unicode character class.

### **Syntax**

cls = unicodeclass(c)

# **Description**

unicodeclass(c) gives the Unicode character class (General\_Category property in the Unicode Character Database) of its argument c, which must be a single-character string. The result is one of the following two-character strings:

Class	Description	Class	Description
'Lu'	Letter, Uppercase	'Pi'	Punctuation, Initial qupte
'Ll'	Letter, Lowercase	'Pf'	Punctuation, Final Quote
'Lt'	Letter, Titlecase	'Po'	Punctuation, Other
'Lm'	Letter, Modifier	'Sm'	Symbol, Math
'Lo'	Letter, Other	'Sc'	Symbol, Currency
'Mn'	Mark, Non-Spcacing	'Sk'	Symbol, Modifier
'Mc'	Mark, Spacing Combining	'So'	Symbol, Other
'Me'	Mark, Enclosing	'Zs'	Separator, Spcace
'Nd'	Number, Decimal Digit	'Zl'	Separator, Line
'Nl'	Number, Letter	'Zp'	Separator, Paragraph
'Noʻ	Number, Other	'Cc'	Other, Control
'Pc'	Punctuation, Connector	'Cf'	Other, Format
'Pd'	Punctuation, Dash	'Cs'	Other, Surrogate
'Ps'	Punctuation, Open	'Co'	Other, Private Use
'Pe'	Punctuation, Close	'Cn'	Other, Not Assigned

### See also

isletter, isdigit, isspace

# upper

Convert all lowercase letters to lowercase.

# Syntax

```
s2 = upper(s1)
```

# **Description**

upper(s1) converts all the lowercase letters of string s1 to uppercase, according to the Unicode Character Database.

# **Example**

```
upper('abcABC123')
  ABCABC123
```

### See also

lower, isletter

### utf32decode

Decode Unicode characters encoded with UTF-32.

# **Syntax**

```
str = utf32decode(b)
```

# **Description**

utf32decode(b) decodes the contents of uint32 or int32 array b which represents Unicode characters encoded with UTF-32 (basically, Unicode code point). The result is a standard character array with a single row, usually encoded with UTF-16. Invalid codes are ignored.

If all the codes in b correspond to the Basic Multilingual Plane (16-bits, and not surrogate 0xd800-0xdfff), the result is equivalent to char(b).

### See also

utf32encode. utf8decode

### utf32encode

Encode a string of Unicode characters using UTF-32.

# **Syntax**

```
b = utf32encode(str)
```

# **Description**

utf32encode(str) encodes the contents of character array str using UTF-32. Each Unicode character in str, made of 1 or 2 UTF-16 words, corresponds to one UTF-32 code. The result is an array of unsigned 32-bit integers.

If all the characters in str correspond to the Basic Multilingual Plane (16-bits, and no surrogate pairs), the result is equivalent to uint32(str).

# **Examples**

```
utf32encode('abc')
  1x3 uint32 array
    97  98  99
str = utf32decode(65872uint32);
double(str)
    55296  56656
utf32encode(str)
  65872uint32
```

### See also

utf32decode, utf8encode

# utf8decode

Decode Unicode characters encoded with UTF-8.

# **Syntax**

```
str = utf8decode(b)
```

# **Description**

utf8decode(b) decodes the contents of uint8 or int8 array b which represents Unicode characters encoded with UTF-8. Each Unicode character corresponds to up to 4 bytes of UTF-8 code. The result is a standard character array with a single row; characters are usually encoded as UTF-16, with 1 or 2 words per character. Invalid codes (for example when the beginning of the decoded data does not correspond to a character boundary) are ignored.

### See also

utf8encode, utf32decode

### utf8encode

Encode a string of Unicode characters using UTF-8.

# **Syntax**

```
b = utf8encode(str)
```

# Description

utf8encode(str) encodes the contents of character array str using UTF-8. Each Unicode character in str corresponds to up to 4 bytes of UTF-8 code. The result is an array of unsigned 8-bit integers.

If the input string does not contain Unicode characters, the output is invalid.

# **Example**

```
b = utf8encode(['abc', 200, 2000, 20000])
b =
    1x10 uint8 array
    97   98   99   195  136  223  144  228  184  160
str = utf8decode(b);
double(str)
    97    98    99   200  2000  20000
```

### See also

utf8decode, utf32encode

# 6.24 Quaternions

Quaternion functions support scalar and arrays of quaternions. Basic arithmetic operators and functions are overloaded to support expressions with the same syntax as for numbers and matrices.

Quaternions are numbers similar to complex numbers, but with four components instead of two. The unit imaginary parts are named i, j, and k. A quaternion can be written w + ix + jy + kz. The following relationships hold:

$$i^2 = j^2 = k^2 = ijk = -1$$

It follows that the product of two quaternions is not commutative; for instance, ij = k but ji = -k.

Quaternions are convenient to represent arbitrary rotations in the 3d space. They are more compact than matrices and are easier to

normalize. This makes them suitable to simulation and control of mechanical systems and vehicles, such as flight simulators and robotics.

Functions below are specific to quaternions:

Function	Purpose
isquaternion	test for quaternion type
q2mat	conversion to rotation matrix
q2rpy	conversion to attitude angles
q2str	conversion to string
qimag	imaginary parts
qinv	element-wise inverse
qnorm	scalar norm
qslerp	spherical linear interpolation
quaternion	quaternion creation
rpy2q	conversion from attitude angles

Operators below accept quaternions as arguments:

<b>Function</b> ctranspose	Operator ,	<b>Purpose</b> conjugate transpose
eq	==	element-wise equality
horzcat	[,]	horizontal array concatenation
ldivide	. \	left division
ne	~=`	element-wise inequality
minus	-	difference
mldivide	\	matrix left division
mrdivide	/	matrix right division
mtimes	*	matrix multiplication
plus	+	addition
rdivide	./	division
times	.*	multiplication
transpose	. ′	transpose
uminus	-	unary minus
uplus	+	unary plus
vertcat	[;]	vertical array concatenation

Most of these operators work as expected, like with complex scalars and matrices. Multiplication and left/right division are not commutative. Matrix operations are not supported: operators \*, /,  $^$  and  $^$  are defined as a convenience (they are equivalent to .\*, ./, ., and  $.^$  respectively) and work only element-wise with scalar arguments.

Mathematical functions below accept quaternions as arguments; with arrays of quaternions, they are applied to each element separately.

Function	Purpose
abs	absolute value
conj	conjugate
cos	cosine
exp	exponential
log	natural logarithm
real	real part
sign	quaternion sign (normalization)
sin	sine
sqrt	square root

Functions below performs computations on arrays of quaternions.

Function	Purpose
cumsum	cumulative sum
diff	differences
double	conversion to array of double
mean	arithmetic mean
sum	sum

Functions below are related to array size.

Function	Purpose
beginning	first subscript
cat	array concatenation
end	last subscript
flipdim	flip array
fliplr	flip left-right
flipud	flip upside-down
ipermute	dimension inverse permutation
isempty	test for empty array
length	length of vector
ndims	number of dimensions
numel	number of elements
permute	dimension permutation
repmat	array replication
reshape	array reshaping
rot90	array rotation
size	array size
squeeze	remove singleton dimensions

Finally, functions below are related to output and assignment.

# FunctionPurposedispdisplaydumpvarconversion to stringsubsasgnassignment to subarrays or to quaternion partssubsrefreference to subarrays or to quaternion parts

Function imag is replaced with qimag which gives a quaternion with the real part set to zero, because there are three imaginary components instead of one with complex numbers.

Operators and functions which accept multiple arguments convert automatically double arrays to quaternions, ignoring the imaginary part of complex numbers.

Conversion to numeric arrays with double adds a dimension for the real part and the three imaginary parts. For example, converting a scalar quaternion gives a 4-by-1 double column vector and converting a 2-by-2 quaternion array gives a 2-by-2-by-4 double array. Real and imaginary components can be accessed with the field access notation: q.w is the real part of q, q.x, q.y, and q.z are its imaginary parts, and q.v is its imaginary parts as an array similar to the result of double but without the real part.

Compatibility note: native functions for quaternions replace library quaternion which defined quaternion scalars and matrices. It is much faster and supports arrays of more than two dimensions; on the other hand, matrix-oriented functions are not supported anymore, and the result of dumpvar is not directly compatible.

# isquaternion

Test for a quaternion.

# **Syntax**

```
b = isquaternion(q)
```

# **Description**

isquaternion(q) is true if the input argument is a quaternion and false otherwise.

# **Examples**

```
isquaternion(2)
  false
isquaternion(quaternion(2))
  true
```

### See also

quaternion, isnumeric

# q2mat

Conversion from quaternion to rotation matrix.

# **Syntax**

```
R = q2mat(q)
```

# **Description**

R=q2mat(q) gives the 3x3 orthogonal matrix R corresponding to the rotation given by scalar quaternion q. For a vector a=[x;y;z] and its representation as a pure quaternion aq=quaternion(x,y,z), the rotation can be performed with quaternion multiplication bq=q\*aq/q or matrix multiplication b=R\*a.

Input argument q does not have to be normalized; a quaternion corresponding to a given rotation is defined up to a multiplicative factor.

# **Example**

```
q = rpy2q(0.1, 0.3, 0.2);
R = q2mat(q)
R =
    0.9363 -0.1688   0.3080
    0.1898   0.9810   0.0954
    -0.2955   0.0954   0.9506
aq = quaternion(1, 2, 3);
q * aq / q
    1.5228i+2.0336j+2.7469k
a = [1; 2; 3];
R * a
    1.5228
    2.4380
    2.7469
```

### See also

q2rpy, rpy2q, quaternion

# q2rpy

Conversion from quaternion to attitude angles.

# **Syntax**

```
(pitch, roll, yaw) = q2rpy(q)
```

# **Description**

q2rpy(q) gives the pitch, roll, and yaw angles corresponding to the rotation given by quaternion q. It is the inverse of rpy2q. All angles are given in radians.

If the input argument is a quaternion array, the results are arrays of the same size; conversion from quaternion to angles is performed independently on corresponding elements.

### See also

rpy2q, q2mat, quaternion

# q2str

Conversion from quaternion to string.

# **Syntax**

```
str = q2str(q)
```

# **Description**

q2str(q) converts quaternion q to its string representation, with the same format as disp.

### See also

quaternion, format

# qimag

Quaternion imaginary parts.

### **Syntax**

```
b = qimag(q)
```

# **Description**

qimag(q) gives the imaginary parts of quaternion q as a quaternion, i.e. the same quaternion where the real part is set to zero. real(q) gives the real part of quaternion q as a double number.

### **Example**

```
q = quaternion(1,2,3,4)
  q =
        1+2i+3j+4k
real(q)
  1
qimag(q)
  2i+3j+4k
```

### See also

quaternion

# qinv

Quaternion element-wise inverse.

# **Syntax**

```
b = qinv(q)
```

# **Description**

qinv(q) gives the inverse of quaternion q. If its input argument is a quaternion array, the result is an quaternion array of the same size whose elements are the inverse of the corresponding elements of the input.

The inverse of a normalized quaternion is its conjugate.

# Example

```
q = quaternion(0.4,0.1,0.2,0.2)
q =
    0.4+0.1i+0.2j+0.2k
p = qinv(q)
p =
    1.6-0.4i-0.8j-0.8k
abs(q)
    0.5
```

```
abs(p)
```

### See also

quaternion, qnorm, conj

# qnorm

Quaternion scalar norm.

# **Syntax**

```
n = qnorm(q)
```

# **Description**

qnorm(q) gives the norm of quaternion q, i.e. the sum of squares of its components, or the square of its absolute value. If q is an array of quaternions, qnorm gives a double array of the same size where each element is the norm of the corresponding element of q.

### See also

quaternion, abs

# qslerp

Quaternion spherical linear interpolation.

# **Syntax**

```
q = qslerp(q1, q2, t)
```

# Description

qslerp(q1,q2,t) performs spherical linear interpolation between quaternions q1 and q2. The result is on the smallest great circle arc defined by normalized q1 and q2 for values of real number t between 0 and 1.

If q1 or q2 is 0, the result is NaN. If they are opposite, the great circle arc going through 1, or 1i, is picked.

If input arguments are arrays of compatible size (same size or scalar), the result is a quaternion array of the same size; conversion

from angles to quaternion is performed independently on corresponding elements.

#### **Example**

```
q = qslerp(1, rpy2q(0, 1, -1.5), [0, 0.33, 0.66, 1]);
(roll, pitch, yaw) = q2rpy(q)
roll =
    0.0000   0.1843   0.2272   0.0000
pitch =
    0.0000   0.3081   0.6636   1.0000
yaw =
    0.0000   -0.4261   -0.8605   -1.5000
```

#### See also

quaternion, rpy2q, q2rpy

### quaternion

Ouaternion creation.

## **Syntax**

```
q = quaternion
q = quaternion(w)
q = quaternion(c)
q = quaternion(x, y, z)
q = quaternion(w, x, y, z)
q = quaternion(w, v)
```

## **Description**

With a real argument, quaternion(x) creates a quaternion object whose real part is w and imaginary parts are 0. With a complex argument, quaternion(c) creates the quaternion object real(c)+i\*imag(c).

With four real arguments, quaternion(w,x,y,z) creates the quaternion object w+i\*x+j\*y+k\*z.

With three real arguments, quaternion(x,y,z) creates the pure quaternion object i\*x+j\*y+k\*z.

In all these cases, the arguments may be scalars or arrays of the same size.

With two arguments, quaternion (w, v) creates a quaternion object whose real part is w and imaginary parts is array v. v must have one more dimension than w for the three imaginary parts.

Without argument, quaternion returns the zero quaternion object. The real or imaginary parts of a quaternion can be accessed with field access, such as q.w, q.x, q.y, q.z, and q.v.

#### **Examples**

```
q = quaternion(1, 2, 3, 4)
    1+2i+3j+4k
a + 5
  6+2i+3j+4k
q * q
  -28+4i+6i+8k
Q = [q, 2; 2*q, 5]
  2x2 quaternion array
Q.y
  3 0
  6 0
q = quaternion(1, [5; 3; 7])
    1+5i+3j+7k
q.v
  5
  3
  7
```

#### See also

real, qimaq, q2str, rpy2q

# rpy2q

Conversion from attitude angles to quaternion.

# **Syntax**

```
q = rpy2q(pitch, roll, yaw)
```

# Description

rpy2q(pitch,roll,yaw) gives the quaternion corresponding to a rotation of angle yaw around the z axis, followed by a rotation of angle pitch around the y axis, followed by a rotation of angle roll round the x axis. All angles are given in radians. The result is a normalized quaternion whose real part is  $\cos(9/2)$  and imaginary part  $\sin(9/2)(v_x i + v_y j + v_z k)$ , for a rotation of 9 around unit vector

 $\begin{bmatrix} v_x \ v_y \ v_z \end{bmatrix}^T$ . The rotation is applied to a point  $\begin{bmatrix} x \ y \ z \end{bmatrix}^T$  given as a pure quaternion a = xi + yj + zk, giving point a also as a pure quaternion; then b=q\*a/q and a=q\b\*q. The rotation can also be seen as changing coordinates from body to absolute, where the body's attitude is given by pitch, roll and yaw.

In order to have the usual meaning of pitch, roll and yaw, the x axis must be aligned with the direction of motion, the y axis with the lateral direction, and the z axis with the vertical direction, with the usual sign conventions for cross products. Two common choices are x pointing forward, y to the left, and z upward; or x forward, y to the right, and z downward.

If input arguments are arrays of compatible size (same size or scalar), the result is a quaternion array of the same size; conversion from angles to quaternion is performed independently on corresponding elements.

### **Example**

Conversion of two vectors from aircraft coordinates (x axis forward, y axis to the left, z axis upward) to earth coordinates (x directed to the north, y to the west, z to the zenith). In aircraft coordinates, vectors are [2;0;0] (propeller position) and [0;5;0] (left wing tip). The aircraft attitude has a pitch of 10 degrees upward, i.e. -10 degrees with the choice of axis, and null roll and yaw.

```
q = rpy2q(0, -10*pi/180, 0)
q =
    0.9962-0.0872j
q * quaternion(2, 0, 0) / q
    1.9696i+0.3473k
q * quaternion(0, 5, 0) / q
    5j
```

#### See also

q2rpy, q2mat, quaternion

# 6.25 List Functions

# apply

Function evaluation with arguments in lists.

```
listout = apply(fun, listin)
listout = apply(fun, listin, nargout)
listout = apply(fun, listin, na)
listout = apply(fun, listin, nargout, na)
```

### **Description**

listout=apply(fun, listin) evaluates function fun with input arguments taken from the elements of list listin. Output arguments are grouped in list listout. Function fun is specified by either its name as a string, a function reference, or an anonymous or inline function.

The number of expected output arguments can be specified with an optional third input argument nargout. By default, the maximum number of output arguments is requested, up to 256; this limit exists to prevent functions with an unlimited number of output arguments, such as deal, from filling memory.

With a 4th argument na (or 3rd if nargout is not specified), named arguments can be provided as a structure.

### **Examples**

```
apply(@size, {magic(3)}, 2)
  {3, 3}
apply(@(x,y) 2*x+3*y, {5, 10})
  {40}
```

The maximum number of output arguments of min is 2 (minimum value and its index):

```
apply(@min, {[8, 3, 4, 7]}) {3, 2}
```

Two equivalent ways of calling disp with a named argument fd to specify the standard error file descriptor 2:

```
disp(123, fd=2);
apply(@disp, {123}, 0, {fd=2});
```

#### See also

map, feval, inline, operator @, varargin, namedargin, varargout

# join

List concatenation.

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### **Syntax**

```
list = join(l1, l2, ...)
```

### **Description**

join(l1, l2, ...) joins elements of lists l1, l2, etc. to make a larger list.

### **Examples**

```
join({1,'a',2:5}, {4,2}, {{'xxx'}})
    {1,'a',[2,3,4,5],4,2,{'xxx'}}
join()
    {}
```

#### See also

```
operator ,, operator ;, replist
```

#### islist

Test for a list object.

## **Syntax**

```
b = islist(obj)
```

# **Description**

islist(obj) is true if the object obj is a list, false otherwise.

# **Examples**

```
islist({1, 2, 'x'})
   true
islist({})
   true
islist([])
   false
ischar('')
   false
```

#### See also

isstruct, isnumeric, ischar, islogical, isempty

#### list2num

Conversion from list to numeric array.

### **Syntax**

```
A = list2num(list)
```

### **Description**

list2num(list) takes the elements of list, which must be numbers or arrays, and concatenates them on a row (along second dimension) as if they were placed inside brackets and separated with commas. Element sizes must be compatible.

### **Example**

```
list2num({1, 2+3j, 4:6})
1 2+3j 4 5 6
```

#### See also

```
num2list, operator [], operator ,
```

### map

Function evaluation for each element of a list

# Syntax

```
(listout1,...) = map(fun, listin1, ...)
```

# **Description**

map(fun,listin1,...) evaluates function fun successively for each corresponding elements of the remaining arguments, which must be lists or cell arrays. It returns the result(s) of the evaluation as list(s) or cell array(s) with the same size as inputs. Input lists which contain a single element are repeated to match other arguments if necessary. Function fun is specified by either its name as a string, a function reference, or an anonymous or inline function.

### **Examples**

```
map('max', {[2,6,4], [7,-1], 1:100})
    {6, 7, 100}
map(@(x) x+10, {3,7,16})
    {13, 17, 26}
(nr, nc) = map(@size, {1,'abc',[4,7;3,4]})
    nr =
        {1,1,2}
    nc =
        {1,3,2}
s = map(@size, {1,'abc',[4,7;3,4]})
s =
        {[1,1], [1,3], [2,2]}
map(@disp, {'hello', 'lme'})
    hello
    lme
```

Lists with single elements are expanded to match the size of other lists. The following example computes atan2(1,2) and atan2(1,3):

```
map(@atan2, {1}, {2,3}) {0.4636,0.3218}
```

#### See also

apply, cellfun, for, inline, operator @

#### num2list

Conversion from array to list.

# Syntax

```
list = num2list(A)
list = num2list(A, dim)
```

# **Description**

num2list(A) creates a list with the elements of non-cell array A.
 num2list(A,dim) cuts array A along dimension dim and creates a
list with the result.

# **Examples**

```
num2list(1:5)
  {1, 2, 3, 4, 5}
num2list([1,2;3,4])
```

```
{1, 2, 3, 4}
num2list([1, 2; 3, 4], 1)
{[1, 2], [3, 4]}
num2list([1, 2; 3, 4], 2)
{[1; 3], [2; 4]}
```

#### See also

list2num, num2cell

# replist

Replicate a list.

## **Syntax**

```
listout = replist(listin, n)
```

## **Description**

replist(listin,n) makes a new list by concatenating n copies of list listin.

## Example

```
replist({1, 'abc'}, 3)
{1,'abc',1,'abc',1,'abc'}
```

#### See also

join, repmat

# 6.26 Structure Functions

## cell2struct

Convert a cell array to a structure array.

# Syntax

```
SA = cell2struct(CA, fields)
SA = cell2struct(CA, fields, dim)
```

### **Description**

cell2struct(CA,fields) converts a cell array to a structure array. The size of the result is size(SA)(2:end), where nf is the number of fields. Field SA(i1,i2,...).f of the result contains cell CA{j,i1,i2,...}, where f is field field{j}. Argument fields contains the field names as strings.

With a third input argument, cell2struct(CA, fields, dim) picks fields of each element along dimension dim. The size of the result is the size of CA where dimension dim is removed.

## **Examples**

```
SA = cell2struct({1, 'ab'; 2, 'cde'}, {'a', 'b'});
SA = cell2struct({1, 2; 'ab', 'cde'}, {'a', 'b'}, 2);
```

#### See also

struct2cell

#### fieldnames

List of fields of a structure.

### **Syntax**

```
fields = fieldnames(strct)
```

# **Description**

fieldnames(strct) returns the field names of structure strct as a list of strings.

## **Example**

```
fieldnames({a=1, b=1:5})
    {'a', 'b'}
```

#### See also

struct, isfield, orderfields, rmfield

# getfield

Value of a field in a structure.

```
value = getfield(strct, name)
```

### **Description**

getfield(strct,name) gets the value of field name in structure strct. It is an error if the field does not exist. getfield(s,'f') gives the same value as s.f. getfield is especially useful when the field name is not fixed, but is stored in a variable or is the result of an expression.

#### See also

```
operator ., struct, setfield, rmfield
```

### isfield

Test for the existence of a field in a structure.

### **Syntax**

```
b = isfield(strct, name)
```

## **Description**

isfield(strct, name) is true if the structure strct has a field whose name is the string name, false otherwise.

### **Examples**

```
isfield({a=1:3, x='abc'}, 'a')
    true
isfield({a=1:3, x='abc'}, 'A')
    false
```

#### See also

fieldnames, isstruct, struct

### isstruct

Test for a structure object.

```
b = isstruct(obj)
```

### **Description**

isstruct(obj) is true if its argument obj is a structure or structure array, false otherwise.

### **Examples**

```
isstruct({a=123})
   true
isstruct({1, 2, 'x'})
   false
a.f = 3;
isstruct(a)
   true
```

#### See also

struct, isfield, isa, islist, ischar, isobject, islogical

### orderfields

Reorders the fields of a structure.

# Syntax

```
strctout = orderfields(strctin)
strctout = orderfields(strctin, structref)
strctout = orderfields(strctin, names)
strctout = orderfields(strctin, perm)
(strctout, perm) = orderfields(...)
```

# Description

With a single input argument, orderfields(strctin) reorders structure fields by sorting them by field names.

With two input arguments, orderfields reorders the fields of the first argument after the second argument. Second argument can be a permutation vector containing integers from 1 to length(strctin), another structure with the same field names, or a list of names. In the last cases, all the fields of the structure must be present in the second argument.

The (first) output argument is a structure with the same fields and the same value as the first input argument; the only difference is the field order. An optional second output argument is set to the permutation vector.

### **Examples**

```
s = \{a=123, c=1:3, b='abcde'\}
    a: 123
    c: real 1x3
    b: 'abcde'
(t, p) = orderfields(s)
  t =
    a: 123
    b: 'abcde'
    c: real 1x3
  p =
    1
    3
t = orderfields(s, {'c', 'b', 'a'})
  t =
    c: real 1x3
    b: 'abcde'
    a: 123
```

#### See also

struct, fieldnames

#### rmfield

Deletion of a field in a structure.

# Syntax

```
strctout = rmfield(strctin, name)
```

# Description

strctout=rmfield(strctin, name) makes a structure strctout with the same fields as strctin, except for field named name which is removed. If field name does not exist, strctout is the same as strctin.

### **Example**

```
x = rmfield({a=1:3, b='abc'}, 'a');
fieldnames(x)
b
```

#### See also

struct, setfield, getfield, orderfields

#### setfield

Assignment to a field in a structure.

## Syntax

```
strctout = setfield(strctin, name, value)
```

### **Description**

strctout=setfield(strctin,name,value) makes a structure strctout with the same fields as strctin, except that field named name is added if it does not exist yet and is set to value. s=setfield(s,'f',v) has the same effect as s.f=v; s=setfield(s,str,v) has the same effect as s.(str)=v.

#### See also

```
operator ., struct, getfield, rmfield
```

#### struct

Creation of a structure

# Syntax

```
strct = struct(field1=value1, field2=value2, ...)
strct = struct(fieldname1, value1, fieldname2, value2, ...)
strct = {field1=value1, field2=value2, ...}
```

### **Description**

struct builds a new structure. With named arguments, the name of each argument is used as the field name. Otherwise, input arguments are used by pairs to create the fields; for each pair, the first argument is the field name, provided as a string, and the second one is the field value.

Instead of named arguments, a more compact notation consists in writing named values between braces. In that case, all values must be named; when no value has a name, a list is created, and mixed named and unnamed values are invalid. Fields are separated by commas; semicolons separate elements of n-by-1 struct arrays. See the documentation of braces for more details.

### **Examples**

Three equivalent ways to create a structure with two fields a and b:

```
x = {a=1, b=2:5};
x = struct(a=1, b=2:5);
x = struct('a', 1, 'b', 2:5);
x.a
   1
x.b
   2 3 4 5
```

#### See also

structarray, isstruct, isfield, rmfield, fieldnames, operator {}

### struct2cell

Convert a structure array to a cell array.

# **Syntax**

```
CA = struct2cell(SA)
```

# **Description**

struct2cell(SA) converts a structure or structure array to a cell array. The size of the result is [nf,size(SA)], where nf is the number of fields. Cell CA{j,i1,i2,...} of the result contains field SA(i1,i2,...).f, where f is the j:th field.

### **Example**

```
SA = cell2struct({1, 'ab'; 2, 'cde'}, {'a', 'b'});
CA = struct2cell(SA);
```

#### See also

cell2struct

## structarray

Create a structure array.

### **Syntax**

```
SA = structarray(field1=A1, field2=A2, ...)
SA = structarray(fieldname1, A1, fieldname2, A2, ...)
```

### **Description**

structarray builds a new structure array. With named arguments, the name of each argument is used as the field name, and the value is a cell array whose elements become the corresponding values in the result. Otherwise, input arguments are used by pairs to create the fields; for each pair, the first argument is the field name, provided as a string, and the second one is the field values as a cell array.

In both cases, all cell arrays must have the same size; the resulting structure array has the same size.

## Example

The following assignments produce the same result:

```
SA = structarray(a = \{1,2;3,4\}, b = \{'a', 1:3; 'def', true\});

SA = structarray('a', \{1,2;3,4\}, 'b', \{'a', 1:3; 'def', true\});
```

#### See also

```
struct, cell2struct
```

# structmerge

Merge the fields of two structures.

```
S = structmerge(S1, S2)
```

### **Description**

structmerge(S1,S2) merges the fields of S1 and S2, producing a new structure containing the fields of both input arguments. More precisely, to build the result, structmerge starts with S1; each field which also exists in S2 is set to the value in S2; and finally, fields in S2 which do not exist in S1 are added.

If S1 and/or S2 are structure arrays, they must have the same size or one of them must be a simple structure (size 1x1). The result is a structure array of the same size where each element is obtained separately from the corresponding elements of S1 and S2; a simple structure argument is reused as necessary.

#### **Examples**

```
S = structmerge({a=2}, {b=3})
S =
    a: 2
    b: 3
S = structmerge({a=1:3, b=4}, {a='AB', c=10})
S =
    a: 'AB'
    b: 4
    c: 10
```

#### See also

fieldnames, setfield, cat

# 6.27 Object Functions

#### class

Object creation.

### **Syntax**

```
object = class(strct, 'classname')
object = class(strct, 'classname', parent1, ...)
str = class(object)
```

### **Description**

class(strct,'classname') makes an object of the specified class with the data of structure strct. Object fields can be accessed only from methods of that class, i.e. functions whose name is classname::methodname. Objects must be created by the class constructor classname::classname.

class(strct,'classname',parent1,...) makes an object of the specified class which inherits fields and methods from one or several other object(s) parent1, ... Parent objects are inserted as additional fields in the object, with the same name as the class. Fields of parent objects cannot be directly accessed by the new object's methods, only by the parent's methods.

class(object) gives the class of object as a string. The table below gives the name of native types.

Class	Native type
double	real or complex double scalar or array
single	real or complex single scalar or array
int8/16/32/64	8/16/32/64-bit signed integer scalar or array
uint8/16/32/64	8/16/32/64-bit unsigned integer scalar or array
logical	logical scalar or array
char	character or character array
list	list
cell	cell array
struct	scalar structure
structarray	structure array
inline	inline function
funref	function reference
null	null value

### **Examples**

```
o1 = class({fld1=1, fld2=rand(4)}, 'c1');
o2 = class({fld3='abc'}, 'c2', o1);
class(o2)
  c2
```

#### See also

struct, inferiorto, superiorto, isa, isobject, methods

#### inferiorto

Set class precedence.

```
inferiorto('class1', ...)
```

### Description

Called in a constructor, inferiorto('class1',...) specifies that the class has a lower precedence than classes whose names are given as input arguments. Precedence is used when a function has object arguments of different classes: the method defined for the class with the highest precedence is called.

#### See also

superiorto, class

#### isa

Test for an object of a given class.

### **Syntax**

```
b = isa(object,'classname')
```

## **Description**

isa(object, 'classname') returns true of object is an object of class class, directly or by inheritance. In addition to the class names given by class, the following classes are supported:

```
Class cell list or cell array double, single or integer scalar or array double or single scalar or array integer integer scalar or array
```

## Example

```
isa(pi,'double')
  true
```

#### See also

class, isobject, methods

### isnull

Test for a null value.

### **Syntax**

```
b = isnull(a)
```

## **Description**

isnull(a) returns true if a is the null value created with null, or false for any value of any other type.

#### See also

class, null

# isobject

Test for an object.

# **Syntax**

```
b = isobject(a)
```

# **Description**

object(a) returns true if a is an object created with class.

#### See also

```
class, isa, isstruct
```

#### methods

List of methods for a class.

### **Syntax**

```
methods classname
list = methods('classname')
```

### **Description**

methods classname displays the list of methods defined for class classname. Inherited methods and private methods are ignored. With an output argument, methods gives produces a list of strings.

#### See also

class, info

#### null

Null value.

### **Syntax**

```
obj = null
```

## **Description**

null gives the only value of the null data type. It stands for the lack of any value. Null values can be tested with isnull or with equality or inequality operators == and  $\tilde{}=$ .

With an input argument, null(A) gives the null space of matrix A.

### **Examples**

```
n = null
n =
    null
isnull(n)
    true
n == null
    true
n ~= null
    false
class(n)
    null
```

#### See also

isnull, null (linear algebra)

# superclasses

Get list of superclasses.

```
list = superclasses(obj)
```

### **Description**

superclasses (obj) gives the list of the names of parent classes (superclasses) of the object obj. Parent classes are specified as additional arguments to class when the object is constructed.

### **Example**

```
use lti;
G = tf(1, [1, 2]);
class(G)
   tf
superclasses(G)
   {'lti'}
isa(G, 'lti')
   true
```

#### See also

class, isa

# 6.28 Logical Functions

#### all

Check whether all the elements are true.

# Syntax

```
v = all(A)
v = all(A,dim)
b = all(v)
```

# **Description**

all(A) performs a logical AND on the elements of the columns of array A, or the elements of a vector. If a second argument dim is provided, the operation is performed along that dimension.

all can be omitted if its result is used by if or while, because these statements consider an array to be true if all its elements are nonzero.

#### **Examples**

```
all([1,2,3] == 2)
  false
all([1,2,3] > 0)
  true
```

#### See also

any, operator &, bitall

### any

Check whether any element is true.

### **Syntax**

```
v = any(A)
v = any(A,dim)
b = any(v)
```

## **Description**

any (A) performs a logical OR on the elements of the columns of array A, or the elements of a vector. If a second argument dim is provided, the operation is performed along that dimension.

### **Examples**

```
any([1,2,3] == 2)
  true
any([1,2,3] > 5)
  false
```

#### See also

all, operator |, bitany

### bitall

Check whether all the corresponding bits are true.

```
v = bitall(A)
v = bitall(A,dim)
b = bitall(v)
```

### **Description**

bitall(A) performs a bitwise AND on the elements of the columns of array A, or the elements of a vector. If a second argument dim is provided, the operation is performed along that dimension. A can be a double or an integer array. For double arrays, bitall uses the 32 least-significant bits.

### **Examples**

```
bitall([5, 3])
  1
bitall([7uint8, 6uint8; 3uint8, 6uint8], 2)
  2x1 uint8 array
  6
  2
```

#### See also

bitany, all, bitand

#### bitand

Bitwise AND.

# **Syntax**

```
c = bitand(a, b)
```

# **Description**

Each bit of the result is the binary AND of the corresponding bits of the inputs.

The inputs can be scalar, arrays of the same size, or a scalar and an array. If the input arguments are of type double, so is the result, and the operation is performed on 32 bits.

### **Examples**

```
bitand(1,3)
  1
bitand(1:6,1)
  1 0 1 0 1 0
bitand(7uint8, 1234int16)
  2int16
```

#### See also

bitor, bitxor, bitall, bitget

# bitany

Check whether any of the corresponding bits is true.

### **Syntax**

```
v = bitany(A)
v = bitany(A,dim)
b = bitany(v)
```

### **Description**

bitany(A) performs a bitwise OR on the elements of the columns of array A, or the elements of a vector. If a second argument dim is provided, the operation is performed along that dimension. A can be a double or an integer array. For double arrays, bitany uses the 32 least-significant bits.

# **Examples**

```
bitany([5, 3])
  7
bitany([0uint8, 6uint8; 3uint8, 6uint8], 2)
  2x1 uint8 array
  6
  7
```

#### See also

bitall, any, bitor

# bitcmp

Bit complement (bitwise NOT).

```
b = bitcmp(i)
b = bitcmp(a, n)
```

### **Description**

bitcmp(i) gives the 1-complement (bitwise NOT) of the integer i.

bitcmp(a,n), where a is an integer or a double, gives the 1-complement of the n least-significant bits. The result has the same type as a.

The inputs can be scalar, arrays of the same size, or a scalar and an array. If a is of type double, so is the result, and the operation is performed on at most 32 bits.

### **Examples**

```
bitcmp(1,4)
   14
bitcmp(0, 1:8)
   1 3 7 15 31 63 127 255
bitcmp([Ouint8, luint8, 255uint8])
   1x3 uint8 array
   255 254 0
```

#### See also

bitxor, operator ~

# bitget

Bit extraction.

# **Syntax**

```
b = bitget(a, n)
```

# **Description**

bitget(a, n) gives the n:th bit of integer a. a can be an integer or a double. The result has the same type as a. n=1 corresponds to the least significant bit.

The inputs can be scalar, arrays of the same size, or a scalar and an array. If a is of type double, so is the result, and n is limited to 32.

#### **Examples**

```
bitget(123,5)
  1
bitget(7, 1:8)
  1 1 1 0 0 0 0 0
bitget(5uint8, 2)
  0uint8
```

#### See also

bitset, bitand, bitshift

#### bitor

Bitwise OR.

#### **Syntax**

```
c = bitor(a, b)
```

### **Description**

The input arguments are converted to 32-bit unsigned integers; each bit of the result is the binary OR of the corresponding bits of the inputs.

The inputs can be scalar, arrays of the same size, or a scalar and an array. If the input arguments are of type double, so is the result, and the operation is performed on 32 bits.

### **Examples**

```
bitor(1,2)
   3
bitor(1:6,1)
   1 3 3 5 5 7
bitor(7uint8, 1234int16)
   1239int16
```

#### See also

bitand, bitxor, bitany, bitget

#### bitset

Bit assignment.

```
b = bitset(a, n)
b = bitset(a, n, v)
```

### **Description**

bitset(a,n) sets the n:th bit of integer a to 1. a can be an integer or a double. The result has the same type as a. n=1 corresponds to the least significant bit. With 3 input arguments, bitset(a,n,v) sets the bit to 1 if v is nonzero, or clears it if v is zero.

The inputs can be scalar, arrays of the same size, or a mix of them. If a is of type double, so is the result, and n is limited to 32.

### **Examples**

```
bitset(123,10)
  635
bitset(123, 1, 0)
  122
bitset(7uint8, 1:8)
  1x8 uint8 array
  7  7  15  23  39  71  135
```

#### See also

bitget, bitand, bitor, bitxor, bitshift

#### bitshift

Bit shift.

### **Syntax**

```
b = bitshift(a, shift)
b = bitshift(a, shift, n)
```

# **Description**

The first input argument is converted to a 32-bit unsigned integer, and shifted by shift bits, to the left if shift is positive or to the right if it is negative. With a third argument n, only n bits are retained.

The inputs can be scalar, arrays of the same size, or a mix of both.

#### **Examples**

```
bitshift(1,3)
   8
bitshift(8, -2:2)
   2 4 8 16 32
bitshift(15, 0:3, 4)
   15 14 12 8
```

#### See also

bitget

### bitxor

Bitwise exclusive OR.

### **Syntax**

```
c = bitxor(a, b)
```

### **Description**

The input arguments are converted to 32-bit unsigned integers; each bit of the result is the binary exclusive OR of the corresponding bits of the inputs.

The inputs can be scalar, arrays of the same size, or a scalar and an array.

### **Examples**

```
bitxor(1,3)
  2
bitxor(1:6,1)
  0 3 2 5 4 7
bitxor(7uint8, 1234int16)
  1237int16
```

#### See also

bitcmp, bitand, bitor, bitget

### false

Boolean constant false.

```
b = false
B = false(n)
B = false(n1, n2, ...)
B = false([n1, n2, ...])
```

### **Description**

The boolean constant false can be used to set the value of a variable. It is equivalent to logical(0). The constant 0 is equivalent in many cases; indices (to get or set the elements of an array) are an important exception.

With input arguments, false builds a logical array whose elements are false. The size of the array is specified by one integer for a square matrix, or several integers (either as separate arguments or in a vector) for an array of any size.

### **Examples**

```
false
  false
islogical(false)
  true
false(2,3)
  F F F
F F
```

#### See also

```
true, logical, zeros
```

# graycode

Conversion to Gray code.

## **Syntax**

```
g = graycode(n)
```

# **Description**

graycode(n) converts the integer number n to Gray code. The argument n can be an integer number of class double (converted to an unsigned integer) or any integer type. If it is an array, conversion is

performed on each element. The result has the same type and size as the input.

Gray code is an encoding which maps each integer of s bits to another integer of s bits, such that two consecutive codes (i.e. graycode(n) and graycode(n+1) for any n) have only one bit which differs.

### **Example**

```
graycode(0:7)
0 1 3 2 6 7 5 4
```

#### See also

igraycode

# igraycode

Conversion from Gray code.

### **Syntax**

```
n = igraycode(g)
```

# **Description**

igraycode(n) converts the Gray code g to the corresponding integer. It is the inverse of graycode. The argument n can be an integer number of class double (converted to an unsigned integer) or any integer type. If it is an array, conversion is performed on each element. The result has the same type and size as the input.

## **Example**

```
igraycode(graycode(0:7))
    0 1 2 3 4 5 6 7
```

#### See also

graycode

# islogical

Test for a boolean object.

```
b = islogical(obj)
```

### **Description**

islogical(obj) is true if obj is a logical value, and false otherwise. The result is always a scalar, even if obj is an array. Logical values are obtained with comparison operators, logical operators, test functions, and the function logical.

### **Examples**

```
islogical(eye(10))
  false
islogical(~eye(10))
  true
```

#### See also

logical, isnumeric, isinteger, ischar

# logical

Transform a number into a boolean.

## **Syntax**

```
B = logical(A)
```

# **Description**

logical(x) converts array or number A to logical (boolean) type. All nonzero elements of A are converted to true, and zero elements to false.

Logical values are stored as 0 for false or 1 for true in unsigned 8-bit integers. They differ from the uint8 type when they are used to select the elements of an array or list.

### **Examples**

```
a=1:3; a([1,0,1])
Index out of range
a=1:3; a(logical([1,0,1]))
    1 3
```

#### See also

```
islogical, uint8, double, char, setstr, operator ()
```

#### true

Boolean constant true.

### **Syntax**

```
b = true
B = true(n)
B = true(n1, n2, ...)
B = true([n1, n2, ...])
```

### **Description**

The boolean constant true can be used to set the value of a variable. It is equivalent to logical(1). The constant 1 is equivalent in many cases; indices (to get or set the elements of an array) are an important exception.

With input arguments, true builds a logical array whose elements are true. The size of the array is specified by one integer for a square matrix, or several integers (either as separate arguments or in a vector) for an array of any size.

## **Examples**

```
true
   true
islogical(true)
   true
true(2)
   T T
   T T
```

#### See also

```
false, logical, ones
```

#### xor

Exclusive or.

```
b3 = xor(b1,b2)
```

### **Description**

xor(b1,b2) performs the exclusive or operation between the corresponding elements of b1 and b2. b1 and b2 must have the same size or one of them must be a scalar.

### **Examples**

```
xor([false false true true],[false true false true])
  F T T F
xor(pi,8)
  false
```

#### See also

operator &, operator |

# **6.29 Dynamical System Functions**

This section describes functions related to linear time-invariant dynamical systems.

#### c2dm

Continuous-to-discrete-time conversion.

### **Syntax**

```
(numd,dend) = c2dm(numc,denc,Ts)
dend = c2dm(numc,denc,Ts)
(numd,dend) = c2dm(numc,denc,Ts,method)
dend = c2dm(numc,denc,Ts,method)
(Ad,Bd,Cd,Dd) = c2dm(Ac,Bc,Cc,Dc,Ts,method)
```

# **Description**

(numd,dend) = c2dm(numc,denc,Ts) converts the continuous-time transfer function numc/denc to a discrete-time transfer function numd/dend with sampling period Ts. The continuous-time transfer function is given by two polynomials in s, and the discrete-time

transfer function is given by two polynomials in z, all as vectors of coefficients with highest powers first.

c2dm(numc,denc,Ts,method) uses the specified conversion
method.method is one of

```
'zoh' or 'z' zero-order hold (default)
'foh' or 'f' first-order hold
'tustin' or 't' Tustin (bilinear transformation)
'matched' or 'm' Matched zeros, poles and gain
```

The input and output arguments numc, denc, numd, and dend can also be matrices; in that case, the conversion is applied separately on each row with the same sampling period Ts.

c2dm(Ac,Bc,Cc,Dc,Ts,method) performs the conversion from continuous-time state-space model (Ac,Bc,Cc,Dc) to discrete-time state-space model (Ad,Bd,Cd,Dd), defined by

$$\frac{dx}{dt}(t) = A_c x(t) + B_c u(t)$$

$$y(t) = C_c x(t) + D_c u(t)$$

and

$$x(k+1) = A_dx(k) + B_du(k)$$
  
$$y(k) = C_dx(k) + D_du(k)$$

Method 'matched' is not supported for state-space models.

#### **Examples**

```
(numd, dend) = c2dm(1, [1, 1], 0.1)
  numd =
    0.0952
dend =
    1 -0.9048
(numd, dend) = c2dm(1, [1, 1], 0.1, 'foh')
  numd =
    0.0484
dend =
    1 -0.9048
(numd, dend) = c2dm(1, [1, 1], 0.1, 'tustin')
  numd =
    0.0476  0.0476
dend =
    1 -0.9048
```

#### See also

d2cm

#### d2cm

Discrete-to-continuous-time conversion.

### **Syntax**

```
(numc,denc) = d2cm(numd,dend,Ts)
denc = d2cm(numd,dend,Ts)
(numc,denc) = d2cm(numd,dend,Ts,method)
denc = d2cm(numd,dend,Ts,method)
```

### **Description**

(numc,denc) = d2cm(numd,dend,Ts,method) converts the discrete-time transfer function numd/dend with sampling period Ts to a continuous-time transfer function numc/denc. The continuous-time transfer function is given by two polynomials in s, and the discrete-time transfer function is given by two polynomials in z, all as vectors of coefficients with highest powers first.

Method is

```
'tustin' or 't' Tustin (bilinear transformation) (default)
```

The input and output arguments nume, denc, numd, and dend can also be matrices; in that case, the conversion is applied separately on each row with the same sampling period Ts.

d2cm(Ad,Bd,Cd,Dd,Ts,method) performs the conversion from discrete-time state-space model (Ad,Bd,Cd,Dd) to continuous-time state-space model (Ac,Bc,Cc,Dc), defined by

$$x(k+1) = A_dx(k) + B_du(k)$$
  
$$y(k) = C_dx(k) + D_du(k)$$

and

$$\frac{dx}{dt}(t) = A_c x(t) + B_c u(t)$$

$$y(t) = C_c x(t) + D_c u(t)$$

#### **Example**

```
(numd, dend) = c2dm(1, [1, 1], 5, 't')
numd =
    0.7143  0.7143
dend =
    1  0.4286
(numc, denc) = d2cm(numd, dend)
numc =
    -3.8858e-17  1
denc =
    1  1
```

#### See also

c2dm

# dmargin

Robustness margins of a discrete-time system.

## **Syntax**

```
(gm,psi,wc,wx) = dmargin(num,den,Ts)
(gm,psi,wc,wx) = dmargin(num,den)
```

## **Description**

The open-loop discrete-time transfer function is given by the two polynomials num and den, with sampling period Ts (default value is 1). If the closed-loop system (with negative feedback) is unstable, all output arguments are set to an empty matrix. Otherwise, dmargin calculates the gain margins gm, which give the interval of gain for which the closed-loop system remains stable; the phase margin psi, always positive if it exists, which defines the symmetric range of phases which can be added to the open-loop system while keeping the closed-loop system stable; the critical frequency associated to the gain margins, where the open-loop frequency response intersects the real axis around -1; and the cross-over frequency associated to the phase margin, where the open-loop frequency response has a unit magnitude. If the Nyquist diagram does not cross the unit circle, psi and wx are empty.

## **Examples**

Stable closed-loop, Nyquist inside unit circle:

```
(gm,psi,wc,wx) = dmargin(0.005,poly([0.9,0.9]))
  gm = [-2, 38]
  psi = []
  wc = [0, 0.4510]
  wx = []
```

Stable closed-loop, Nyquist crosses unit circle:

```
(gm,psi,wc,wx) = dmargin(0.05,poly([0.9,0.9]))

gm = [-0.2, 3.8]

psi = 0.7105

wc = [0, 0.4510]

wx = 0.2112
```

Unstable closed-loop:

```
(gm,psi,wc,wx) = dmargin(1,poly([0.9,0.9]))
  gm = []
  psi = []
  wc = []
  wx = []
```

#### **Caveats**

Contrary to many functions, dmargin cannot be used with several transfer functions simultaneously, because not all of them may correspond simultaneously to either stable or unstable closed-loop systems.

#### See also

margin

# margin

Robustness margins of a continuous-time system.

# **Syntax**

```
(gm,psi,wc,wx) = margin(num,den)
```

# Description

The open-loop continuous-time transfer function is given by the two polynomials num and den. If the closed-loop system (with negative feedback) is unstable, all output arguments are set to an empty matrix. Otherwise, margin calculates the gain margins qm, which give the

interval of gain for which the closed-loop system remains stable; the phase margin psi, always positive if it exists, which defines the symmetric range of phases which can be added to the open-loop system while keeping the closed-loop system stable; the critical frequency associated to the gain margins, where the open-loop frequency response intersects the real axis around -1; and the cross-over frequency associated to the phase margin, where the open-loop frequency response has a unit magnitude. If the Nyquist diagram does not cross the unit circle, psi and wx are empty.

## **Examples**

Stable closed-loop, Nyquist inside unit circle:

```
(gm,psi,wc,wx) = margin(0.5,poly([-1,-1,-1]))
  gm = [-2, 16]
  psi = []
  wc = [0, 1.7321]
  wx = []
```

Stable closed-loop, Nyquist crosses unit circle:

```
(gm,psi,wc,wx) = margin(4,poly([-1,-1,-1]))
  gm = [-0.25 2]
  psi = 0.4737
  wc = [0, 1.7321]
  wx = 1.2328
```

Unstable closed-loop:

```
(gm,psi,wc,wx) = margin(10,poly([-1,-1,-1]))
  gm = []
  psi = []
  wc = []
  wx = []
```

#### **Caveats**

Contrary to many functions, margin cannot be used with several transfer functions simultaneously, because not all of them may correspond simultaneously to either stable or unstable closed-loop systems.

#### See also

dmargin

#### movezero

Change the position of a real or complex zero in a real-coefficient polynomial.

#### **Syntax**

```
pol2 = movezero(pol1, p0, p1)
(pol2, p1actual) = movezero(pol1, p0, p1)
```

## **Description**

movezero should be used in the mousedrag handle when the user drags the zero of a polynomial with real coefficients. It insures a consistent user experience.

If p0 is a real or complex zero of the polynomial pol1, movezero computes a new polynomial pol2, with real coefficients, a zero at p1, and most other zeros unchanged. If p0 and p1 are real,

```
pol2 = conv(deconv(pol1, [1, -p0]), [1, -p1])
```

If p0 and p1 are complex and their imaginary part has the same sign,

Otherwise, a real pole p0 is moved to complex pole p1 if imag(p1) > 0 and there is another real pole in pol0. A complex pole p0 can be moved to real(p1) if imag(p1)\*imag(p0) < 0; in that case, conj(p0) is moved to real(p0).

If it exists, the second output argument is set to the actual value of the displaced pole. It can be used to provide feedback to the user during the drag.

## **Examples**

```
roots(movezero(poly([1,3,2+3j,2-3j]),1,5))
5
3
2+3j
2-3j
roots(movezero(poly([1,3,2+3j,2-3j]),1,2j))
2j
-2j
2+3j
2-3j
roots(movezero(poly([1,3,2+3j,2-3j]),2+3j,5+8j))
1
```

```
3
5+8j
5-8j
roots(movezero(poly([1,3,2+3j,2-3j]),2+3j,5-8j))
1
3
5
2
(pol, newPole) = movezero(poly([1,3,2+3j,2-3j]),2+3j,5-8j);
newPole
5
```

#### See also

roots, conv, deconv

#### ss2tf

Conversion from state space to transfer function.

## **Syntax**

```
(num,den) = ss2tf(A,B,C,D)
den = ss2tf(A,B,C,D)
(num,den) = ss2tf(A,B,C,D,iu)
den = ss2tf(A,B,C,D,iu)
```

# Description

A continuous-time linear time-invariant system can be represented by the state-space model

$$\frac{dx}{dt}(t) = Ax(t) + Bu(t)$$

$$y(t) = Cx(t) + Du(t)$$

where x(t) is the state, u(t) the input, y(t) the output, and ABCD four constant matrices which characterize the model. If it is a single-input single-output system, it can also be represented by its transfer function num/den. (num,den) = ss2tf(A,B,C,D) converts the model from state space to transfer function. If the state-space model has multiple outputs, num is a matrix whose lines correspond to each output (the denominator is the same for all outputs). If the state-space model has multiple inputs, a fifth input argument is required and specifies which one to consider.

For a sampled-time model, exactly the same function can be used. The derivative is replaced by a forward shift, and the variable s of the Laplace transform is replaced by the variable z of the z transform. But as long as coefficients are concerned, the conversion is the same.

The degree of the denominator is equal to the number of states, i.e. the size of A. The degree of the numerator is equal to the number of states if D is not zero, and one less if D is zero.

If D is zero, it can be replaced by the empty matrix [].

## **Example**

Conversion from the state-space model dx/dt = -x + u, y = x to the transfer function Y(s)/U(s) = 1/(s+1):

```
(num, den) = ss2tf(-1, 1, 1, 0)
  num =
    1
  den =
    1 1
```

#### See also

tf2ss

#### tf2ss

Conversion from transfer function to state space.

# **Syntax**

```
(A,B,C,D) = tf2ss(num,den)
```

# **Description**

tf2ss(num,den) returns the state-space representation of the transfer function num/den, which is given as two polynomials. The transfer function must be causal, i.e. num must not have more columns than den. Systems with several outputs are specified by a num having one row per output; the denominator den must be the same for all the outputs.

tf2ss applies to continuous-time systems (Laplace transform) as well as to discrete-time systems (z transform or delta transform).

#### **Example**

```
(A,B,C,D) = tf2ss([2,5],[2,3,8])
A =
-1.5 -4
1 0
B =
1
0
C =
1 2.5
D =
0
```

#### See also

ss2tf, zp2ss

## zp2ss

Conversion from transfer function given by zeros and poles to state space.

## **Syntax**

```
(A,B,C,D) = zp2ss(z,p,k)
```

# **Description**

zp2ss(z,p,k) returns the state-space representation of the transfer function with zeros z, poles p and gain k (ratio of leading coefficients of numerator and denominator in decreasing powers). The transfer function must be causal, i.e. the number of zeros must not be larger than the number of poles. zp2ss supports only systems with one input and one output. Complex zeros and complex poles must make complex conjugate pairs, so that the corresponding polynomials have real coefficients.

zp2ss applies to continuous-time systems (Laplace transform) as well as to discrete-time systems (z transform or delta transform).

## **Example**

```
(A,B,C,D) = zp2ss([1;2],[3;4-1j;4+1j],5)

A =

8 -17    1

1   0   0

0   0   3
```

```
B = 0 0 1 C = 25 -75 5 D = 0
```

#### See also

tf2ss

# 6.30 Input/Output Functions

#### **bwrite**

Store data in an array of bytes.

## **Syntax**

```
s = bwrite(data)
s = bwrite(data, precision)
```

## **Description**

bwrite(data) stores the contents of the matrix data into an array of class uint8. The second parameter is the precision, whose meaning is the same as for fread. Its default value is 'uint8'.

# **Examples**

```
bwrite(12345, 'uint32;l')
  1x4 uint8 array
    57   48    0    0
bwrite(12345, 'uint32;b')
  1x4 uint8 array
    0    0   48   57
```

#### See also

swrite, sread, fwrite, sprintf, typecast

#### clc

Clear the text window or panel.

## **Syntax**

```
clc
clc(fd)
```

## **Description**

clc (clear console) clears the contents of the command-line window or panel.

clc(fd) clears the contents of the window or panel associated with file descriptor fd.

# disp

Simple display on the standard output.

# **Syntax**

```
disp(obj)
disp(obj, fd=fd)
```

## **Description**

disp(obj) displays the object obj. Command format may be used to control how numbers are formatted.

With named argument fd, disp(obj,fd=fd) writes obj to the file descriptor fd.

# Example

```
disp('hello')
hello
```

#### See also

format, fprintf

## fclose

Close a file.

# **Syntax**

```
fclose(fd)
fclose('all')
```

## **Description**

fclose(fd) closes the file descriptor fd which was obtained with functions such as fopen. Then fd should not be used anymore. fclose('all') closes all the open file descriptors.

### feof

Check end-of-file status.

## **Syntax**

```
b = feof(fd)
```

## **Description**

feof(fd) is false if more data can be read from file descriptor fd, and true if the end of the file has been reached.

## **Example**

Count the number of lines and characters in a file (fopen and fclose are not available in all LME applications):

```
fd = fopen('data.txt');
lines = 0;
characters = 0;
while ~feof(fd)
    str = fgets(fd);
    lines = lines + 1;
    characters = characters + length(str);
end
fclose(fd);
```

#### See also

ftell

#### fflush

Flush the input and output buffers.

# **Syntax**

```
fflush(fd)
```

## **Description**

fflush(fd) discards all the data in the input buffer and forces data out of the output buffer, when the device and their driver permits it. fflush can be useful to recover from errors.

# fgetl

Reading of a single line.

## **Syntax**

```
line = fgetl(fd)
line = fgetl(fd, n)
```

## **Description**

A single line (of at most n characters) is read from a text file. The end of line character is discarded. Upon end of file, fgetl gives an empty string.

#### See also

fgets, fscanf

# fgets

Reading of a single line.

# **Syntax**

```
line = fgets(fd)
line = fgets(fd, n)
```

# **Description**

A single line (of at most n characters) is read from a text file. Unless the end of file is encountered before, the end of line (always a single line feed) is preserved. Upon end of file, fgets gives an empty string.

#### See also

fgetl, fscanf

#### fionread

Number of bytes which can be read without blocking.

## **Syntax**

```
n = fionread(fd)
```

## **Description**

fionread(fd) returns the number of bytes which can be read without blocking. For a file, all the data until the end of the file can be read; but for a device or a network connection, fionread gives the number of bytes which have already been received and are stored in the read buffer.

If the number of bytes cannot be determined, fionread returns -1.

#### See also

fread

#### **format**

Default output format.

## **Syntax**

```
format
format short
format short e
format short eng
format short q
format long
format long e
format long eng
format long g
format int
format int d
format int u
format int x
format int o
format int b
format bank
format rat
format '+'
format i
format i
```

format loose
format compact

## **Description**

format changes the format used by command disp and for output produced with expressions which do not end with a semicolon. The following arguments are recognized:

Arguments	Meaning
(none)	fixed format with 0 or 4 digits, loose spacing
short	fixed format with 0 or 4 digits
short e	exponential format with 4 digits
short eng	engineering format with 4 digits
short g	general format with up to 4 digits
long	fixed format with 0 or 15 digits
long e	exponential format with 15 digits
long eng	engineering format with 15 digits
long g	general format with up to 15 digits
int	signed decimal integer
int d	signed decimal integer
int u	unsigned decimal integer
int x	hexadecimal integer
int o	octal integer
int b	binary integer
bank	fixed format with 2 digits (for currencies)
rat	rational approximation
+	'+', '-' or 'l' for nonzero, space for zero
i	symbol i to represent the imaginary unit
j	symbol j to represent the imaginary unit
loose	empty lines to improve readability
compact	no empty line

Format for numbers, for imaginary unit symbol and for spacing is set separately. Format rat displays rational approximations like rat with the default tolerance, but also displays the imaginary part if it exists. Format '+' displays compactly numeric and boolean arrays: positive numbers and complex numbers with a positive real part are displayed as +, negative numbers or complex numbers with a negative real part as -, pure imaginary nonzero numbers as I, and zeros as spaces.

The default format is format short g, format j, and format compact.

#### See also

disp, fprintf, rat

# **fprintf**

Formatted output.

## **Syntax**

```
n = fprintf(fd,format,a,b,...)
n = fprintf(format,a,b,...)
n = fprintf(..., fd=fd, NPrec=nPrec)
```

## **Description**

fprintf(format,a,b,...) converts its arguments to a string and writes it to the standard output.

fprintf(fd,format,a,b,...) specifies the output file descriptor. The file descriptor can also be specified as a named argument fd.

In addition to fd, fprintf also accepts named argument NPrec for the default number of digits in floating-point numbers.

See sprintf for a description of the conversion process.

## **Example**

```
fprintf('%d %.2f %.3E %g\n',1:3,pi)
1 2.00 3.000E0 3.1416
22
```

#### See also

sprintf, fwrite

#### fread

Raw input.

# **Syntax**

```
(a, count) = fread(fd)
(a, count) = fread(fd, size)
(a, count) = fread(fd, size, precision)
```

# **Description**

fread(fd) reads signed bytes from the file descriptor fd until it reaches the end of file. It returns a column vector whose elements are signed bytes (between -128 and 127), and optionally in the second output argument the number of elements it has read.

fread(fd,size) reads the number of bytes specified by size. If size is a scalar, that many bytes are read and result in a column vector. If size is a vector of two elements [m,n], m\*n elements are read row by row and stored in an m-by-n matrix. If the end of the file is reached before the specified number of elements have been read, the number of rows is reduced without throwing an error. The optional second output argument always gives the number of elements in the result. If size is the empty array [], elements are read until the end of the file: it must be specified if there is a third argument.

With a third argument, fread(fd,size,precision) reads integer words of 1, 2, or 4 bytes, or IEEE floating-point numbers of 4 bytes (single precision) or 8 bytes (double precision). The meaning of the string precision is described in the table below.

precision	meaning
int8	signed 8-bit integer (-128 $\leq$ x $\leq$ 127)
char	signed 8-bit integer (-128 $\leq$ x $\leq$ 127)
int16	signed 16-bit integer (-32768 $\le x \le 32767$ )
int32	signed 32-bit integer (-2147483648 $\leq x \leq$ 2147483647)
int64	signed 64-bit integer (-9.223372e18 $\leq x \leq 9.223372e18$ )
uint8	unsigned 8-bit integer ( $0 \le x \le 255$ )
uchar	unsigned 8-bit integer ( $0 \le x \le 255$ )
uint16	unsigned 16-bit integer ( $0 \le x \le 65535$ )
uint32	unsigned 32-bit integer (0 $\leq$ x $\leq$ 4294967295)
uint64	unsigned 64-bit integer (0 $\leq$ x $\leq$ 18.446744e18)
single	32-bit IEEE floating-point
double	64-bit IEEE floating-point

By default, multibyte words are encoded with the least significant byte first (little endian). The characters '; b' can be appended to specify that they are encoded with the most significant byte first (big endian); for symmetry, '; l' is accepted and ignored.

By default, the output is a double array. To get an output which has the same type as what is specified by precision, the character \* can be inserted at the beginning. For instance '\*uint8' reads bytes and stores them in an array of class uint8, '\*int32;b' reads signed 32-bit words and stores them in an array of class int32 after performing byte swapping if necessary, and '\*char' reads bytes and stores them in a character row vector (i.e. a plain string).

Precisions 'int64' and 'uint64' are supported only if types int64 and uint64 are supported.

#### See also

fwrite, sread

#### frewind

Rewind current read or write position in a file.

## **Syntax**

```
frewind(fd)
```

## **Description**

frewind(fd) sets the position in an open file where the next input/output commands will read or write data to the beginning of the file. The argument fd is the file descriptor returned by fopen or similar functions (fopen is not available in all LME applications).

frewind(fd) has the same effect as fseek(fd,0) or fseek(fd,0,'b').

#### See also

fseek, ftell

# fscanf

Reading of formatted numbers.

# Syntax

```
r = fscanf(fd, format)
(r, count) = fscanf(fd, format)
```

# Description

A single line is read from a text file, and numbers, characters and strings are decoded according to the format string. The format string follows the same rules as sscanf.

The optional second output argument is set to the number of elements decoded successfully (may be different than the length of the first argument if decoding strings).

# Example

Read a number from a file (fopen and fclose are not available in all LME applications):

```
fd = fopen('test.txt', 'rt');
fscanf(fd, '%f')
  2.3
fclose(fd);
```

#### See also

sscanf

## fseek

Change the current read or write position in a file.

## **Syntax**

```
status = fseek(fd, position)
status = fseek(fd, position, mode)
```

## **Description**

fseek(fd,position,mode) changes the position in an open file where the next input/output commands will read or write data. The first argument fd is the file descriptor returned by fopen or similar functions (fopen is not available in all LME applications). The second argument is the new position. The third argument mode specifies how the position is used:

- b absolute position from the beginning of the file
- c relative position from the current position
- e offset from the end of the file (must be  $\leq 0$ )

The default value is 'b'. Only the first character is checked, so 'beginning' is a valid alternative for 'b'. fseek returns 0 if successful or -1 if the position is outside the limits of the file contents.

#### See also

frewind, ftell

#### ftell

Get the current read or write position in a file.

## **Syntax**

```
position = ftell(fd)
```

## **Description**

ftell(fd) gives the current file position associated with file descriptor fd. The file position is the offset (with respect to the beginning of the file) at which the next input function will read or the next output function will write. The offset is expressed in bytes. With text files, ftell may not always correspond to the number of characters read or written.

#### See also

fseek, feof

#### **fwrite**

Raw output.

## **Syntax**

```
count = fwrite(fd, data)
count = fwrite(fd, data, precision)
```

## **Description**

fwrite(fd, data) writes the contents of the matrix data to the output referenced by the file descriptor fd. The third parameter is the precision, whose meaning is the same as for fread. Its default value is 'uint8'.

#### See also

fread, swrite, bwrite

#### redirect

Redirect or copy standard output or error to another file descriptor.

# **Syntax**

```
redirect(fd, fdTarget)
redirect(fd, fdTarget, copy)
redirect(fd)
R = redirect(fd)
redirect
R = redirect
```

## **Description**

redirect(fd,fdTarget) redirects output from file descriptor fd to fdTarget. fd must be 1 for standard output or 2 for standard error. If fdTarget==fd, the normal behavior is restored.

redirect(fd,fdTarget,copy) copies output to both fd and fdTarget if copy is true, instead of redirecting it only to fdTarget. If copy is false, the result is the same as with two input arguments.

With zero or one input argument and without output argument, redirect displays the current redirection for the specified file descriptor (1 or 2) or for both of them. Note that the redirection itself may alter where the result is displayed.

With an output argument, redirect returns a 1-by-2 row vector if the file descriptor is specified, or a 2-by-2 matrix otherwise. The first column contains the target file descriptor and the second column, 1 for copy mode and 0 for pure redirection mode.

#### **Examples**

Create a new file diary.txt and copy to it both standard output and error:

```
fd = fopen('diary.txt', 'w');
redirect(1, fd, true);
redirect(2, fd, true);
```

Stop copying standard output and error and close file:

```
redirect(1, 1);
redirect(2, 2);
fclose(fd);
```

Redirect standard error to standard output and get the redirection state:

```
redirect(2, 1)
redirect
   stdout (fd=1) -> fd=1
   stderr (fd=2) -> fd=1
redirect(2)
   stderr (fd=2) -> fd=1
R = redirect
   R =
        1 0
        1 0
R = redirect(2)
R =
        1 0
```

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# sprintf

Formatted conversion of objects into a string.

## **Syntax**

```
s = sprintf(format,a,b, ...)
s = sprintf(..., NPrec=nPrec)
```

## **Description**

sprintf converts its arguments to a string. The first parameter is the format string. All the characters are copied verbatim to the output string, except for the control sequences which all begin with the character '%'. They have the form

%fn.dt

where f is zero, one or more of the following flags:

Flag	Meaning
-	left alignment (default is right alignment)
+	display of a + sign for positive numbers
0	zero padding instead of spaces
#	alternate format (see below)
space	sign replaced with space for positive numbers

n is the optional width of the field as one or more decimal digits (default is the minimum width to display the data).

d is the number of digits after the decimal separator for a number displayed with a fractional part (default is 4 or what is specified by named argument NPrec), the minimum number of displayed digits for a number displayed as an integer, or the number of characters for a string (one or more decimal digits).

t is a single character denoting the type of conversion. In most cases, each control sequence corresponds to an additional argument.

All elements of arrays are used sequentially as if they were provided separately; strings are used as a whole. The table below gives the valid values of t.

Char.	Conversion
%	single %
d	decimal number as an integer
i	same as d
Χ	hexadecimal number (for integers between 0 and 2 <sup>32</sup> -1)
Χ	same as x, with uppercase digits
0	octal number (for integers between 0 and 2 <sup>32</sup> -1)
f	fixed number of decimals (exp. notation if $abs(x)>1e18$ )
F	same as f, with an uppercase E
е	scientific notation such as 1e5
E	scientific notation such as 1E5
n	engineering notation such as 100e3
N	engineering notation such as 100E3
g	decimal or scientific notation
Ğ	same as g, with an uppercase E
k	same as g, with as few characters as possible
K	same as k, with an uppercase E
Р	SI prefix (k=1e3, u=1e-6) or engineering notation
С	character
S	string

The # flag forces octal numbers to begin with 0, nonzero hexadecimal numbers to begin with 0x, and floating-point numbers to always have a decimal point even if they do not have a fractional part.

Integer formats %d, %i, %o and %x round fractional numbers to the nearest integer.

Instead of decimal digits, the width n and/or the precision d can be replaced with character \*; then one or two additional arguments (or elements of an array) are consumed and used as the width or precision.

# **Examples**

Numbers:

```
sprintf('%d %.2f %.2e %.2E %.2g',pi*ones(1,5))
    3 3.14 3.14e0 3.14E0 3.14

Compact representation with '%k':
    sprintf('%.1k ', 0.001, 0.11, 111, 1000)
        1e-3 0.11 111 1e3

Width and precision:
    sprintf('*%8.3f*%8.6s*%-8.6s*',pi,'abcdefgh','abcdefgh')
        * 3.142* abcdef*abcdef *
```

Repetition of format string to convert all values:

```
sprintf('%c_','a':'z')
   a_b_c_d_e_f_g_h_i_j_k_l_m_n_o_p_q_r_s_t_u_v_w_x_y_z_
```

Width and precision provided as expressions:

```
sprintf('%*.*f', 15, 7, pi)
3.1415927
```

Zero padding for integer format:

```
sprintf('%.3d,%.3d', 12, 12345)
012,12345
```

Default precision:

```
sprintf('%f %e', pi, pi)
  3.1416 3.1416e0
sprintf('%f %e', pi, pi, NPrec=2)
  3.14 3.14e0
```

#### See also

fprintf, sscanf, swrite

#### sread

Raw input from a string or an array of bytes.

## **Syntax**

```
(a, count) = sread(str, size, precision)
(a, count) = sread(str, [], precision)
(a, count) = sread(bytes, ...)
```

# **Description**

sread(str) reads data from string str or array of class uint8 or int8
the same way as fread reads data from a file.

# **Examples**

```
(data, count) = sread('abc')
  data =
    97
    98
    99
  count =
    3
```

```
(data, count) = sread('abcdef',[2,2])
  data =
    97 98
    99 100
  count =
    4
(data, count) = sread('abcd',[inf,3])
  data =
    97 98 99
  count =
    3
```

#### See also

swrite, bwrite, fread, typecast

#### sscanf

Decoding of formatted numbers.

## **Syntax**

```
r = sscanf(str, format)
(r, count) = scanf(str, format)
(r, count, nchar) = scanf(str, format)
```

# **Description**

Numbers, characters and strings are extracted from the first argument. Exactly what is extracted is controlled by the second argument, which can contain the following elements:

Substring in format	Meaning
%C	single character
%S	string
%d	integer number in decimal
%X	unsigned integer number in hexadecimal
%0	unsigned integer number in octal
%i	integer number
%f	floating-point number
%e	floating-point number
%g	floating-point number
%%	%
other character	exact match

%i recognizes an optional sign followed by a decimal number, an hexadecimal number prefixed with 0x or 0X, a binary number prefixed with 0b or 0B, or an octal number prefixed with 0.

The decoded elements are accumulated in the output argument, either as a column vector if the format string contains %d, %o, %x, %i, %f, %e or %g, or a string if the format string contains only %c, %s or literal values. If a star is inserted after the percent character, the value is decoded and discarded. A width (as one or more decimal characters) can be inserted before s, d, x, o, i, f, e or g; it limits the number of characters to be decoded. In the input string, spaces and tabulators are skipped before decoding %s, %d, %x, %o, %i, %f, %e or %g.

The format string is recycled as many times as necessary to decode the whole input string. The decoding is interrupted if a mismatch occurs.

The optional second output argument is set to the number of elements decoded successfully (may be different than the length of the first argument if decoding strings). The optional third output argument is set to the number of characters which were consumed in the input string.

## **Examples**

```
sscanf('f(2.3)', 'f(%f)')
sscanf('12a34x778'. '%d%c')
  12
  97
  34
 120
778
sscanf('abc def', '%s')
 abcdef
sscanf('abc def', '%c')
 abc def
sscanf('12,34','%*d,%d')
sscanf('0275a0ff', '%2x')
 117
 160
255
```

#### See also

sprintf

#### swrite

Store data in a string.

## **Syntax**

```
s = swrite(data)
s = swrite(data, precision)
```

## Description

swrite(data) stores the contents of the matrix data into a string. The second parameter is the precision, whose meaning is the same as for fread. Its default value is 'uint8'.

### **Examples**

```
swrite(65:68)
  ABCD
double(swrite([1,2], 'int16'))
  1 0 2 0
double(swrite([1,2], 'int16;b'))
  0 1 0 2
```

## See also

bwrite, fwrite, sprintf, sread

# 6.31 File System Functions

Access to any kind of file can be useful to analyze data which come from other applications (such as experimental data) and to generate results in a form suitable for other applications (such as source code or HTML files). Functions specific to files are described in this section. Input, output, and control are done with the following generic functions:

<b>Function</b>	Description
fclose	close the file
feof	check end of file status
fflush	flush I/O buffers
fgetl	read a line
fgets	read a line
fprintf	write formatted data
fread	read data
frewind	reset the current I/O position
fscanf	read formatted data
fseek	change the current I/O position
ftell	get the current I/O position
fwrite	write data
redirect	redirect output

# fopen

Open a file.

## **Syntax**

```
fd = fopen(path)
fd = fopen(path, mode)
```

# **Description**

fopen opens a file for reading and/or writing. The first argument is a path whose format depends on the platform. If it is a plain file name, the file is located in the current directory; what "current" means also depends on the operating system. The output argument, a real number, is a file descriptor which can be used by many input/output functions, such as fread, fprintf, or dumpvar.

The optional second input argument, a string of one or two characters, specifies the mode. It can take one of the following values:

Mode	Meaning
(none)	same as 'r'
r'	read-only, binary mode, seek to beginning
'w'	read/write, binary mode, create new file
'a'	read/write, binary mode, seek to end
rt'	read-only, text mode, seek to beginning
'wt'	read/write, text mode, create new file
'at'	read/write, text mode, seek to end

In text mode, end-of-line characters LF, CR and CRLF are all converted to LF ( $'\n'$ ) on input. On output, they are converted to the

native sequence for the operating system, which is CRLF on Windows and LF elsewhere. To force the output end-of-line to be LF irrespectively of the operating system, use 'q' instead of 't' (e.g. 'wq' to write to a file); to force it to be CRLF, use 'T' (e.g. 'aT' to append to a file).

#### **Examples**

```
Reading a whole text file into a string:
```

```
fd = fopen('file.txt', 'rt');
str = fread(fd, inf, '*char');
fclose(fd);
```

Reading a whole text file line by line:

```
fd = fopen('file.txt', 'rt');
while ~feof(fd)
    str = fgets(fd)
end
fclose(fd);
```

Writing a matrix to a CSV (comma-separated values) text file:

```
M = magic(5);
fd = fopen('file.txt', 'wt');
for i = 1:size(M, 1)
    for j = 1:size(M, 2)-1
        fprintf(fd, '%g,', M(i,j));
    end
    fprintf(fd, '%g\n', M(i,end));
end
fclose(fd);
```

Reading 5 bytes at offset 3 in a binary file, giving an 5-by-1 array of unsigned 8-bit integers:

```
fd = fopen('file.bin');
fseek(fd, 3);
data = fread(fd, 5, '*uint8');
fclose(fd);
```

#### See also

fclose

# 6.32 Path Manipulation Functions

# fileparts

File path splitting into directory, filename and extension.

## **Syntax**

```
(dir, filename, ext) = fileparts(path)
```

## **Description**

fileparts (path) splits string path into the directory (initial part until the last file separator), the filename without extension (substring after the last file separator before the last dot), and the extension (substring starting from the last dot after the last file separator).

The directory is empty if path does not contain any file separator, and the extension is empty if the remaining substring does not contain a dot. When these three strings are concatenated, the result is always equal to the initial path.

The separator depends on the operating system: a slash on unix (including Mac OS X and Linux), and a backslash or a slash on Windows.

#### **Examples**

```
(dir, filename, ext) = fileparts('/home/tom/report.txt')
 dir =
    /home/tom/
 filename =
    report
 ext =
(dir, filename, ext) = fileparts('/home/tom/')
    /home/tom/
  filename =
 ext =
(dir, filename, ext) = fileparts('results.txt.back')
 dir =
  filename =
    results.txt
 ext =
    . back
```

#### See also

```
fullfile, filesep
```

# filesep

File path separator.

## **Syntax**

```
ch = filesep
```

## **Description**

filesep gives the character used as separator between directories and files in paths. It depends on the operating system: a slash on unix (including Mac OS X and Linux), and a backslash on Windows.

#### See also

```
fullfile, fileparts
```

## fullfile

File path construction.

#### **Syntax**

```
path = fullfile(p1, p2, ...)
```

## **Description**

fullfile constructs a file path by concatenating all its string arguments, removing separators when missing. At least one input argument is required.

## **Examples**

```
fullfile('/home/tom/', 'report.txt')
  /home/tom/report.txt
fullfile('/home/tom', 'data', 'meas1.csv')
  /home/tom/data/meas1.csv
```

#### See also

```
fileparts, filesep
```

# 6.33 XML Functions

This section describes functions which import XML data. Two separate sets of functions implement two approaches to parse XML data:

- Document Object Model (DOM): XML is loaded entirely in memory from a file (xmlread) or a character string (xmlreadstring).
   Additional functions permit to traverse the DOM tree and to get its structure, the element names and attributes and the text.
- Simple API for XML (SAX): XML is parsed from a file descriptor (saxnew) and events are generated for document start and end, element start and end, and character sequences.

With both approaches, creation and modification of the document are not possible.

#### DOM

Two opaque types are implemented: DOM nodes (including document, element and text nodes), and attribute lists. A document node object is created with the functions xmlreadstring (XML string) or xmlread (XML file or other input channel). Other DOM nodes and attribute lists are obtained by using DOM methods and properties.

# Methods and properties of DOM node objects

# fieldnames getElementById getElementsByTagName

subsref xmlrelease

Method

#### **Description**

List of property names Get a node specified by id

Get a list of all descendent nodes of the given tag nam Get a property value

Release a document node

**Property** Description attributes Attribute list (opaque object) childElementCount Number of element children childNodes List of child nodes children List of element child nodes depth Node depth in document tree Root element of a document node documentElement firstChild First child node firstElementChild First element child node lastChild Last child node lastElementChild Last element child node line Line number in original XML document Next sibling element node nextElementSibling Next sibling node nextSibling nodeName Node tag name, '#document', or '#text' nodeValue Text of a text node offset Offset in original XML document

ownerDocument Owner DOM document node
parentNode Parent node
previousElementSibling Previous sibling element node

previousSibling Previous sibling element

Previous sibling node

textContent Concatenated text of all descendent text nodes XML representation, including all children

A document node object is released with the xmlrelease method. Once a document node object is released, all associated node objects become invalid. Attribute lists and native LME types (strings and numbers) remain valid.

## Methods and properties of DOM attribute list objects

Method	Description
fieldnames	List of attribute names
length	Number of attributes
subsref	Get an attribute

Properties of attribute lists are the attribute values as strings. Properties whose name is compatible with LME field name syntax can be retrieved with the dot syntax, such as attr.id. For names containing invalid characters, such as accented letters, or to enumerate unknown attributes, attributes can be accessed with indexing, with either parenthesis or braces. The result is a structure with two fields name and value.

#### SAX

XML is read from a file descriptor, typically obtained with fopen. The next event is retrieved with saxnext which returns its description in a structure.

# getElementById

Get a node specified by id.

## **Syntax**

```
node = getElementById(root, id)
```

## **Description**

getElementById(root,id) gets the node which is a descendant of node root and whose attribute id matches argument id. It throws an error if the node is not found.

In valid XML documents, every id must be unique. If the document is invalid, the first element with the specified id is obtained.

#### See also

xmlread, getElementsByTagName

# getElementsByTagName

Get a list of all descendent nodes of the given tag name.

# **Syntax**

```
node = getElementsByTagName(root, name)
```

# **Description**

getElementsByTagName(root,name) collects a list of all the element nodes which are direct or indirect descendants of node root and whose name matches argument name.

#### **Examples**

```
doc = xmlreal('Abc <b>de</b> <i>fg <b>hijk</b></i>');
b = getElementsByTagName(doc, 'b')
b =
    {DOMNode,DOMNode}
b2 = b{2}.xml
b2 =
    <b>hijk</b>
xmlrelease(doc);
```

#### See also

xmlread, getElementById

## saxcurrentline

Get current line number of SAX parser.

## **Syntax**

```
n = saxcurrentline(sax)
```

# **Description**

saxcurrentline(sax) gets the current line of the XML file parsed by the SAX parser passed as argument. It can also be used after an error.

#### See also

saxcurrentpos, saxnew, saxnext

# saxcurrentpos

Get current position in input stream of SAX parser.

# **Syntax**

```
n = saxcurrentpos(sax)
```

## **Description**

saxcurrentpos(sax) gets the current position of the XML file parsed by the SAX parser passed as argument (the number of bytes consumed thus far). It can also be used after an error.

The value given by saxcurrentpos differs from the result of ftell on the file descriptor, because the SAX parser input is buffered.

#### See also

saxcurrentline, saxnew, saxnext

#### saxnew

Create a new SAX parser.

## **Syntax**

```
sax = saxnew(fd)
sax = saxnew(fd, Trim=t, HTML=h)
```

## **Description**

saxnew(fd) create a new SAX parser to parse XML from file descriptor fd. The parser is an opaque (non-numeric) type. Once it is not needed anymore, it should be released with the saxrelease function.

Named argument Trim (a boolean value) specifies if white spaces are trimmed around tags. The default value is false.

Named argument HTML (a boolean value) specifies HTML mode. The default value is false (XML mode). In HTML mode, unknown entities and less-than characters not followed by tag names are considered as plain text, and attribute values can be missing (same as attribute names) or unquoted. This can be used for the lowest level of a rudimentary HTML parser.

# **Example**

```
fd = fopen('data.xml');
sax = saxnew(fd);
while true
  ev = saxnext(sax);
  switch ev.event
    case 'docBegin'
    // beginning of document
    case 'docEnd'
    // end of document
```

```
break;
case 'elBegin'
    // beginning of element ev.tag with attr ev.attr
case 'elEnd'
    // end of element ev.tag
case 'elEmpty'
    // empty element ev.tag with attr ev.attr
case 'text'
    // text element ev.text
end
end
saxrelease(sax);
fclose(fd);
```

#### See also

saxrelease, saxnext, xmlread

#### saxnext

Get next SAX event.

# Syntax

```
event = saxnext(sax)
```

## **Description**

saxnext(sax) gets the next SAX event and returns its description in a structure. Argument sax is the SAX parser created with saxnew.

The event structure contains the following fields:

```
event Event type as a string: 'docBegin', 'docEnd', 'elBegin',
  'elEnd', 'elEmpty', or 'text'.
```

```
tag For 'elBegin', 'elEnd' and 'elEmpty', element tag.
```

**attr** For 'elBegin' and 'elEmpty', structure array containing the element attributes. Each attribute is defined by two string fields, name and value.

```
text For 'text', text string.
```

#### See also

saxnew, saxcurrentline

#### saxrelease

Release a SAX parser.

## **Syntax**

```
saxrelease(sax)
```

## **Description**

saxrelease(sax) releases the SAX parser sax created with saxnew.

#### See also

saxnew

#### **xmlread**

Load a DOM document object from a file descriptor.

## **Syntax**

```
doc = xmlread(fd)
```

# Description

xmlread(fd) loads XML to a new DOM document node object by reading a file descriptor until the end, and returns a new document node object. The file descriptor can be closed before the document node object is used. Once the document is not needed anymore, it should be released with the xmlrelease method.

# **Example**

Load an XML file 'doc.xml' (this assumes support for files with the function fopen).

```
fd = fopen('doc.xml');
doc = xmlread(fd);
fclose(fd);
root = doc.documentElement;
...
xmlrelease(doc);
```

#### See also

xmlreadstring, xmlrelease, saxnew

# xmlreadstring

Parse an XML string into a DOM document object.

## **Syntax**

```
doc = xmlreadstring(str)
```

## **Description**

xmlreadstring(str) parses XML from a string to a new DOM document node object. Once the document is not needed anymore, it should be released with the xmlrelease method.

#### **Examples**

```
xml = '<a>one < b id="x">two</b> < c id="y" num="3">three</c></a>';
doc = xmlreadstring(xml)
 doc =
  DOM document
root = doc.documentElement;
root.nodeName
  ans =
root.childNodes{1}.nodeValue
  ans =
    one
root.childNodes{2}.xml
    <b id="x">two</b>
a = root.childNodes{2}.attributes
 DOM attributes (1 item)
a.id
  Х
qetElementById(doc,'y').xml
  <c id="y" num="3">three</c>
xmlrelease(doc);
```

#### See also

xmlread, xmlrelease

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## **xmlrelease**

Release a DOM document object.

## **Syntax**

xmlrelease(doc)

## **Description**

xmlrelease(doc) releases a DOM document object. All DOM node objects obtained directly or indirectly from it become invalid.

Releasing a node which is not a document has no effect.

### See also

xmlreadstring, xmlread

# 6.34 Time Functions

## clock

Current date and time.

## **Syntax**

t = clock

# **Description**

clock returns a 1x6 row vector, containing the year (four digits), the month, the day, the hour, the minute and the second of the current local date and time. All numbers are integers, except for the seconds which are fractional. The absolute precision is plus or minus one second with respect to the computer's clock; the relative precision is plus or minus 1 microsecond on a Macintosh, and plus or minus 1 millisecond on Windows.

## **Example**

```
clock
1999 3 11 15 37 34.9167
```

### See also

posixtime, tic, toc

# posixtime

Current Posix time.

## **Syntax**

```
t = posixtime
```

# Description

posixtime returns the Posix time, the integral number of seconds since 1 February 1970 at 00:00:00 UTC, based on the value provided by the operating system.

## **Example**

```
posixtime
1438164887
```

### See also

clock

### tic

Start stopwatch.

### **Syntax**

```
tic
t0 = tic
tic(CPUTime=true)
t0 = tic(CPUTime=true)
```

# **Description**

Without output argument, tic resets the stopwatch. Typically, tic is used once at the beginning of the block to be timed, and toc at the end. toc can be called multiple times to get intermediate times.

With an output argument, tic gets the current state of the stopwatch. Multiple independent time measurements can be performed by passing this value to toc.

By default, tic and toc are based on the real, "wall-clock" time. tic(CPUTime=true) is based on CPU time instead, which gives more accurate results for non-multithreaded programs. Measurements made with wall-clock time and CPU time are independent and can be mixed freely.

### See also

toc, clock

### toc

Elapsed time of stopwatch.

## **Syntax**

```
elapsed_time = toc
elapsed_time = toc(t0)
elapsed_time = toc(CPUTime=true)
elapsed_time = toc(t0, CPUTime=true)
```

# **Description**

Without input argument, toc gets the time elapsed since the last execution of tic without output argument. Typically, toc is used at the end of the block of statements to be timed.

With an input argument, toc(t0) gets the time elapsed since the execution of t0=tic. Multiple independent time measurements, nested or overlapping, can be performed simultaneously.

With а named argument. toc(CPUTime=true) or toc(t0,CPUTime=true) use CPU time: toc measures only the time spent in the LME application. Other processes do not have typing tic(CPUTime=true) large impact. For instance. at the command-line prompt, waiting 5 seconds, and typing toc(CPUTime=true) will show a value much smaller than 5: while typing tic and toc will show the same elapsed time a chronograph would. CPU time is usually more accurate for non-multithreaded code, and wall-clock time for multi-threaded code, or measurements involving devices or network communication.

### **Examples**

Time spent to compute the eigenvalues of a random matrix:

```
tic; x = eig(rand(200)); toc
  0.3046
```

Percentage of time spent in computing eigenvalues in a larger program:

```
eigTime = 0;
s = [];
tic(CPUTime=true);
for i = 1:100
    A = randn(20);
    eigT0 = tic(CPUTime=true);
    s1 = eig(A);
    eigTime = eigTime + toc(eigT0, CPUTime=true);
    s = [s, s1];
end
totalTime = toc(CPUTime=true);
100 * eigTime / totalTime
    78.4820
```

### See also

tic. clock

# 6.35 Date Functions

Date functions perform date and time conversions between the calendar date and the julian date.

The calendar date is the date of the proleptic Gregorian calendar, i.e. the calendar used in most countries today where centennial years are not leap unless they are a multiple of 400. This calendar was introduced by Pope Gregory XIII on October 5, 1582 (Julian Calendar, the calendar used until then) which became October 15. The calendar used in this library is proleptic, which means the rule for leap years is applied back to the past, before its introduction. Negative years are permitted; the year 0 does exist.

The julian date is the number of days since the reference point, January 1st -4713 B.C. (Julian calendar) at noon. The fractional part corresponds to the fraction of day after noon: a fraction of 0.25, for instance, is 18:00 or 6 P.M. The julian date is used by astronomers with GMT; but using a local time zone is fine as long as an absolute time is not required.

# cal2julian

Calendar to julian date conversion.

## **Syntax**

```
jd = cal2julian(datetime)
jd = cal2julian(year, month, day)
jd = cal2julian(year, month, day, hour, minute, second)
```

## **Description**

cal2julian(datetime) converts the calendar date and time to the julian date. Input arguments can be a vector of 3 components (year, month and day) or 6 components (date and hour, minute and seconds), or scalar values provided separately. The result of clock can be used directly.

## **Example**

Number of days between October 4 1967 and April 18 2005:

```
cal2julian(2005, 4, 18) - cal2julian(1967, 10, 4) 14624
```

### See also

```
julian2cal, clock
```

# julian2cal

Julian date to calendar conversion.

# **Syntax**

```
datetime = julian2cal(jd)
(year, month, day, hour, minute, second) = julian2cal(jd)
```

# **Description**

julian2cal(jd) converts the julian date to calendar date and time. With a single output, the result is given a a row vector of 6 values for the year, month, day, hour, minute and second; with more output arguments, values are given separately.

### **Example**

```
Date 1000 days after April 18 2005:
```

### See also

cal2julian

# 6.36 MAT-files

### matfiledecode

Decode the contents of a MATLAB MAT-file.

## **Syntax**

```
var = matfiledecode(fd)
var = matfiledecode(data)
var = matfiledecode(..., ignoreErr)
```

# **Description**

matfiledecode(fd) reads data from file descriptor fd until the end of the file. The data must be the contents of a MATLAB-compatible MAT-file. They are made of 8-bit bytes; no text conversion must take place. The result is a structure whose fields have the name and the contents of the variables saved in the MAT-file.

Instead of a file descriptor, the data can be provided directly as the argument. In that case, the argument data must be an array, which can be read from the actual file with fread or obtained from a network connection.

Only arrays are supported (scalar, matrices, arrays of more than two dimensions, real or complex, numeric, logical or char). A second input argument can be used to specify how to handle data of unsupported types: with false (default value), unsupported types cause an error; with true, they are ignored.

# **Example**

```
fd = fopen('data.mat');
s = matfiledecode(fd);
fclose(fd);
s
```

```
s =
    x: real 1x1024
    y: real 1x1024
```

### See also

matfileencode

# matfileencode

Encode the contents of a MATLAB MAT-file.

## **Syntax**

```
matfileencode(fd, s)
matfileencode(s)
```

## **Description**

matfileencode(fd,s) writes the contents of structure s to file descriptor fd as a MATLAB-compatible MAT-file. Each field of s corresponds to a separate variable in the MAT-file. With one argument, matfileencode(s) writes to the standard output (which should be uncommon since MAT-files contain non-printable bytes).

Only arrays are supported (scalar, matrices, arrays of more than two dimensions, real or complex, numeric, logical or char).

# **Examples**

```
s.a = 123;
s.b = 'abc';
fd = fopen('data.mat', 'wb');
matfileencode(fd, s);
fclose(fd);
```

Function variables can be used to save all variables:

```
v = variables;
fd = fopen('var.mat', 'wb');
matfileencode(fd, v);
fclose(fd);
```

#### See also

matfiledecode, variables

# 6.37 Shell

This section describes functions related to the Unix or Windows shell. They are available only on Windows and on Unix (or Unix-like) systems, such as macOS.

The versions for Unix and Windows have significant differences:

Most functions described here are defined on both Unix and Windows, to avoid errors when loading functions which contain conditional code for Unix and Windows. Functions with an empty implementation return the error "Not supported". Table below gives the status of all commands.

Command	Unix	Windows
cd	supported	supported
cputime	supported	undefined
dir	supported	supported
dos	not supported	supported
getenv	supported	supported
pwd	supported	supported
setenv	supported	not supported
sleep	supported	supported
unix	supported	not supported
unsetenv	supported	not supported

On Windows, some of the functionality of unix is provided by dos.
 The main difference is that dos does not give any output, except for the status code of the command.

# **Functions**

### cd

Set or get the current directory.

## **Syntax**

cd(str) str = cd

# **Description**

cd(str) changes the current directory. Without input argument, cd gives the current directory, like pwd.

The current directory is the root directory where files specified by relative paths are searched by functions like fopen and dir. LME libraries are specified by name, not by path; the places where they are searched is specified by a list of search paths, typically specified with a path command or a dialog box in a graphical user interface.

## **Example**

```
cd('/usr/include');
```

### See also

pwd, dir

# cputime

Amount of processing time since the beginning of the process.

## **Syntax**

```
t = cputime
```

# **Description**

cputime gives the amount of processing time spent since the application has been launched.

### See also

```
posixtime, clock
```

### dir

List of files and directories.

## **Syntax**

```
dir
dir(path)
r = dir
r = dir(path)
```

# **Description**

dir displays the list of files and directories in the current path. A string input argument can specify the path.

With an output argument, dir returns the list of files and directories as a structure array with the following fields:

Name	Value
name	file name or directory name
isdir	false for files, true for directories
altname	alternate name (Windows only)

Field isdir may be missing on some patforms. On Windows, altname contains the DOS-compatible name (a.k.a. "8.3") if it exists, or an empty string otherwise.

### See also

cd, pwd

### dos

Execute a command under Windows.

## Syntax

```
status = dos(str)
```

# **Description**

dos(str) executes a command with the system Windows function. No input can be provided, and output is discarded. dos returns the status code of the command, which is normally 0 for successful execution.

# Example

```
dos('del C:/tmp/data.txt');
```

### See also

unix

# getenv

Get the value of an environment variable.

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## **Syntax**

```
value = getenv(name)
```

## **Description**

getenv(name) gives the value of the environment variable of the specified name. If no such environment variable exists, getenv returns an empty string.

## **Example**

```
user = getenv('USER');
```

### See also

setenv, unsetenv

## pwd

Get the current directory.

## **Syntax**

```
str = pwd
```

# Description

pwd ("print working directory") gives the current directory. It has the same effect as cd without input argument.

### See also

cd, dir

### setenv

Set the value of an environment variable.

# **Syntax**

```
setenv(name, value)
setenv(name)
```

## **Description**

setenv(name, value) sets the value of the environment variable of the specified name. Both arguments are strings. If no such environment variable exists, it is created.

With a single input argument, setenv creates an empty environment variable (or remove the value of an exisisting environment variable).

Environment variables are defined in the context of the application; they can be accessed in the application or in processes it launches. Environment variables of the calling process (command shell, for instance) are not changed.

seteny is not defined for Windows.

## **Example**

```
setenv('CONTROLDEBUG', '1');
```

### See also

getenv, unsetenv

# sleep

Suspend execution for a specified amount of time.

### **Syntax**

```
sleep(t)
```

## **Description**

sleep(t) suspend execution during t seconds with a resolution of a microsecond.

### **Example**

```
sleep(1e-3);
```

## unix

Execute a Unix command.

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## **Syntax**

```
unix(str)
```

## **Description**

unix(str) executes a command with the default shell. No input can be provided, and output is directed to the standard output of LME.

## **Examples**

```
unix ls
unix('cc -o calc calc.c; ./calc')
```

### See also

dos

### unsetenv

Remove an environment variable.

# **Syntax**

```
unsetenv(name)
```

# **Description**

unsetenv(name) removes the definition of the environment variable of the specified name. Argument is a string. If no such environment variable exists, unsetenv does nothing.

Environment variables are defined in the context of the application; they can be accessed in the application or in processes it launches. Environment variables of the calling process (command shell, for instance) are not changed.

unsetenv is not defined for Windows.

# **Example**

```
unsetenv('CONTROLDEBUG');
```

#### See also

getenv, setenv

# 6.38 Graphics

LME provides low-level commands for basic shapes as well as high-level commands for more specialized plots:

**Low-level commands** Low-level commands add simple shapes such as lines, marks, polygons, circles and images. With them, you can display virtually everything you want. Arguments of these commands are such that it is very easy to work globally with matrices without computing each value sequentially in a loop.

**High-level commands** High-level commands perform some computation of their own to process their arguments before displaying the result. This has two benefits: first, the code is simpler, more compact, and faster to develop. Second, command execution is faster, because the additional processing is not interpreted by LME, but implemented as native machine code. The information related to interactive manipulation is often easier to use, too. Most of these functions are related to automatic control and signal processing.

Here is the list of these commands:

2D low-level drawing commands

activeregion	colormap	pcolor
area	contour	plot
bar	fplot	polar
barh	image	quiver
circle	line	text

### 2D high-level drawing commands

bodemag	dsigma	nyquist
bodephase	dstep	nyquist
dbodemag	erlocus	plotroots
dbodephase	hgrid	rlocus
dimpulse	hstep	sgrid
dinitial	impulse	sigma
dlsim	initial	step
dnichols	lsim	zgrid
dnyquist	ngrid	

altscale label figurestyle legend	plotopt scale e scalefa scaleov	ctor	ticks title
3D contour3 line3 mesh	plot3 plotpoly sensor3	surf	
2D!:	and liabting		

## 3D scaling and lighting

camdolly	camroll	daspect
camorbit	camtarget	lightangle
campan	camup	material
campos	camva	
camproj	camzoom	

# 6.39 Remarks on graphics

Many functions which produce the display of graphical data accept two optional arguments: one to specify the style of lines and symbols, and one to identify the graphical element for interactive manipulation.

# **Style**

The style defines the color, the line dash pattern (for continuous traces) or the shape (for discrete points) of the data.

There are two different ways to specify the style. The first one, described below, is with a single string. The second one, introduced with Sysquake 5, is with an option structure built with plotset or directly with named arguments; it is more verbose, hence easier to understand, and gives access to more settings, such as line width or marker colors.

The possible values in a style string are given below. Note that the color is ignored on some output devices (such as black and white printers) and the dash pattern is used only on high-resolution devices (such as printers or EPS output). The color code is lowercase for thin lines and uppercase for thicker lines on devices which support it.

Color	String
black	k
blue	b
green	g
cyan	С
red	r
magenta	m
yellow	У
white	W
RGB	h(rrggbb)
RGB	h(rgb)

### Dash Pattern String

solid	_ (underscore)
dashed	-
dotted	:
dash-dot	1

Shape	String
none (invisible)	(space)
point	
circle	0
cross	Χ
plus	+
star	*
triangle up	^
triangle down	V
square	[
diamond	<

Miscellaneous	String
stairs	S
stems	t
fill	f
arrow at end	a
arrows at beginning and end	Α

Color 'h(rrggbb)' specifies a color by its red, green, and blue components; each of them is given by two hexadecimal digits from 00 (minimum brightness) to ff (maximum brightness). Color 'h(rgb)' specifies each component with a single hexadecimal digit. For example, 'h(339933)' and 'h(393)' both specify the same greenish gray. Like for other colors, an uppercase 'H' means that the line is thick.

Style 's' (stairs) is supported only by the plot, dimpulse, dstep, dlsim, and dinitial functions. It is equivalent to a zero-order hold, i.e. two points are linked with a horizontal segment followed by a vertical segment.

Style 't' (stems) draws for each value a circle like 'o' and a vertical line which connects it to the origin (in 2D plots, y=0 for linear scale or y=1 for logarithmic scale; in 3D plots, z=0). In polar plots, stems connects points to x=y=0.

Style 'f' (fill) fills the shape instead of drawing its contour. Exactly how the shape is filled depends on the underlying graphics architecture; if the contour intersects itself, there may be holes.

Style 'a' adds an arrow at the end of lines drawn by plot, and style 'A' adds arrows to the beginning and the end. The arrow size depends only on the default character size, neither on the line length nor on the plot scale. Its color and thickness are the same as the line's.

Many graphical commands accept data for more than one line. If the style string contains several sequences of styles, the first line borrows its style from the first sequence, the second line, from the second sequence, and so on. If there are not enough styles, they are recycled. A sequence is one or two style specifications, one of them for the color and the other one for the dash pattern or the symbol shape, in any order. Sequences of two specifications are used if possible. Commas may be used to remove ambiguity. Here are some examples:

```
plot([0,1;0,1;0,1],[1,1;2,2;3,3],'k-r!')
```

The first line (from (0,1) to (1,1)) is black and dashed, the second line (from (0,2) to (1,2)) is red and dash-dot, and the third line (from (0,3) to (1,3)) is black and dashed again.

```
plot([0,1;0,1;0,1],[1,1;2,2;3,3],'rbk')
```

The first line is red, the second line is blue, and the third line is black.

```
plot([0,1;0,1;0,1],[1,1;2,2;3,3],'-br')
```

The first and third lines are blue and dashed, and the second line is red and solid.

```
plot([0,1;0,1;0,1],[1,1;2,2;3,3],':,H(ccccc)')
```

The first and third lines are dotted, and the second line is gray, solid, and thick.

# **Graphic ID**

The second optional argument is the graphic ID. It has two purposes. First, it specifies that the graphic element can be manipulated by the user. When the user clicks in a figure, Sysquake scans all the curves which have a non-negative graphic ID (the default value of all commands is -1, making the graphical object impossible to grasp) and sets

 $_{z0}$ ,  $_{x0}$ ,  $_{y0}$ ,  $_{id}$ , and  $_{ix}$  such that they correspond to the nearest element if it is close enough to the mouse coordinates. Second, the argument  $_{id}$  is set to the ID value so that the mousedown, mousedrag, and mouseup handlers can identify the different objects the user can manipulate.

In applications without live interactivity, such as Sysquake Remote, the graphic ID argument is accepted for compatibility reasons, but ignored.

# Scale

Before any figure can be drawn on the screen, the scale (or equivalently the portion of the plane which is represented on the screen) must be determined. The scale depends on the kind of graphics, and consequently is specified in the draw handler, but can be changed by the user with the zoom and shift commands. What the user specifies has always the priority. If he or she has not specified a new scale, the scale command found in the draw handler is used:

```
scale([xMin,xMax,yMin,yMax])
```

If scale is not used, or if some of the limits are NaN (not an number), a default scale is given by the plot commands themselves. If used, the scale command should always be executed before any plot command, because several of them use the scale to calculate traces only over the visible range or to adjust the density of the calculated points of the traces.

If you need to know the limits of the displayed area in your draw handler, use scale to get them right after setting the default scale, so that you take into account the zoom and shift specified by the user:

```
scale(optString, [defXMin, defXMax, defYMin, defYMax]);
sc = scale;
xMin = sc(1);
xMax = sc(2);
yMin = sc(3);
yMax = sc(4);
```

# **Grids**

In addition to the scale ticks displayed along the bounding frame, grids can be added to give visual clues and make easier the interpretation of graphics. X and Y grids are vertical or horizontal lines displayed in the figure background. They can be switched on and off by the user in the Grid menu, or switched on by programs with the plotoption

command (they are set off by default). In the example below, both X and Y grids are switched on:

```
plotoption xgrid
plotoption ygrid
plot(rand(1,10));
```

Commands which display grids for special kind of graphics are also available:

Command	Intended use
hgrid	nyquist, dnyquist
ngrid	nichols, dnichols
sgrid	plotroots, rlocus (continuous-time)
zgrid	plotroots, rlocus (discrete-time)

They can be used without argument, to let the user choose the level of details: *none* means the command does not produce any output; *basic* is the default value and gives a simple, non-obstructive hint (a single line or a circle); and *full* gives more details. To change by program the default level of details (basic), plotoption is used. In the example below, the grid for the complex plane of the z transform is displayed with full details. Once the figure is displayed, the user is free to reduce the level of details with the Grid menu.

```
scale('equal', [-2,2,-2,2]);
zgrid;
plotoption fullgrid;
plotroots([1,-1.5,0.8]);
```

# **6.40 Base Graphical Functions**

# activeregion

Region associated with an ID.

# **Syntax**

```
activeregion(xmin, xmax, ymin, ymax, id)
activeregion(X, Y, id)
```

# **Description**

The command activeregion defines invisible regions with an ID for interactive manipulations in Sysquake. Contrary to most other graphical objects, a hit is detected when the mouse is inside the region, not close like with points and lines.

activeregion(xmin,xmax,ymin,ymax,id) defines a rectangular shape.

activeregion(X,Y,id) defines a polygonal shape. The start and end points do not have to be the same; the shape is closed automatically.

## **Example**

Rectangular button. If an ID was given to plot without activeregion, a hit would be detected when the mouse is close to any of the four corners; with activeregion, a hit is detected when the mouse is inside the rectangle.

```
plot([50, 70, 70, 50, 50], [10, 10, 30, 30, 10]); activeregion(50, 70, 10, 30, id=1);
```

### See also

plot, image

## altscale

Alternative y scale for 2D plots.

### **Syntax**

```
altscale(b)
```

## **Description**

altscale(b) selects an alternative y scale whose axis and labels are displayed on the right of the rectangular frame of 2D plots. Its input argument is a logical value which is true to select the alternative scale and false to revert to the primary scale.

# Example

```
bar(1:5, rand(1, 5));
altscale(true);
plot(1:5, 3 * rand(1,5), 'R');
label('', 'y1', 'y2');
legend('y1\ny2', 'bfR');
```

#### See also

scale, label

#### area

Area plot.

## **Syntax**

```
area(y)
area(x, y)
area(x, y, y0)
area(..., style)
area(..., style, id)
```

## **Description**

With column vector arguments, area(x,y) displays the area between the horizontal axis y=0 and the points given by x and y. When the second argument is an array with as many rows as elements in x, area(x,Y) displays the contribution of each column of Y, summed along each row. When both the first and second arguments are arrays of the same size, area(X,Y) displays independent area plots for corresponding columns of X and Y without summation.

With a single argument, area(y) takes integers 1, 2, ..., n for the horizontal coordinates.

With a third argument, area(x, y, y0) displays the area between the horizontal line y=y0 and values defined by y.

The optional arguments style and id have their usual meaning. area uses default colors when argument style is missing.

## **Examples**

Red area defined by points (1,2), (2,3), (3,1), and (5,2) above y=0; on top of it, blue area defined by points (1,2+1), (2,3+2) etc.

```
area([1;2;3;5],[2,1;3,2;1,5;2,1], 0, 'rb');
```

Two separate areas above y=0.2 defined by points (1,2), (2,3), (3,1), (5,2); and (6,1), (7,2), (8,5), and (9,1).

```
area([1,6;2,7;3,8;5,9],[2,1;3,2;1,5;2,1], 0.2, 'rb');
```

### See also

```
plot, bar, hbar
```

### bar

Vertical bar plot.

## **Syntax**

```
bar(y)
bar(x, y)
bar(x, y, w)
bar(..., kind)
bar(..., kind, style)
bar(...., id)
```

## **Description**

bar(x,y) plots the columns of y as vertical bars centered around the corresponding value in x. If x is not specified, its default value is 1:size(y,2).

bar(x, y, w), where w is scalar, specifies the relative width of each bar with respect to the horizontal distance between the bars; with values smaller than 1, bars are separated with a gap, while with values larger than 1, bars overlap. If w is a vector of two components [w1, w2], w1 corresponds to the relative width of each bar in a group (columns of y), and w2 to the relative width of each group. Default values, used if w is missing or is the empty matrix [], is 0.8 for both w1 and w2.

bar(...,kind), where kind is a string, specifies the kind of bar plot. The following values are recognized:

```
'grouped' Columns of y are grouped horizontally (default)
```

'stacked' Columns of y are stacked vertically

'interval' Bars defined with min and max val.

With 'interval', intervals are defined by two consecutive rows of y, which must have an even number of rows.

The optional arguments style and id have their usual meaning. bar uses default colors when argument style is missing.

# **Examples**

```
Simple bar plot (see Fig. 6.5):

bar([2,4,3,6;3,5,4,1]);

Stacked bar plot:

bar(1:4, magic(4), [], 'stacked');

Interval plot:

bar(1:4, [2,4,3,1;5,6,4,6], [], 'interval');
```

### See also

barh, plot

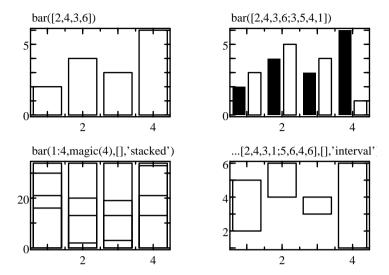


Figure 6.5 Example of bar with different options

### barh

Horizontal bar plot.

## **Syntax**

```
barh(x)
barh(y, x)
barh(y, x, w)
barh(..., kind)
barh(..., kind, style)
barh(..., id)
```

# **Description**

barh plots a bar plot with horizontal bars. Please see bar for a description of its behavior and arguments.

# **Examples**

Simple horizontal bar plot:

```
barh([2,4,3,6;3,5,4,1]);
```

Stacked horizontal bar plot:

```
barh(1:4, magic(4), [], 'stacked');
Horizontal interval plot:
barh(1:4, [2,4,3,1;5,6,4,6], [], 'interval');
```

### See also

bar, plot

### circle

Add circles to the figure.

## **Syntax**

```
circle(x,y,r)
circle(x,y,r,style)
circle(x,y,r,style,id)
```

## **Description**

circle(x,y,r) draws a circle of radius r centered at (x,y). The arguments can be vectors to display several circles. Their dimensions must match; scalar numbers are repeated if necessary. The optional fourth and fifth arguments are the style and object ID (cf. their description above).

In mouse handlers,  $\_x0$  and  $\_y0$  correspond to the projection of the mouse click onto the circle;  $\_nb$  is the index of the circle in x, y and r, and  $\_ix$  is empty.

Circles are displayed as circles only if the scales along the x and y axes are the same, and linear. With different linear scales, circles are displayed as ellipses. With logarithmic scales, they are not displayed.

# **Examples**

```
circle(1, 2, 5, 'r', 1);
circle(zeros(10,1), zeros(10, 1), 1:10);
```

### See also

plot, line

# colormap

Current colormap from scalar to RGB.

### **Syntax**

```
colormap(clut)
clut = colormap
```

## **Description**

Command colormap(clut) changes the color mapping from scalar values to RGB values used by commands such as pcolor, image and surf.

Colormaps are arrays of size n-by-3. Each row corresponds to a color; the first column is the intensity of red from 0 (no red component) to 1 (maximum intensity), the second column the intensity of green, and the third column the intensity of blue. Input values are mapped uniformly to one of the discrete color entries, 0 to the first row and 1 to the last row.

With an input argument, colormap(clut) sets the colormap to clut. With an output argument, colormap returns the current colormap.

### See also

pcolor, image

### contour

Level curves.

## **Syntax**

```
contour(z)
contour(z, [xmin, xmax, ymin, ymax])
contour(z, [xmin, xmax, ymin, ymax], levels)
contour(z, [xmin, xmax, ymin, ymax], levels, style)
```

# **Description**

contour(z) plots seven contour lines corresponding to the surface whose samples at equidistant points 1:size(z,2) in the x direction and 1:size(z,1) on the y direction are given by z. Contour lines are at equidistant levels. With a second non-empty argument [xmin, xmax, ymin, ymax], the samples are at equidistant points between xmin and xmax in the x direction and between ymin and ymax in the y direction.

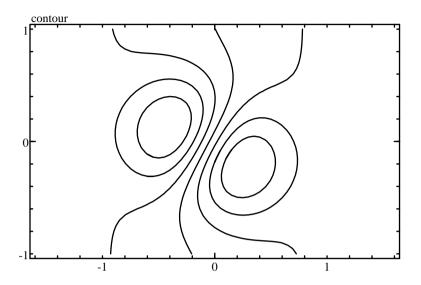


Figure 6.6 Example of contour

The optional third argument levels, if non-empty, gives the number of contour lines if it is a scalar or the levels themselves if it is a vector.

The optional fourth argument is the style of each line, from the minimum to the maximum level (styles are recycled if necessary). The default style is 'kbrmgcy'.

When the style is f for a filled region, the corresponding level is filled on the side with a lower value of z. If the style argument is the single character 'f', all levels are filled with the default colors. Regions with a value of z smaller than the lowest level are left transparent; an explicit lower level should be specified to fill the whole rectangle.

## **Examples**

A function is evaluated over a grid of two variables x and y, and is drawn with contour (see Fig. 6.6):

```
(x, y) = meshgrid(-2 + (0:40) / 10);

z = exp(-((x-0.2).^2+(y+0.3).^2)) ...

- exp(-((x+0.5).^2+(y-0.1).^2)) + 0.1 * x;

scale\ equal;

contour(z, [-1,1,-1,1]);
```

Filled contours:

### See also

image, quiver

# figurestyle

Figure style.

# **Syntax**

```
figurestyle(name, style)
style = figurestyle(name)
```

# **Description**

figurestyle sets or gets the style of figures. The same settings apply to all subplots; settings for specific subplots are changed with subplotstyle. Styles are set or got separately for each feature of the graphics (plot background, drawing, title, etc.). They are specified with the same structures as plotset or fontset (except for 'plotmargin'), or with the corresponding named arguments.

The first argument, name, is the name of the style feature:

ıres

figurestyle(name, style) changes the specified style. The style can be specified with a style structure, like what is returned by plotset or fontset, or with named arguments. Settings which are not specified keep their default values.

With a single argument, figurestyle(name) returned the current specified style.

The value for 'plotmargin' is a structure which describes the margin width around plots or subplots. It contains the following fields:

Value
left margin in multiple of a digit width
right margin in multiple of a digit width
top margin in multiple of line height
bottom margin in multiple of line height
see below
see below
see below

The fonts the widths are based on are the title font for Top, and the label font for the other fields. When an alternative y scale is used with altscale, the width of the right margin is based on Left instead of Right.

If field CenteredLabelWidth is larger than 0, it specifies the width of an additional margin (in multiple of a digit width) where the label of the Y axis is displayed, centered vertically. If field CenteredLabelHeight is larger than 0, it specifies the height of an additional bottom margin (in multiple of a line height) where the label of the X axis is displayed, centered horizontally. The default location of axis labels is at the end of the tick labels.

If field FixedControlVPos is false, controls (buttons, sliders etc.) are centered vertically in the subplot content area, or can be scrolled vertically by the user if they exceed the available space. If it is true, controls are aligned at the top and cannot be scrolled.

Considered as a whole, styles should be chosen such that they provide enough contrast to make all features visible. In particular, the font color should be changed when a dark background is selected. Some combinations, such as red on green, are difficult to distinguish for color-blind persons.

In Sysquake, figurestyle should not be used in figure draw handlers, because it applies to all subplots. It should typically be placed in init or menu handlers. To change the default figure styles which are used in all figures unless they are overridden by figurestyle, defaultstyle should be called instead.

### **Example**

Blue appearance with different dark shades for the backgrounds, and large fonts.

```
figurestyle('winbg', FillColor='#002');
figurestyle('plotbg', FillColor='#005');
figurestyle('legend', FillColor='#00a');
figurestyle('draw', Size=18, LineWidth=4, Color='#88f');
figurestyle('grid', Size=18, LineWidth=2, Color='#66c');
figurestyle('xygrid', LineWidth=2, Color='#66c');
figurestyle('frame', LineWidth=3, Color='#44f');
figurestyle('figfont', Size=20, Color='white');
figurestyle('controlfont', Size=20, Color='white');
figurestyle('legendfont', Size=20, Color='white');
figurestyle('titlefont', Size=32, Bold=true, Color='white');
figurestyle('tickfont', Size=18, Color='white');
figurestyle('labelfont', Size=18, Color='white');
```

#### See also

subplotstyle, plotset, plotfont, plotoption

### fontset

Options for fonts.

## **Syntax**

```
options = fontset
options = fontset(name1=value1, ...)
```

```
options = fontset(name1, value1, ...)
options = fontset(options0, name1, value1, ...)
```

## **Description**

fontset(name1, value1,...) creates the font description used by text. Options are specified with name/value pairs, where the name is a string which must match exactly the names in the table below. Case is significant. Alternatively, options can be given with named arguments. Options which are not specified have a default value. The result is a structure whose fields correspond to each option. Without any input argument, fontset creates a structure with all the default options. Options can also be passed directly to text or math as named arguments.

When its first input argument is a structure, fontset adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options (empty arrays mean "automatic"):

Name	Default	Meaning
Font	, ,	font name
Size	10	character size in points
Bold	false	true for bold font
Italic	false	true for italic font
Underline	false	true for underline characters
Color	[0,0,0]	text color

The default font is used if the font name is not recognized. The color is specified as an empty array (black), a scalar (gray) or a 3-element vector (RGB) of class double (0=black, 1=maximum brightness) or uint8 (0=black, 255=maximum brightness).

## **Examples**

Default font:

```
fontset
  Font: ''
  Size: 10
  Bold: false
  Italic: false
  Underline: false
  Color: real 1x3
```

Named argument directly in text:

```
text(0, 0, 'Text', Font='Times', Italic=true, Bold=true)
```

### See also

text

# fplot

Function plot.

## **Syntax**

```
fplot(fun)
fplot(fun, limits)
fplot(fun, limits, style)
fplot(fun, limits, style, id)
fplot(fun, limits, style, id, p1, p2, ...)
```

## **Description**

Command fplot(fun, limits) plots function fun, specified by its name as a string, a function reference, or an inline or anonymous function. The function is plotted for x between limit(1) and limit(2); the default limits are [-5,5].

The optional third and fourth arguments are the same as for all graphical commands.

Remaining input arguments of fplot, if any, are given as additional input arguments to function fun. They permit to parameterize the function. For example fplot('fun',[0,10],'',-1,2,5) calls fun as y=fun(x,2,5) and displays its value for x between 0 and 10.

# **Examples**

fplot(@(x)  $(x+0.3)^2+a*exp(-3*x^2)$ , [-2,3], 'r', 1);

### See also

plot, inline, operator @

# image

Raster RGB or grayscale image.

## **Syntax**

```
image(gray)
image(red, green, blue)
image(rgb)
image(..., [xmin, xmax, ymin, ymax])
image(..., mode)
image(..., id)
```

## **Description**

image displays a raster image (an image defined by a rectangular array of patches of colors called *pixels*). The raster image can be either grayscale or color. A grayscale image is defined by a double matrix of pixel values in the range 0 (black) to 1 (white), by a uint8 matrix in the range 0 (black) to 255 (white), or by a uint16 matrix in the range 0 (black) to 65535 (white). A color image is defined by three matrices of equal size, corresponding to the red, green, and blue components, or by an array with three planes along the 3rd dimension. Each component is defined between 0 (black) to 1 (maximum intensity) with double values, between 0 (black) to 255 (maximum intensity) with uint8 values, or between 0 (black) and 65535 (maximum intensity) with uint16 values. If a colormap has been defined, grayscale image rendering uses it.

The position is defined by the the minimum and maximum along the coordinates horizontal and vertical axes. The raster image is scaled to fit. The first line of the matrix or matrices is displayed at the top. The position can be specified an argument [xmin,xmax,ymin,ymax]; by default, [0,size(im,2),0,size(im,1)] where im stands for the image array or one of its RGB components.

If mode is 'e', the raster image is scaled down such that each pixel has the same size; otherwise, the specified position is filled with the raster image. You should use 'e' when you want a better quality, but do not add other elements in the figure (such as marks or lines) and do not have interaction with the mouse.

Pixels on the screen are interpolated using the bilinear method if mode is '1', and the bicubic method if mode is '3'.

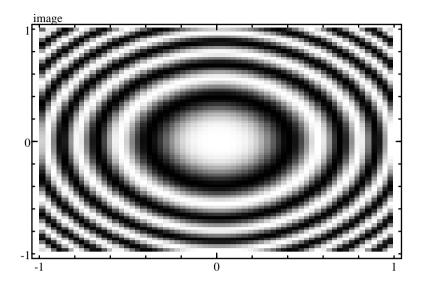


Figure 6.7 Example of image

## **Examples**

Two ways to display a table of 10-by-10 random color cells (see Fig. 6.7):

```
image(rand(10), rand(10), rand(10));
image(rand(10, 10, 3));
```

A ramp of gray shades:

```
image(uint8(0:255));
```

Operator : and function meshgrid can be used to create the x and y coordinates used to display a function z(x,y) as an image.

```
(X, Y) = meshgrid(-pi:0.1:pi);

Z = cos(X.^2 + Y.^2).^2;

image(Z, [-1,1,-1,1], '3');
```

### See also

contour, quiver, colormap, pcolor

### label

Plot labels.

## **Syntax**

```
label(label_x)
label(label_x, label_y)
label(label_x, label_y, label_y2)
```

## **Description**

label(label\_x, label\_y) displays labels for the x and y axes. Its arguments are strings. The label for the y axis may be omitted.

When an alternative y scale is used with altscale, its label can be specified with a third argument.

For a dB scale, an additional label [dB] is automatically displayed below the text specified by label\_y; it is not displayed if there is no label\_y (or an empty label\_y). If label\_y is a single-space string, it is replaced by [dB] for a dB scale (i.e. [dB] is aligned correctly with the top of the figure).

With plotoption math, labels can contain MathML or LaTeX.

## **Examples**

```
step(1,[1,2,3,4]);
label('t [s]', 'y [m]');

With literal strings, the command syntax may be more convenient:
label Re Im;
dB scale with only a [dB] label:
scale logdb;
bodemag(1, [1, 2, 3]);
label('', '');
```

### See also

text, legend, title, ticks, altscale, plotoption

# legend

Plot legend.

## **Syntax**

```
legend(str)
legend(str, style)
```

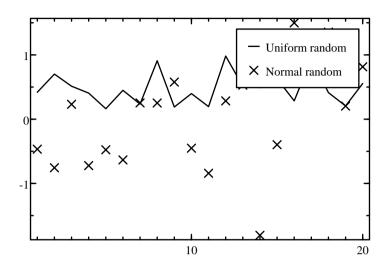


Figure 6.8 Example of legend

## **Description**

legend(str,style) displays legends for styles defined in string style. In string str, legends are separated by linefeed characters \n. Legends are displayed at the top right corner of the figure in a frame. All styles are permitted: symbols, lines, and filling. They are recycled if more legends are defined in str. If str is empty, no legend is displayed.

With a single input argument, legend(str) uses the default style 'k'.

With plotoption math, legend lines in first argument can contain MathML or LaTeX.

## **Example**

Legend for two traces (see Fig. 6.8).

```
plot(1:20, [rand(1,20); randn(1,20)], '_x');
legend('Uniform random\nNormal random', '_x');
```

### See also

label, ticks, title, plotoption

### line

Plot lines.

### **Syntax**

```
line(A, b)
line(V, P0)
line(..., style)
line(..., style, id)
```

## **Description**

line displays one or several straight line(s). Each line is defined by an implicit equation or an explicit, parametric equation.

**Implicit equation:** Lines are defined by equations of the form  $a_1x + a_2y = b$ . The first argument of line is a matrix which contains the coefficients  $a_1$  in the first column and  $a_2$  in the second column. The second argument is a column vector which contains the coefficients b.

**Explicit equations:** Lines are defined by equations of the form  $P = P_0 + \lambda V$  where  $P_0$  is a point of the line, V a vector which defines its direction, and  $\lambda$  a real parameter. The first argument of line is a matrix which contains the coefficients  $v_x$  in the first column and  $v_y$  in the second column. The second argument is a matrix which contains the coefficients  $x_0$  in the first column and  $y_0$  in the second column.

In both cases, each row corresponds to a different line. If one of the arguments has one row and the other has several (or none), the same row is duplicated to match the other size.

In figures with a logarithmic scale, only horizontal and vertical lines are allowed.

The optional third and fourth arguments are the same as for all graphical commands.

In mouse handlers, \_x0 and \_y0 correspond to the projection of the mouse position onto the line; \_nb is the index of the line in A and b, and \_ix is empty.

# **Examples**

```
Vertical line at x=5:
line([1,0],5)

Draggable horizontal blue lines at y=2 and y=3:
line([0,1], [2;3], 'b', 1)

The same lines with named arguments:
line([0,1], [2;3], Color='blue', id=1)
```

#### See also

plot, circle

#### math

Display MathML or LaTeX in a figure.

## **Syntax**

```
math(x, y, string)
math(x, y, string, justification)
math(..., font)
math(..., id=id)
```

## **Description**

With three arguments, math(x,y,string) renders a string as MathML or LaTeX, centered at the specified position. The third argument is assumed to be MathML unless it starts with a dollar character; in that case, it is converted to MathML as if it was processed by latex2mathml.

An optional fourth argument specifies how the MathML equation should be aligned with respect to the position (x,y). It is a string of one or two characters from the following set:

# Char. Alignment Center (may be omitted) Left Right Top Bottom

For instance, 'l' means that the MathML equation is displayed to the right of the given position and is centered vertically, and 'rt', that the equation is to the bottom left of the given position.

An optional trailing argument specifies the font. It is a structure which is typically created with fontset; but only the base font size is used. Alternatively, the base font size can be specified with a named argument.

An ID can be specified with a named argument (not with a normal, unnamed argument).

The following MathML elements are supported: math, merror, mfenced, mfrac, mi, mn, mo, mpadded, mphantom, mroot, mrow, msqrt, mspace, msub, msubsup, msup, mtable, mtd, mtext, mtr.

## **Examples**

```
math(0, 0, mathml([1,pi,1e30]));
math(0, 0, mathml(1e-6, Format='e', NPrec=2), Size=20);
math(0, 0, '$\\rho=\\sqrt{x^2+y^2}$');
```

#### See also

text, mathml, latex2mathml, fontset

# pcolor

Pseudocolor plot.

## **Syntax**

```
pcolor(C)
pcolor(X, Y, C)
pcolor(..., style)
pcolor(..., style, id)
```

## **Description**

Command pcolor(C) displays a pseudocolor plot, i.e. a rectangular array where the color of each cell corresponds to the value of elements of 2-D array C. These values are real numbers between 0 and 1. The color used by pcolor depends on the current color map; the default is a grayscale from black (0) to white (1).

pcolor(X,Y,C) displays the plot on a grid whose vertex coordinates are given by arrays X and Y. Arrays X, Y and C must all have the same size.

With an additional string input argument, pcolor(..., style) specifies the style of the lines drawn between the cells.

The following argument, if it exists, is the ID used for interactivity. During interactive manipulation, the index obtained with \_ix corresponds to the corner of the patch under the mouse with the smallest index.

## **Example**

```
use colormaps;
n = 11;
(x, y) = meshgrid(1:n);
phi = pi/8;
X = x*cos(phi)-y*sin(phi);
Y = x*sin(phi)+y*cos(phi);
C = magic(n)/n^2;
```

```
pcolor(X, Y, C, 'k');
colormap(blue2yellow2redcm);
plotoption noframe;
```

#### See also

plot, colormap, image

# plot

Generic plot.

#### **Syntax**

```
plot(y)
plot(x, y)
plot(..., style)
plot(..., style, id)
```

# **Description**

The command plot displays graphical data in the current figure. The data are given as two vectors of coordinates x and y. If x is omitted, its default value is 1:size(y,2). Depending on the style, the points are displayed as individual marks or are linked with lines. The stairs style ('s') can be used to link two successive points with a horizontal line followed by a vertical line. If x and y are matrices, each row is considered as a separate line or set of marks; if only one of them is a matrix, the other one, a row or column vector, is replicated to match the size of the other argument.

The optional fourth argument is an identification number which is used for interactive manipulation. It should be equal or larger than 1. If present and a mousedown, mousedrag and/or mouseup handler exists, the position of the mouse where the click occurs is mapped to the closest graphical element which has been displayed with an ID; for the command plot, the closest point is considered (lines linking the points are ignored). If such a point is found at a small distance, the built-in variables  $_{x}0$ ,  $_{y}0$ , and  $_{z}0$  are set to the position of the point before it is moved; the variable  $_{i}1$  is set to the ID as defined by the command plot; the variable  $_{n}1$  is set to the number of the row, and the variable  $_{i}1$  is set to the index of the column of the matrix x and y.

# **Examples**

Sine between 0 and  $2\pi$ :

A complete SQ file for displaying a red triangle whose corners can be moved interactively on Sysquake:

```
variables x, y
                  // x and y are 1-by-3 vectors
init (x,y) = init // init handler
figure "Triangle"
  draw drawTri(x, y)
  mousedrag (x, y) = dragTri(x, y, _ix, _x1, _y1)
functions
{@
function (x,y) = init
  x = [-1,1,0];
  y = [-1, -1, 2];
  subplots('Triangle');
function drawTri(x,y)
  scale('equal', [-3, 3, -3, 3]);
  plot(x, y, FillColor='red', id=1);
function (x, y) = dragTri(x, y, ix, x1, y1)
  if isempty(ix)
    cancel; // not a click over a point
  end
  x(ix) = x1;
  y(ix) = y1;
@}
```

#### See also

fplot, line, circle

# plotoption

Set plot options.

```
plotoption(str1, str2, ...)
plotoption opt1 opt2 ...
```

plotoption sets the initial value of the plot options the user can change. Its arguments, character strings, can each take one of the following values.

<sup>&#</sup>x27;frame' Rectangular frame with tick marks and a white background around the plot (default).

<sup>&#</sup>x27;noframe' No frame, no tickmarks, no white background.

<sup>&#</sup>x27;label' Subplot name above the frame (default).

<sup>&#</sup>x27;nolabel' No subplot name.

<sup>&#</sup>x27;legend' Legend (if it has been set with legend).

<sup>&#</sup>x27;nolegend' Hidden legend.

<sup>&#</sup>x27;trlegend' Legend in top right corner (default).

<sup>&#</sup>x27;tllegend' Legend in top left corner.

<sup>&#</sup>x27;brlegend' Legend in bottom right corner.

<sup>&#</sup>x27;bllegend' Legend in bottom left corner.

<sup>&#</sup>x27;margin' Margin for title and labels (default).

<sup>&#</sup>x27;nomargin' No margin.

<sup>&#</sup>x27;math' MathML or LaTeX rendering in title, label, legend, and controls like button and slider. The string (or substring in legend) is parsed as MathML if the first character is '<', or as LaTeX if it is '\$'. Otherwise, it is displayed as if it was the text content of an <mtext> element, to guarantee that there is no font mismatch with mathematical expressions.

<sup>&#</sup>x27;nomath' No math (default).

<sup>&#</sup>x27;xticks' Ticks and labels for the x axis.

<sup>&#</sup>x27;noxticks' No ticks and labels for the x axis.

<sup>&#</sup>x27;yticks' Ticks and labels for the y axis.

<sup>&#</sup>x27;noyticks' No ticks and labels for the y axis.

<sup>&#</sup>x27;xyticks' Ticks and labels for the x and y axes (default).

<sup>&#</sup>x27;noxyticks' No ticks and labels for the x and y axes.

<sup>&#</sup>x27;xgrid' Grid of vertical lines for the x axis.

```
'noxgrid' No grid for the x axis.
```

## **Examples**

Display of a photographic image without frame:

```
plotoption noframe;
image(photo);

Math in a title:

plotoption math;
title '$\\hbox{Solution of}\\;\\dot{x}=f(x,t)$';
```

#### See also

figurestyle, scale, legend

# plotset

Options for plot style.

```
options = plotset
options = plotset(name1, value1, ...)
options = plotset(options0, name1, value1, ...)
```

<sup>&#</sup>x27;ygrid' Grid of horizontal lines for the y axis.

<sup>&#</sup>x27;noygrid' No grid for the y axis.

<sup>&#</sup>x27;xygrid' Grid of vertical and horizontal lines for the x and y axes.

<sup>&#</sup>x27;noxygrid' No grid for the x and y axes (default).

<sup>&#</sup>x27;grid' Normal details for grids displayed by sgrid, zgrid, etc. (default).

<sup>&#</sup>x27;nogrid' Removal of grids displayed by sgrid, zgrid, etc.

<sup>&#</sup>x27;fullgrid' More details for grids displayed by sgrid, zgrid, etc.

<sup>&#</sup>x27;fill3d' In 3D graphics, zoom in so that the bounding box fills the figure.

plotset(name1, value1,...) creates the style option argument used by functions which display graphics, such as plot and line. Options are specified with name/value pairs, where the name is a string which must match exactly the names in the table below. Case is significant. Options which are not specified have a default value. The result is a structure whose fields correspond to each option. Without any input argument, plotset creates a structure with the default style.

When its first input argument is a structure, plotset adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

Name	Default	Meaning
ArrowEnd	false	arrow at end
ArrowStart	false	arrow at start
Color	[]	line color
Fill	false	fill
FillColor	[]	filling color
LineStyle	, ,	line style
LineWidth	[]	line width
Marker	, ,	marker style
MarkerEdgeColor	[]	marker edge color
MarkerFaceColor	[]	marker face (filling) color
Stairs	false	stairs
Stems	false	stems

Colors are specified by value or by name. An empty array means the default color (usually black for lines and marker edge, none for filling, and white for marker face). A scalar number represents a shade of gray, an array of 3 numbers an RGB color. An additional element (last element of array of 2 or 4 numbers) represents the alpha component (transparency) where 0 is completely transparent; it is ignored on some platforms. Color type can be uint8 from 0 to 255, uint16 from 0 to 65535, or single or double from 0 to 1. In all cases, 0 represents black and the largest value, the maximum brightness.

Color names can be one of the following:

Name	Value as uint8
'black'	[0,0,0]
'blue'	[0,0,255]
'green'	[0,128,0]
'cyan'	[0,255,255]
'red'	[255,0,0]
'magenta'	[255,0,255]
'yellow'	[255,255,0]
'white'	[255,255,255]
'aqua'	[0,255,255]
'darkgray'	[169,169,169]
'darkgrey'	[169,169,169]
'darkgreen'	[0,64,0]
'fuchsia'	[255,0,255]
'gray'	[128,128,128]
'grey'	[128,128,128]
'lime'	[0,255,0]
'maroon'	[128,0,0]
'navy'	[0,0,128]
'olive'	[128,128,0]
'orange'	[255,165,0]
'purple'	[128,0,128]
'silver'	[192,192,192]
'teal'	[0,128,128]

Option LineStyle is an empty string for the default line style (solid line unless FillColor is set), or one of the following one-character strings:

```
Dash Pattern LineStyle
solid '_' (underscore)
dashed '-' (hyphen)
dotted ':'
dash-dot '!'
hidden ' '(space)
```

Option Marker is an empty string for the default symbol (usually no symbol, or a cross for plotroots), or one of the following strings:

```
Marker Shape
                 Marker
                  ' '(space)
none
point
                  0'
circle
                  'x'
cross
                  '+'
plus
star
                  , ^ ,
triangle up
triangle down
                  'v'
triangle left
                  ' < '
triangle right
                  '>'
                  '[]' or '['
square
                  '<>'
diamond
```

An explicit Fill=true is usefull only for filling with the default color or colors, with functions such contour. Otherwise, specifying a filling color with FillColor implies Fill=true.

When Stems is true, a marker is drawn for each point and is linked with a vertical line which connects it to the origin (in 2D plots, y=0 for linear scale or y=1 for logarithmic scale; in 3D plots, z=0). In polar plots, stems connects points to x=y=0.

Functions which support multiple styles, such as plot where each trace can have a different style, accept a structure array or a list of structures. If there are less elements in the style array or list than there are traces to plot, styles are recycled, restarting from the first one. If there are too many, superfluous styles are ignored.

When Stairs is true, for functions wihich support it, points are connected with a horizontal line followed by a vertical line.

#### **Examples**

Default options:

```
plotset
   ArrowEnd: false
   ArrowStart: false
   Color: []
   FillColor: []
   LineStyle: ''
   LineWidth: []
   Marker: ''
   MarkerEdgeColor: []
   MarkerFaceColor: []
```

Plot of 5 random lines defined by 10 points each, odd and even ones with different styles:

```
data = rand(5, 10);
styleOdd = plotset(ArrowStart=true,
```

```
LineWidth=2,
Color='red',
Size=0);
styleEven = plotset(ArrowEnd=true,
LineWidth=2,
Size=10,
Color='blue',
MarkerEdgeColor='black',
MarkerFaceColor='yellow');
plot(data, {styleOdd, styleEven});
```

Multiple styles can also be built directly as a structure array, without plotset; missing fields take their default values.

```
styles = {
   LineWidth=2, Color='red';
   LineStyle='-', Color='blue'
};
plot(data, styles);
```

#### See also

plot, plotoption, figurestyle

# polar

Polar plot.

## Syntax

```
polar(theta, rho)
polar(..., style)
polar(..., style, id)
```

# **Description**

Command polar displays graphical data in the current figure with polar coordinates. The data are given as two vectors of coordinates theta (in radians) and rho. Depending on the style, the points are displayed as individual marks or are linked with lines. If x and y are matrices, each row is considered as a separate line or set of marks; if only one of them is a matrix, the other one, a vector, is reused for each line.

Automatic scaling is performed the same way as for cartesian plots after polar coordinates have been converted. The figure axes, ticks and grids are specific to polar plots. Polar plots can be mixed with other graphical commands based on cartesian coordinates such as plot, line and circle.

## **Example**

```
theta = 0:0.01:20*pi;
rho = exp(0.1 * theta) .* sin(5 * theta);
polar(theta, rho);
```

#### See also

plot

# quiver

Quiver plot.

## **Syntax**

```
quiver(x, y, u, v)
quiver(u, v)
quiver(..., scale)
quiver(..., style)
```

# **Description**

quiver(x,y,u,v) displays vectors (u,v) starting at (x,y). If the four arguments are matrices of the same size, an arrow is drawn for each corresponding element. If x and y are vectors, they are repeated: x is transposed to a row vector if necessary and repeated to match the number of rows of u and v; and y is transposed to a column vector if necessary and repeated to match their number of columns. The absolute size of arrows is scaled with the average step of the grid given by x and y, so that they do not overlap if the grid is uniform.

If x and y are missing, their default values are [1,2,...,m] and [1,2,...,n] respectively, where m and n are the number of rows and columns of u and v.

With a 5th (or 3rd) argument, quiver(...,scale) multiplies the arrow lengths by the scalar number scale. If scale is zero, arrows are not scaled at all: u and v give directly the absolute value of the vectors.

With a 6th (or 4th) string argument, quiver(...,style) uses the specified style to draw the arrows.

## **Example**

Force field; complex numbers are used to simplify computation.

```
scale equal;
z = fevalx(@plus, -5:0.5:5, 1j*(-5:0.5:5)');
z0 = 0.2+0.3j;
f = 1+20*sign(z-z0)./(max(abs(z-z0).^2,3));
x = real(z);
y = imag(z);
u = real(f);
v = imag(f);
quiver(x, y, u, v);
```

#### See also

plot, image contour

#### scale

Set the scale.

## **Syntax**

```
scale([xmin,xmax,ymin,ymax])
scale([xmin,xmax,ymin,ymax,zmin,zmax])
scale(features)
scale(features, usersettablefeatures)
scale(features, [xmin,xmax,ymin,ymax])
scale(features, usersettablefeatures, [xmin,xmax,ymin,ymax])
sc = scale
(sc, type) = scale
```

# **Description**

Without output argument, the scale command, which should be placed before any other graphical command, sets the scale and scale options. The last parameter contains the limits of the plot, either for both x and y axes or only for the x axis in 2D graphics, or for x, y and z axes for 3D graphics. The limits are used only if the user has not changed them by zooming.

The first parameter(s) specify some properties of the scale, and which one can be changed by the user. There are two ways to specify them: with a string or with one or two integer numbers. The recommended way is with a string. The list below enumerates the possible values.

<sup>&#</sup>x27;equal' Same linear scale for x and y axes. Typically used for representation of the complex plane, such as the roots of a poly-

nomial or a Nyquist diagram. For 3D graphics, same effect as daspect([1,1,1]).

- 'pixel' Pixel (unit) linear scale for x and y axes. Used for diagrams which cannot be scaled, such as block diagrams, Venn diagrams, or special use interface. The y axis is always oriented upward.
- 'lock' See below.
- 'linlin' Linear scale for both axes.
- 'linlog' Linear scale for the x axis, and logarithmic scale for the y axis.
- 'loglin' Logarithmic scale for the x axis, and linear scale for the y axis.
- 'loglog' Logarithmic scale for both axes.
- 'lindb' Linear scale for the x axis, and dB scale for the y axis.
- 'logdb' Logarithmic scale for the x axis, and dB scale for the y axis.
- 'lindb/logdb' Linear scale for the x axis, and dB scale for the y axis. The user can choose a logarithmic scale for the x axis, and a logarithmic or linear scale for the y axis.
- 'loglog/set' Logarithmic scale for the x and y axes, without possibility for the user to change them.

The last-but-one setting shows how to enable the options the user can choose in Sysquake. The setting and the enabled options are separated by a dash; if a simple setting is specified, the enabled options are assumed to be the same. Enabling dB always permits the user to choose a logarithmic or linear scale, and enabling a logarithmic scale always permits to choose a linear scale. The 'equal' option cannot be combined with anything else. Changing the options in subsequent redraws is ignored, because options are under the user control.

The last setting ending with /set shows how to force options without letting the user override them. In this case, options can be changed during redraws. SQ files with customs ways to change the kind of scale must use this method.

When the properties are specified with one or two integer numbers, each bit corresponds to a property. Only the properties in bold in the table below can be set by the user, whatever the setting is.

## Bit Meaning

- 0 **log x**
- 2 tick on x axis
- 3 grid for x axis
- 4 labels on x axis
- 6 **log y**
- 7 **dB y**
- 8 tick on y axis
- 9 grid for y axis
- 10 labels on y axis
- 12 same scale on both axes
- 13 minimum grid
- 14 maximum grid

scale lock locks the scale as if the user had done it by hand. It fixes only the initial value; the user may change it back afterwards.

The scale is usually limited to a range of 1e-6 for linear scales and a ratio of 1e-6 for logarithmic scales. This avoids numeric problems, such as when a logarithmic scale is chosen and the data contain the value 0. In some rare cases, a large scale may be required. The 'lock' option is used to push the limits from 1e-6 to 1e-24 for both linear and logarithmic scales. A second argument must be provided:

```
scale('lock', [xmin,xmax,ymin,ymax]);
```

The command must be used in a draw handler (or from the command line interface). To add other options, use a separate scale command:

```
scale logdb;
scale('lock', [1e-5, 1e8, 1e-9, 1e9]);
```

The scale is locked, and the user may not unlock it. In the example above, note also that a single string argument can be written without quote and parenthesis if it contains only letters and digits.

With output arguments, scale returns the current scale as a vector [xmin, xmax, ymin, ymax]. If the scale is not fixed, the vector is empty. If only the horizontal scale is set, the vector is [xmin, xmax]. During a mouse drag, both the horizontal and vertical scales are fixed. The values returned by scale reflect the zoom chosen by the user. They can be used to limit the computation of data displayed by plot to the visible area. The optional second output argument type tells whether a linear or a logarithmic scale is set for axis x and y; it is a string such as 'linlin' or 'loglin'.

# **Examples**

Here are some suggestions for the most usual graphics:

```
Time response
               (default linlin is fine)
               scale loadb
Bode mag
               scale loglin
Bode phase
               scale('lindb/logdb',[0,pi/Ts])
D bode mag
D bode phase
               scale('linlin/loglin',[0,pi/Ts])
Poles
               scale equal
D poles
               scale('equal',[-1,1,-1,1])
               scale('equal',[-1.5,1.5,-1.5,1.5])
Nyquist
Nichols
               scale lindh
 Use of scale to display a sine in the visible x range:
scale([0,10]);
               % default x range between 0 and 10
sc = scale;
                % maybe changed by the user (1x2 or 1x4)
xmin = sc(1);
xmax = sc(2);
x = xmin + (xmax - xmin) * (0:0.01:1);
               % 101 values between xmin and xmax
y = \sin(x);
plot(x, y);
```

#### See also

plotoption, scalefactor

#### scalefactor

Change the scale displayed in axis ticks and labels.

#### Syntax

```
scalefactor(f)
f = scalefactor
```

# **Description**

scalefactor(f) sets the factor used to display the ticks and the labels. Its argument f can be a vector of two or three real positive numbers to set separately the x, y, and z axes, or a real positive scalar to set the same factor for all axes. scalefactor([fx,fy]) is equivalent to scalefactor([fx,fy,1]). The normal factor value is 1, so that the ticks correspond to the graphical contents. With a different factor, the contents are displayed with the same scaling, but the ticks and labels are changed as if the graphical data had been scaled by the factor. For instance, you can plot data in radians (the standard angle unit in LME) and display ticks and labels in degrees by using a factor of 180/pi.

With an output argument, scalefactor gives the current factors as a 2-elements vector.

#### **Example**

Display the sine with a scale in degrees:

```
phi = 0:0.01:2*pi;
plot(phi, sin(phi));
scalefactor([180/pi, 1]);
```

#### See also

scale, plotoption

#### scaleoverview

Set the scale overview rectangle.

#### **Syntax**

```
scaleoverview([xmin,xmax,ymin,ymax])
scaleoverview([xmin,xmax,ymin,ymax],'xy')
scaleoverview([xmin,xmax],'x')
scaleoverview([ymin,ymax],'y')
```

# **Description**

scaleoverview sets the limits of a rectangular region used to provide an overview of the scale used in another plot. Typically, the same data are displayed in two subplots: one with a large, fixed displayed area (set with scale) with a smaller scale overview rectangle set with scaleoverview, and one with a smaller displayed area (set with scale) which matches the limits set with scaleoverview in the first plot. In Sysquake, scale synchronization is used to keep both subplots synchronized when the user zooms or drags the data in the second subplot or manipulates directly the scale overview rectangle.

By default, limits on axis x and y are provided. A second argument can specify which axis has limits: 'xy' (default), 'x' or 'y' (then the first argument is an array of two elements).

#### See also

scale

# subplotstyle

Subplot style.

## **Syntax**

```
subplotstyle(name, style)
style = subplotstyle(name)
```

## **Description**

subplotstyle sets or gets the style of the current subplot. In Sysquake's SQ files, it should be used in draw handlers. It has the same arguments as figurestyle, which handles the settings globally for all subplots, or the default settings when both figurestyle and subplotstyle are used.

## **Example**

```
subplot 211;
subplotstyle('plotbg', FillColor='yellow');
subplotstyle('frame', LineWidth=2);
step(1, 1:3);
subplot 212;
subplotstyle('plotbg', FillColor='orange');
step(1, 1:4);
```

#### See also

figurestyle, plotset, plotfont, plotoption

#### text

Display text in a figure.

# **Syntax**

```
text(x, y, string)
text(x, y, string, justification)
text(..., font)
text(..., id=id)
```

# **Description**

With three arguments, text(x,y,string) displays a string centered at the specified position. An optional fourth argument specifies how the string should be aligned with respect to the position (x,y). It is a string of one or two characters from the following set:

# Char. Alignment c Center (may be omitted) l Left r Right t Top b Bottom

For instance, 'l' means that the string is displayed to the right of the given position and is centered vertically, and 'rt', that the string is to the bottom left of the given position.

An optional trailing argument specifies the font, size, type face, and color to use. It is a structure which is typically created with fontset. Alternatively, named arguments can be used directly, without fontset.

An ID can be specified with a named argument (not with a normal, unnamed argument).

# **Examples**

A line is drawn between (-1,-1) and (1,1) with labels at both ends.

```
plot([-1,1], [-1,1]);
text(-1,-1, 'p1', 'tr');
text(1, 1, 'p2', 'bl');

Text with font specification:

font = fontset(Font='Times',
    Bold=true,
    Size=18,
    Color=[1,0,0]);
text(1.1, 4.2, 'Abc', font);

Same font with named arguments:

text(1.1, 4.2, 'Abc', font,
    Font='Times',
    Bold=true,
    Size=18,
    Color=[1,0,0]);
```

#### See also

label, fontset, sprintf

#### tickformat

Subplot tick format.

## **Syntax**

```
tickformat(axis, format)
```

## **Description**

tickformat(axis, format) specifies the format to be used for tick labels. The first argument, axis, specified which axis is affected: it is 1 or 'x' for the first axis (horizontal) or 2 or 'y' for the second axis (vertical, on the left or the right depending on the last call to altscale if any). Second argument, format, is a string similar to the first argument of sprintf. Only numeric formats are supported (%d, %e, %f, %g, %h, %i, %k, %n, %o, %P, %x), with their options, width and precision. Double % gives a single %, and other characters are used literally.

The format is not used if ticks are specified with function ticks.

#### **Examples**

General format with a unit for the x axis, and fixed format with two fractional digits for the y axis:

```
step(1, [1, 2, 3, 4]);
tickformat('x', '%g h');
tickformat('y', '%.2f');
```

#### See also

ticks, sprintf

#### ticks

Subplot ticks and tick labels.

## **Syntax**

```
ticks(axis, majorTicks)
ticks(axis, majorTicks, minorTicks)
ticks(axis, majorTicks, minorTicks, tickLabels)
```

# **Description**

ticks replaces default ticks (small scale marks along the plot frame and their labels outside the frame) with custom ones.

ticks(axis,majorTicks) specifies the value of major ticks (large ones). The first argument, axis, specified which axis is affected: it

is 1 or 'x' for the first axis (horizontal) or 2 or 'y' for the second axis (vertical, on the left or the right depending on the last call to altscale if any). Second argument, majorTicks, is an array of values where ticks are displayed; the same scaling as the one applied to the plot contents is used.

With a third argument, ticks(axis,majorTicks,minorTicks) also displays minor ticks (smaller ones, typically used with a finer spacing) specified by array minorTicks.

With a fourth argument, ticks(axis,majorTicks,minorTicks,labels) displays labels at the position of major ticks. Labels are given as a string of linefeed-separated substrings, such as 'one\ntwo. If more values are specified for major ticks than for labels, labels are reused, starting from the first one. Superfluous labels are ignored. If no minor tick is displayed, argument minorTicks is optional.

Values out of range are not displayed. If axis labels are specified with function label, tick labels which would overlap are not displayed. 3D plots have always default ticks.

#### **Examples**

Bar plot where bars correspond to months:

```
bar([2,4,3,6]);
ticks('x', 1:4, 'Jan\nFeb\nMar\nApr');

Tick labels with units and axis label:

scale([0, 10]);
step(1, [1, 2, 3, 4]);
majorTicks = 0:2:10;
minorTicks = 0:0.5:10;
labels = sprintf('%g s\n', majorTicks);
ticks('x', majorTicks, minorTicks, labels);
ticks('y', 0:0.1:1);
label Time;

Plot without any tick and label:

step(1, [1, 2, 3, 4]);
ticks('x', []);
ticks('y', []);
```

#### See also

label, tickformat, legend, title, text, sprintf

#### title

Subplot title.

#### **Syntax**

title(string)

## **Description**

title(string) sets or changes the title of the current subplot.
With plotoption math, the title can contain MathML or LaTeX.

#### See also

label, legend, ticks, text, sprintf, plotoption

# 6.41 3D Graphics

Three-dimension graphic commands enable the representation of objects defined in three dimensions x, y and z on the two-dimension screen. The transform from the 3D space to the screen is performed as if there were a virtual camera in the 3D space with a given position, orientation, and angle of view (related to the focal length in a real camera).

# **Projection**

The projection is defined by the following parameters:

**Target point** The target point is a 3D vector which defines the position where the camera is oriented to.

**Projection kind** Two kinds of projections are supported: orthographic and perspective.

**View point** The view point is a 3D vector which defines the position of the camera. For orthographic projection, it defines a direction independent from the target position; for perspective projection, it defines a position, and the view orientation is defined by the vector from view point to target point.

**Up vector** The up vector is a 3D vector which fixes the orientation of the camera around the view direction. The projection is such that the up vector is in a plane which is vertical in the 2D projection. Changing it makes the projection rotate around the image of the target.

**View angle** The view angle defines the part of the 3D space which is projected onto the image window in perspective projections. It is zero in orthographic mode.

All of these parameters can be set automatically. Here is how the whole projection and scaling process is performed:

- Scale data separately along each direction according to daspect
- Find bounding box of all displayed data, or use limits set with scale
- Find radius of circumscribed sphere of bounding box
- If the target point is automatic, set it to the center of the bounding box; otherwise, use position set with camtarget
- If the view point is automatic, set it to direction [-3;-2;1] at infinity in orthographic mode, or in that direction with respect to the target point at a distance such that the view angle of the circumscribed sphere is about 6 degrees; otherwise, use position set with campos
- If the up vector is automatic, set it to [0,0,1] (vertical, pointing upward); otherwise, use position set with camup
- Compute the corresponding homogeneous matrix transform
- Set the base scaling factor so that the circumscribed sphere fits the display area
- Apply an additional zoom factor which depends on camva and camzoom

# **Surface shading**

Surface and mesh colors add information to the image, helping the viewer in interpreting it. Colors specified by the style argument also accepted by 2D graphical commands are used unchanged. Colors specified by a single-component value, RGB colors, or implicit, are processed differently whether lightangle and/or material have been executed, or not. In the first case, colors depend directly on the colors specified or the default value; in the second case, the Blinn-Phong reflection model is used with flat shading. In both cases, single-color values are mapped to colors using the current color map (set with colormap). Commands which accept a color argument are mesh, surf, and plotpoly.

#### **Direct colors**

If neither lightangle nor material has been executed, colors depend only on the color argument provided with x, y, and z coordinates. If the this argument is missing, color is obtained by mapping linearly the z coordinates to the full range of the current color map.

## **Blinn-Phong reflection model**

In the Blinn-Phong reflexion model, the color of a surface depends on the intrinsic object color, the surface reflexion properties, and the relative positions of the surface, the viewer, and light sources.

# camdolly

Move view position and target.

## **Syntax**

camdolly(d)

# **Description**

camdolly(d) translates the camera by 3x1 or 1x3 vector d, moving the target and the view point by the same amount.

#### See also

campan, camorbit, campos, camproj, camroll, camtarget, camup,
camva, camzoom

## camorbit

Camera orbit around target.

# **Syntax**

camorbit(dphi, dtheta)

camorbit(dphi,dtheta) rotates the camera around the target point by angle dphi around the up vector, and by angle dtheta around the vector pointing to the right of the projection plane. Both angles are given in radians. A positive value of dphi makes the camera move to the right, and a positive value of dtheta makes the camera move down.

#### See also

camdolly, campan, campos, camproj, camroll, camtarget, camup, camva, camzoom

## campan

Tilt and pan camera.

## **Syntax**

campan(dphi, dtheta)

# **Description**

campan(dphi,dtheta) pans the camera by angle dphi and tilts it by angle dtheta. Both angles are in radians. More precisely, the target point is changed so that the vector from view point to target is rotated by angle dphi around the up vector, then by angle dtheta around a "right" vector (a vector which is horizontal in view coordinates).

#### See also

camdolly, camorbit, campos, camproj, camroll, camtarget, camup,
camva. camzoom

# campos

Camera position.

#### **Syntax**

campos(p)
campos auto
campos manual
p = campos

campos(p) sets the view position to p. p is a 3D vector.

campos auto sets the view position to automatic mode, so that it follows the target. campos manual sets the view position to manual mode.

With an output argument, campos gives the current view position.

#### See also

camdolly, camorbit, campan, camproj, camroll, camtarget, camup, camva, camzoom

# camproj

Projection kind.

## **Syntax**

```
camproj(str)
str = camproj
```

# **Description**

camproj(str) sets the projection mode; string str can be either 'orthographic' (or 'o') for a parallel projection, or 'perspective' (or 'p') for a projection with a view point at a finite distance.

With an output argument, camproj gives the current projection mode.

#### See also

camdolly, camorbit, campan, campos, camroll, camtarget, camup, camva, camzoom

#### camroll

Camera roll around view direction.

```
camroll(dalpha)
```

camroll(dalpha) rotates the up vector by angle dalpha around the vector from view position to target. dalpha is given in radians. A positive value makes the scene rotate counterclockwise.

#### See also

camdolly, camorbit, campan, campos, camproj, camtarget, camup, camva.camzoom

# camtarget

Target position.

## **Syntax**

```
camtarget(p)
camtarget auto
camtarget manual
p = camtarget
```

## **Description**

camtarget(p) sets the target to p. p is a 3D vector.

camtarget auto sets the target to automatic mode, so that it follows the center of the objects which are drawn. camtarget manual sets the target to manual mode.

With an output argument, camtarget gives the current target.

#### See also

camdolly, camorbit, campan, campos, camproj, camroll, camup, camva, camzoom

# camup

Up vector.

```
camup(p)
camup auto
camup manual
p = camup
```

camup(p) sets the up vector to p. p is a 3D vector.

camup auto sets the up vector to [0,0,1]. camup manual does nothing.

With an output argument, camup gives the current up vector.

#### See also

camdolly, camorbit, campan, campos, camproj, camroll, camtarget,
camva, camzoom

#### camva

View angle.

## **Syntax**

```
camva(va)
va = camva
```

# **Description**

camva(va) sets the view angle to va, which is expressed in degrees. The projection mode is set to 'perspective'. The scale is adjusted so that the graphics have about the same size.

With an output argument, camva gives the view angle in degrees, which is 0 for an orthographic projection.

#### See also

camdolly, camorbit, campan, campos, camproj, camroll, camtarget,
camup, camzoom

#### camzoom

Zoom in or out.

```
camzoom(f)
```

camzoom(f) scales the projection by a factor f. The image grows if f is larger than one, and shrinks if it is smaller.

#### See also

camdolly, camorbit, campan, campos, camproj, camroll, camtarget, camup, camva

## contour3

Level curves in 3D space.

## **Syntax**

```
contour3(z)
contour3(z, [xmin, xmax, ymin, ymax])
contour3(z, [xmin, xmax, ymin, ymax], levels)
contour3(z, [xmin, xmax, ymin, ymax], levels, style)
```

## **Description**

contour3(z) plots in 3D space seven contour lines corresponding to the surface whose samples at equidistant points 1:size(z,2) in the x direction and 1:size(z,1) on the y direction are given by z. Contour lines are at equidistant levels. With a second non-empty argument [xmin, xmax, ymin, ymax], the samples are at equidistant points between xmin and xmax in the x direction and between ymin and ymax in the y direction.

The optional third argument levels, if non-empty, gives the number of contour lines if it is a scalar or the levels themselves if it is a vector.

The optional fourth argument is the style of each line, from the minimum to the maximum level (styles are recycled if necessary). The default style is 'kbrmgcy'.

#### See also

contour, mesh, surf

# daspect

Scale ratios along x, y and z axis.

## **Syntax**

```
daspect([rx,ry,rz])
daspect([])
R = daspect
```

## **Description**

daspect(R) specifies the scale ratios along x, y and z axis. Argument R is a vector of 3 elements rx, ry and rz. Coordinates in the 3D space are divided by rx along the x axis, and so on, before the projection is performed. For example, a box whose size is [2;5;3] would be displayed as a cube with daspect([2;5;3]).

daspect([]) sets the scale ratios so that the bounding box of 3D elements is displayed as a cube.

With an output argument, R=daspect gives the current scale ratios as a vector of 3 elements.

#### See also

scale

# lightangle

Set light sources in 3D world.

## **Syntax**

```
lightangle
lightangle(az, el)
```

# **Description**

lightangle(az,el) set lighting source(s) at infinity, with asimuth az and elevation el, both in radians. With missing input argument, the default azimuth is 4 and the default elevation is 1. If az and el are vectors, they must have the same size (except if one of them is a scalar, then it is replicated as needed); lightangle sets multiple light sources.

#### See also

material

#### line3

Plot straight lines in 3D space.

#### **Syntax**

```
line3(A, b)
line3(V, P0)
line3(A, b, style)
line3(A, b, style, id)
```

## **Description**

line3 displays one or several straight line(s) in the 3D space. Each line is defined by two implicit equations or one explicit, parametric equation.

**Implicit equation:** Lines are defined by two equations of the form  $a_1x + a_2y + a_3z = b$ . The first argument of line3 is a matrix which contains the coefficients  $a_1$  in the first column,  $a_2$  in the second column, and  $a_3$  in the third column; two rows define a different line. The second argument is a column vector which contains the coefficients b. If one of these arguments has two rows and the other has several pairs, the same rows are reused multiple times.

**Explicit equations:** Lines are defined by equations of the form  $P = P_0 + \lambda V$  where  $P_0$  is a point of the line, V a vector which defines its direction, and  $\lambda$  a real parameter. The first argument of line3 is a matrix which contains the coefficients  $v_x$  in the first column,  $v_y$  in the second column and  $v_z$  in the third column. The second argument is a matrix which contains the coefficients  $x_0$  in the first column,  $y_0$  in the second column and  $z_0$  in the third column.

The optional third and fourth arguments are the same as for all graphical commands.

## **Example**

Implicit or parametric forms of a vertical line at x=5, y=6:

```
line3([1,0,0;0,1,0], [5;6])
line3([0, 0, 1], [5, 6, 0])
```

#### See also

plot3, line

#### material

Surface reflexion properties.

## **Syntax**

```
material(p)
```

## **Description**

material(p) sets the reflexion properties of the Blinn-Phong model of following surfaces drawn with surf and plotpoly. Argument p is a scalar or a vector of two real values between 0 and 1. The first or only element, ka, is the weight of ambiant light; the second element, kd, is the weight of diffuse light reflected from all light sources.

#### See also

lightangle

#### mesh

Plot a mesh in 3D space.

## **Syntax**

```
mesh(x, y, z)
mesh(z)
mesh(x, y, z, color)
mesh(z, color)
mesh(..., kind)
mesh(..., kind, style)
mesh(..., kind, style, id)
```

# **Description**

mesh(x,y,z) plots a mesh defined by 2-D arrays x, y and z. Arguments x and y must have the same size as z or be vectors of size(z,2) and size(z,1) elements, respectively. If x and y are missing, their default values are coordinates from 1 to size(z,2) along x axis and from 1 to size(z,1) along y axis. Color is obtained by mapping the full range of z values to the color map.

mesh(x,y,z,color) maps values of array color to the color map. color must have the same size as z and contain values between 0 and 1, which are mapped to the color map.

mesh(...,kind) specifies which side of the mesh is visible. kind is a string of 1 or 2 characters: 'f' if the front side is visible (the side where increasing y are on the left of increasing x coordinates), and 'b' if the back side is visible. Default '' is equivalent to 'fb'.

mesh(...,style) specifies the line or symbol style of the mesh.
The default '' is to map z or color values to the color map.
mesh(...,id) specifies the ID used for interactivity in Sysquake.

#### **Example**

```
(X, Y) = meshgrid([-2:0.2:2]);
Z = X.*exp(-X.^2-Y.^2);
mesh(X, Y, Z);
```

#### See also

plot3, surf, plotpoly

# plot3

Generic 3D plot.

## **Syntax**

```
plot3(x, y, z)
plot3(x, y, z, style)
plot3(x, y, z, style, id)
```

# **Description**

The command plot3 displays 3D graphical data in the current figure. The data are given as three vectors of coordinates x, y and z. Depending on the style, the points are displayed as individual marks or are linked with lines.

If x, y and z are matrices, each row is considered as a separate line or set of marks; row or column vectors are replicated to match the size of matrix arguments if required.

plot3(...,id) specifies the ID used for interactivity in Sysquake.

## **Example**

Chaotic attractor of the Shimizu-Morioka system:

```
 \begin{array}{lll} (t,x) &=& 0 \\ \text{de} 45(@(t,x) \ [x(2); \ (1-x(3))*x(1)-0.75*x(2); \ x(1)^2-0.45*x(3)], \\ [0,300], \ [1;1;1]); \\ \text{plot} 3(x(:,1)', \ x(:,2)', \ x(:,3)', \ 'r'); \\ \text{label } x \ y \ z; \\ \text{campos}([-1.5; \ -1.4; \ 3.1]); \end{array}
```

#### See also

line3, plotpoly, plot

# plotpoly

Plot polygons in 3D space.

## **Syntax**

```
plotpoly(x, y, z, ind)
plotpoly(x, y, z, 'strip')
plotpoly(x, y, z, 'fan')
plotpoly(x, y, z, color, ind)
plotpoly(x, y, z, color, 'strip')
plotpoly(x, y, z, color, 'fan')
plotpoly(..., vis)
plotpoly(..., vis, style)
plotpoly(..., vis, style, id)
```

## **Description**

plotpoly(x,y,z,ind) plots polygons whose vertices are given by vectors x, y and z. Rows of argument ind contain the indices of each polygon in arrays x, y, and z. Vertices can be shared by several polygons. Color of each polygon is mapped linearly from the z coordinate of the center of gravity of its vertices to the color map. Each polygon can be concave, but must be planar and must not self-intersect (different polygons may intersect).

plotpoly(x,y,z,'strip') plots a strip of triangles. Triangles are made of three consecutive vertices; their indices could be defined by the following array  $ind_strip$ :

```
ind_strip = ...
[ 1 2 3
    3 2 4
    3 4 5
    5 4 6
    5 6 7
    etc. ];
```

Ordering is such that triangles on the same side of the strip have the same orientation.

plotpoly(x,y,z,'fan') plots triangles which share the first vertex and form a fan. Their indices could be defined by the following array  $ind_fan$ :

```
ind_fan = ...
[ 1 2 3
    1 3 4
    1 4 5
    etc. ];
```

plotpoly(x,y,z,color,...) uses color instead of z to set the filling color of each polygon. color is always a real double array (or scalar) whose elements are between 0 and 1. How it is interpreted depends on its size:

- A scalar defines the color of all polygons; it is mapped to the color map.
- A vector of three elements defines the RGB color of all polygons (row vector if there are 3 vertices to avoid ambiguity).
- A vector with as many elements as x, y and z defines the color of each vertex (column vector if there are 3 vertices to avoid ambiguity). Polygons have the mean value of all their vertices, which is mapped to the color map.
- An array with as many columns as elements in x, y and z defines the RGB color of each vertex. Polygons have the mean value of all their vertices.

plotpoly(..., vis) uses string vis to specify which side of the surface is visible: 'f' for front only, 'b' for back only, or 'fb' or 'bf' for both sides. The front side is defined as the one where vertices have an anticlockwise orientation. The default is 'f'.

plotpoly(...,vis,style) uses string style to specify the style of edges.

plotpoly(...,id) specifies the ID used for interactivity in Sysquake.

#### See also

plot3, surf

#### sensor3

Make graphical element sensitivive to 3D interactive displacement.

```
sensor3(type, param, id)
sensor3(type, param, typeAlt, paramAlt, id)
```

sensor3(type,param,id) specifies how a 3D element can be dragged interactively. Contrary to 2D graphics where the mapping between the mouse cursor and the graphical coordinates depends on two separate scaling factors, manipulation in 3D space must use a surface as an additional constraint. sensor3 specifies this surface for a graphical object whose ID is the same as argument id.

The constraint surface is specified with string type and numeric array param. It always contains the selected point. For instance, if the user clicks the second point of plot3([1,2],[5,3],[2,4],'',1) and sensor3 defines a horizontal plane, the move lies in horizontal plane z=4. In addition to position  $_p1$ , parameters specific to the constraint surface are provided in special variable  $_q$ , a vector of two elements.

type = 'plane' The constraint surface is the plane defined by
the selected point \_p0 and two vectors [vx1;vy1;vz1] and
[vx2;vy2;vz2] given in argument param = [vx1,vy1,vz1;
vx2,vy2,vz2]. During the drag, \_q contains the coefficients of
these two vectors, such that \_p1 = \_p0+\_q'\*param'.

type = 'sphere' The constraint surface is a sphere whose center is defined by a point param = [px,py,pz]. Its R is such that the surface contains the selected point \_p0. During the drag, \_q contains the spherical coordinates phi and theta, such that \_p1 = param' + R \*  $[\cos(q_{-}(1))*\cos(q_{-}(2)); \sin(q_{-}(2))]$ .

With five input arguments, sensor3(type,param,typeAlt,paramAlt,id) specifies an alternative constraint surface used when the modifier key is held down.

# **Examples**

(simple XY plane...)

(phi/theta without modifier, R with modifier with plane and ignored 2nd param)

#### See also

plot3, mesh, plotpoly, surf

#### surf

Plot a surface defined by a grid in 3D space.

#### Syntax

```
surf(x, y, z)
surf(z)
surf(x, y, z, color)
surf(z, color)
surf(..., vis)
surf(..., vis, style)
surf(..., vis, style, id)
```

## **Description**

surf(x,y,z) plots a surface defined by 2-D arrays x, y and z. Arguments x and y must have the same size as z or be vectors of size(z,2) and size(z,1) elements, respectively. If x and y are missing, their default values are coordinates from 1 to size(z,2) along x axis and from 1 to size(z,1) along y axis. Color of each surface cell is obtained by mapping the average z values to the color map.

surf(x,y,z,color) maps values of array color to the color map. color must have the same size as z and contain values between 0 and 1.

surf(...,vis) specifies which side of the surface is visible. vis is a string of 1 or 2 characters: 'f' if the front side is visible (the side where increasing y are on the left of increasing x coordinates), and 'b' if the back side is visible. Default '' is equivalent to 'fb'.

surf(...,style) specifies the line or symbol style of the mesh between surface cells, or the fill style of the surface. The default '' is to map z or color values to the color map for the surface cells and not to draw cell bounds.

mesh(...,id) specifies the ID used for interactivity in Sysquake.

#### **Example**

```
(X, Y) = meshgrid([-2:0.2:2]);
Z = X.*exp(-X.^2-Y.^2);
surf(X, Y, Z, 'k');
```

#### See also

plot3, mesh, plotpoly

# 6.42 Graphical Functions for Dynamical Systems

Graphical commands described in this section are related to automatic control. They display the time responses and frequency responses of linear time-invariant systems defined by transfer functions or state-space models in continuous time (Laplace transform) or discrete time (z transform).

Some of these functions can return results in output arguments instead of displaying them. These values depend not only on the input arguments, but also on the current scale of the figure. For instance, the set of frequencies where the response of the system is evaluated for the Nyquist diagram is optimized in the visible area. Option Range of responseset can be used when this behavior is not suitable, such as for phase portraits using lsim. Output can be used for uncommon display purposes such as special styles, labels, or export. Evaluation or simulation functions not related to graphics, like polyval, ode45 or filter, are better suited to other usages.

# bodemag

Magnitude Bode diagram of a continuous-time system.

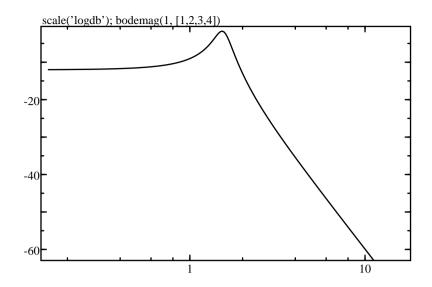
## **Syntax**

```
bodemag(numc, denc)
bodemag(numc, denc, w)
bodemag(numc, denc, opt)
bodemag(numc, denc, w, opt)
bodemag(Ac, Bc, Cc, Dc)
bodemag(Ac, Bc, Cc, Dc, w)
bodemag(Ac, Bc, Cc, Dc, opt)
bodemag(Ac, Bc, Cc, Dc, w, opt)
bodemag(..., style)
bodemag(..., style, id)
(mag, w) = bodemag(...)
```

# **Description**

bodemag(numc,denc) plots the magnitude of the frequency response of the continuous-time transfer function numc/denc. The range of frequencies is selected automatically or can be specified in an optional argument w, a vector of frequencies.

Further options can be provided in a structure opt created with responseset; field Range is utilized. The optional arguments style and id have their usual meaning.



**Figure 6.9** scale('logdb'); bodemag(1, [1,2,3,4])

bodemag(Ac,Bc,Cc,Dc) plots the magnitude of the frequency response  $Y(j\omega)/U(j\omega)$  of the continuous-time state-space model (Ac,Bc,Cc,Dc) defined as

$$j\omega X(j\omega) = A_c X(j\omega) + B_c U(j\omega)$$
  
 $Y(j\omega) = C_c X(j\omega) + D_c U(j\omega)$ 

With output arguments, bodemag gives the magnitude and the frequency as column vectors. No display is produced.

## **Examples**

```
Green plot for |1/(s^3+2s^2+3s+4)| with s=j\omega (see Fig. 6.9): bodemag(1, [1, 2, 3, 4], 'g'); 
The same plot, between \omega=0 and \omega=10: scale([0,10]); bodemag(1, [1, 2, 3, 4], 'g');
```

#### See also

bodephase, dbodemag, sigma, responseset, plotset

# **bodephase**

Phase Bode diagram for a continuous-time system.

## **Syntax**

```
bodephase(numc, denc)
bodephase(numc, denc, w)
bodephase(numc, denc, opt)
bodephase(numc, denc, w, opt)
bodephase(Ac, Bc, Cc, Dc)
bodephase(Ac, Bc, Cc, Dc, w)
bodephase(Ac, Bc, Cc, Dc, opt)
bodephase(Ac, Bc, Cc, Dc, w, opt)
bodephase(..., style)
bodephase(..., style, id)
(phase, w) = bodephase(...)
```

## **Description**

bodephase(numc,denc) plots the phase of the frequency response of the continuous-time transfer function numc/denc. The range of frequencies is selected automatically or can be specified in an optional argument w, a vector of frequencies.

Further options (such as time delay) can be provided in a structure opt created with responseset; fields Delay and Range are utilized. The optional arguments style and id have their usual meaning.

bodephase(Ac,Bc,Cc,Dc) plots the phase of the frequency response  $Y(j\omega)/U(j\omega)$  of the continuous-time state-space model (Ac,Bc,Cc,Dc) defined as

$$j\omega X(j\omega) = A_c X(j\omega) + B_c U(j\omega)$$
  
 $Y(j\omega) = C_c X(j\omega) + D_c U(j\omega)$ 

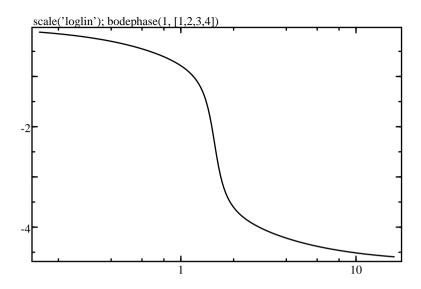
With output arguments, bodephase gives the phase and the frequency as column vectors. No display is produced.

## **Example**

```
Green plot for arg(1/(s^3 + 2s^2 + 3s + 4)), with s = j\omega (see Fig. 6.10): bodephase(1, [1, 2, 3, 4], 'g');
```

#### See also

bodemag, dbodephase, responseset, plotset



**Figure 6.10** scale('loglin'); bodephase(1, [1,2,3,4])

# dbodemag

Magnitude Bode diagram for a discrete-time system.

## **Syntax**

```
dbodemag(numd, dend, Ts)
dbodemag(numd, dend, Ts, w)
dbodemag(numd, dend, Ts, opt)
dbodemag(numd, dend, Ts, w, opt)
dbodemag(Ad, Bd, Cd, Dd, Ts)
dbodemag(Ad, Bd, Cd, Dd, Ts, w)
dbodemag(Ad, Bd, Cd, Dd, Ts, opt)
dbodemag(Ad, Bd, Cd, Dd, Ts, opt)
dbodemag(Ad, Bd, Cd, Dd, Ts, w, opt)
dbodemag(..., style)
dbodemag(..., style, id)
(mag, w) = dbodemag(...)
```

# **Description**

dbodemag(numd,dend,Ts) plots the magnitude of the frequency response of the discrete-time transfer function numd/dend with sampling period Ts. The range of frequencies is selected automatically or can be specified in an optional argument w, a vector of frequencies.

Further options can be provided in a structure opt created with responseset; field Range is utilized. The optional arguments style

and id have their usual meaning.

dbodemag(Ad,Bd,Cd,Dd,Ts) plots the magnitude of the frequency response  $Y(j\omega)/U(j\omega)$  of the discrete-time state-space model (Ad,Bd,Cd,Dd) defined as

$$zX(z) = A_dX(z) + B_dU(z)$$
  
$$Y(z) = C_dX(z) + D_dU(z)$$

where  $z = e^{j\omega T_s}$ .

With output arguments, dbodemag gives the magnitude and the frequency as column vectors. No display is produced.

## **Example**

```
dbodemag(1,poly([0.9,0.7+0.6j,0.7-0.6j]),1);
```

#### See also

bodemag, dbodephase, dsigma, responseset, plotset

# dbodephase

Phase Bode diagram for a discrete-time system.

# **Syntax**

```
dbodephase(numd, dend, Ts)
dbodephase(numd, dend, Ts, w)
dbodephase(numd, dend, Ts, opt)
dbodephase(numd, dend, Ts, w, opt)
dbodephase(Ad, Bd, Cd, Dd, Ts)
dbodephase(Ad, Bd, Cd, Dd, Ts, w)
dbodephase(Ad, Bd, Cd, Dd, Ts, opt)
dbodephase(Ad, Bd, Cd, Dd, Ts, opt)
dbodephase(Ad, Bd, Cd, Dd, Ts, w, opt)
dbodephase(..., style)
dbodephase(..., style, id)
(phase, w) = dbodephase(...)
```

# **Description**

dbodemag(numd,dend,Ts) plots the phase of the frequency response of the discrete-time transfer function numd/dend with sampling period Ts. The range of frequencies is selected automatically or can be specified in an optional argument w, a vector of frequencies.

Further options can be provided in a structure opt created with responseset; field Range is utilized. The optional arguments style and id have their usual meaning.

dbodephase(Ad,Bd,Cd,Dd,Ts) plots the phase of the frequency response  $Y(j\omega)/U(j\omega)$  of the discrete-time state-space model (Ad,Bd,Cd,Dd) defined as

$$zX(z) = A_dX(z) + B_dU(z)$$
  
$$Y(z) = C_dX(z) + D_dU(z)$$

where  $z = e^{j\omega T_s}$ .

With output arguments, dbodephase gives the phase and the frequency as column vectors. No display is produced.

#### **Example**

```
dbodephase(1,poly([0.9,0.7+0.6j,0.7-0.6j]),1);
```

#### See also

bodephase, dbodemag, responseset, plotset

# dimpulse

Impulse response plot of a discrete-time linear system.

## **Syntax**

```
dimpulse(numd, dend, Ts)
dimpulse(numd, dend, Ts, opt)
dimpulse(Ad, Bd, Cd, Dd, Ts)
dimpulse(Ad, Bd, Cd, Dd, Ts, opt)
dimpulse(..., style)
dimpulse(..., style, id)
(y, t) = dimpulse(...)
```

# Description

dimpulse(numd, dend, Ts) plots the impulse response of the discretetime transfer function numd/dend with sampling period Ts.

Further options can be provided in a structure opt created with responseset; field Range is utilized. The optional arguments style and id have their usual meaning.

dimpulse(Ad,Bd,Cd,Dd,Ts) plots the impulse response of the discrete-time state-space model (Ad,Bd,Cd,Dd) defined as

$$x(k+1) = A_dx(k) + B_du(t)$$
  
$$y(k) = C_dx(k) + D_du(k)$$

where u(k) is a unit discrete impulse. The state-space model must have a scalar input, and may have a scalar or vector output.

With output arguments, dimpulse gives the output and the time as column vectors. No display is produced.

## **Example**

```
dimpulse(1, poly([0.9,0.7+0.6j,0.7-0.6j]), 1, 'r');
```

#### See also

impulse, dstep, dlsim, dinitial, responseset, plotset

#### dinitial

Time response plot of a discrete-time linear state-space model with initial conditions.

## **Syntax**

```
dinitial(Ad, Bd, Cd, Dd, Ts, x0)
dinitial(Ad, Cd, Ts, x0)
dinitial(..., opt)
dinitial(..., style)
dinitial(..., style, id)
(y, t) = dinitial(...)
```

# **Description**

dinitial(Ad,Bd,Cd,Dd,Ts,x0) plots the output(s) of the discretetime state-space model (Ad,Bd,Cd,Dd) with null input and initial state x0. The model is defined as

$$x(k+1) = A_dx(k) + B_du(t)$$
  
$$y(k) = C_dx(k) + D_du(k)$$

where u(k) is null. Sampling period is Ts. The state-space model may have a scalar or vector output.

Since there is no system input, matrices Bd and Dd are not used. They can be omitted.

The simulation time range can be provided in a structure opt created with responseset. It is a vector of two elements, the start time and the end time. Such an explicit time range is required when the response is not displayed in a plot where the x axis represents the time.

The optional arguments style and id have their usual meaning.

With output arguments, dinitial gives the output and the time as column vectors. No display is produced.

#### See also

initial, dimpulse, responseset, plotset

#### dlsim

Time response plot of a discrete-time linear system with arbitrary input.

## **Syntax**

```
dlsim(numd, dend, u, Ts)
dlsim(Ad, Bd, Cd, Dd, u, Ts)
dlsim(Ad, Bd, Cd, Dd, u, Ts, x0)
dlsim(..., opt)
dlsim(..., style)
dlsim(..., style, id)
dlsim(..., opt, style)
dlsim(..., opt, style, id)
(y, t) = dlsim(...)
```

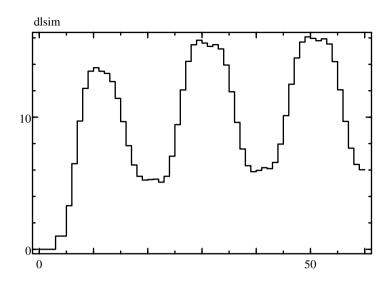
# Description

dlsim(numd,dend,u,Ts) plots the time response of the discrete-time transfer function numd/dend with sampling period Ts. The input is given in real vector u, where the element i corresponds to time (i-1)\*Ts. Input samples before 0 and after length(u)-1 are 0.

dlsim(Ad,Bd,Cd,Dd,u,Ts) plots the time response of the discretetime state-space model (Ad,Bd,Cd,Dd) defined as

$$x(k+1) = A_dx(k) + B_du(t)$$
  
$$y(k) = C_dx(k) + D_du(k)$$

where the system input at time sample k is u(k,:). For single-input systems, u can also be a row vector.



**Figure 6.11** dlsim(1, poly([0.9,0.7+0.6j,0.7-0.6j]), u)

dlsim(Ad,Bd,Cd,Dd,u,Ts,x0) starts with initial state x0 at time t=0. The length of x0 must match the number of states. The default initial state is the zero vector.

The simulation time range can be provided in a structure opt created with responseset. It is a vector of two elements, the start time and the end time. Such an explicit time range is required when the response is not displayed in a plot where the x axis represents the time.

The optional arguments style and id have their usual meaning.

With output arguments, dlsim gives the output and the time as column vectors (or an array for the output of a multiple-output state-space model, where each row represents a sample). No display is produced.

#### **Example**

Simulation of a third-order system with a rectangular input (see Fig. 6.11):

```
u = repmat([ones(1,10), zeros(1,10)], 1, 3);

dlsim(1, poly([0.9,0.7+0.6j,0.7-0.6j]), u, 1, 'rs');
```

#### See also

dstep, dimpulse, dinitial, lsim, responseset, plotset

#### dnichols

Nichols diagram of a discrete-time system.

## **Syntax**

```
dnichols(numd, dend)
dnichols(numd, dend, w)
dnichols(numd, dend, opt)
dnichols(numd, dend, w, opt)
dnichols(..., style)
dnichols(..., style, id)
w = dnichols(...)
(mag, phase) = dnichols(...)
(mag, phase, w) = dnichols(...)
```

## **Description**

dnichols(numd,dend) displays the Nichols diagram of the discrete-time transfer function given by polynomials numd and dend. In discrete time, the Nichols diagram is the locus of the complex values of the transfer function evaluated at  $e^{j\omega}$ , where  $\omega$  is a real number between 0 and  $\pi$  inclusive, displayed in the phase-magnitude plane. Usually, the magnitude is displayed with a logarithmic or dB scale; use scale('lindb') or scale('linlog/lindb') before dnichols.

The range of frequencies is selected automatically between 0 and  $\pi$  or can be specified in an optional argument w, a vector of normalized frequencies.

Further options can be provided in a structure opt created with responseset; fields NegFreq and Range are utilized. The optional arguments style and id have their usual meaning.

With output arguments, dnichols gives the magnitude and phase of the frequency response and the frequency as column vectors. No display is produced.

In Sysquake, when the mouse is over a Nichols diagram, in addition to the magnitude and phase which can be retrieved with  $_{y0}$  and  $_{x0}$ , the normalized frequency is obtained in  $_{q}$ .

## **Example**

```
scale('lindb');
ngrid;
dnichols(3, poly([0.9,0.7+0.6j,0.7-0.6j]))
```

#### See also

nichols, ngrid, dnyquist, responseset, plotset

## dnyquist

Nyquist diagram of a discrete-time system.

## **Syntax**

```
dnyquist(numd, dend)
dnyquist(numd, dend, w)
dnyquist(numd, dend, opt)
dnyquist(numd, dend, w, opt)
dnyquist(..., style)
dnyquist(..., style, id)
w = dnyquist(...)
(re, im) = dnyquist(...)
(re, im, w) = dnyquist(...)
```

## **Description**

The Nyquist diagram of the discrete-time transfer function given by polynomials numd and dend is displayed in the complex plane. In discrete time, the Nyquist diagram is the locus of the complex values of the transfer function evaluated at  $e^{j\omega}$ , where  $\omega$  is a real number between 0 and  $\pi$  inclusive (other definitions include the range between  $\pi$  and  $2\pi$ , which gives a symmetric diagram with respect to the real axis).

The range of frequencies is selected automatically between 0 and  $\pi$  or can be specified in an optional argument w, a vector of normalized frequencies.

Further options can be provided in a structure opt created with responseset; fields NegFreq and Range are utilized. The optional arguments style and id have their usual meaning.

With output arguments, dnichols gives the real and imaginary parts of the frequency response and the frequency as column vectors. No display is produced.

In Sysquake, when the mouse is over a Nyquist diagram, in addition to the complex value which can be retrieved with z0 or x0 and y0, the normalized frequency is obtained in q.

## **Example**

Nyquist diagram with the same scale along both x and y axis and a Hall chart grid (reduced to a horizontal line) (see Fig. 6.12)

```
scale equal;
hgrid;
dnyquist(3, poly([0.9,0.7+0.6j,0.7-0.6j]))
```

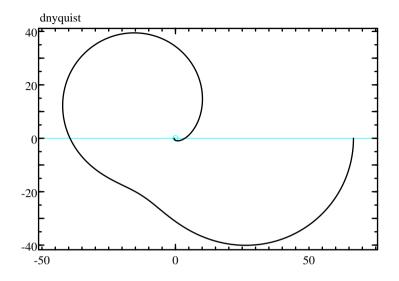


Figure 6.12 dnyquist(3, poly([0.9,0.7+0.6j,0.7-0.6j]))

nyquist, hgrid, dnichols, responseset, plotset

# dsigma

Singular value plot for a discrete-time state-space model.

## **Syntax**

```
dsigma(Ad, Bd, Cd, Dd, Ts)
dsigma(Ad, Bd, Cd, Dd, Ts, w)
dsigma(Ad, Bd, Cd, Dd, Ts, opt)
dsigma(Ad, Bd, Cd, Dd, Ts, w, opt)
dsigma(..., style)
dsigma(..., style, id)
(sv, w) = dsigma(...)
```

# Description

dsigma(Ad,Bd,Cd,Dd,Ts) plots the singular values of the frequency response of the discrete-time state-space model (Ad,Bd,Cd,Dd) defined as

$$zX(z) = A_dX(z) + B_dU(z)$$

$$Y(z) = C_d X(z) + D_d U(z)$$

where  $z = e^{j\omega T_s}$  and Ts is the sampling period.

Further options can be provided in a structure opt created with responseset; field Range is utilized. The optional arguments style and id have their usual meaning.

dsigma is the equivalent of dbodemag for multiple-input systems. For single-input systems, it produces the same plot.

The range of frequencies is selected automatically or can be specified in an optional argument w. a vector of frequencies.

With output arguments, dsigma gives the singular values and the frequency as column vectors. No display is produced.

#### See also

dbodemag, dbodephase, sigma, responseset, plotset

## dstep

Step response plot of a discrete-time linear system.

## **Syntax**

```
dstep(numd, dend, Ts)
dstep(numd, dend, Ts, opt)
dstep(Ad, Bd, Cd, Dd, Ts)
dstep(Ad, Bd, Cd, Dd, Ts, opt)
dstep(..., style)
dstep(..., style, id)
(y, t) = dstep(...)
```

# **Description**

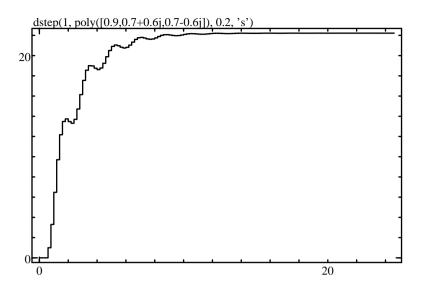
dstep(numd,dend,Ts) plots the step response of the discrete-time transfer function numd/dend with sampling period Ts.

Further options can be provided in a structure opt created with responseset; field Range is utilized. The optional arguments style and id have their usual meaning.

dstep(Ad,Bd,Cd,Dd,Ts) plots the step response of the discrete-time state-space model (Ad,Bd,Cd,Dd) defined as

$$x(k+1) = A_dx(k) + B_du(t)$$
  
$$y(k) = C_dx(k) + D_du(k)$$

where u(k) is a unit step. The state-space model must have a scalar input, and may have a scalar or vector output.



**Figure 6.13** dstep(1,poly([.9,.7+.6j,.7-.6j]),0.2,'s')

With output arguments, dstep gives the output and the time as column vectors. No display is produced.

#### **Examples**

Step response of a discrete-time third-order system (see Fig. 6.13): dstep(1, poly([0.9,0.7+0.6j,0.7-0.6j]), 1, 'g');

Step response of a state-space model with two outputs, and a style argument which is a struct array of two elements to specify two different styles:

```
A = [-0.3,0.1;-0.8,-0.4];
B = [2;3];
C = [1,3;2,1];
D = [2;1];
style = {Color='navy',LineWidth=3; Color='red',LineStyle='-'};
step(A, B, C, D, style);
```

#### See also

dimpulse, dlsim, step, hstep, responseset, plotset

#### erlocus

Root locus of a polynomial with coefficients bounded by an ellipsoid.

## **Syntax**

```
erlocus(CO, P)
erlocus(CO, P, sizes, colors)
```

## **Description**

erlocus displays the set of the roots of all the polynomial whose coefficients are bounded by an ellipsoid defined by C0 and P. The polynomials are defined as C0 + [0,dC], where dC\*inv(P)\*dC' < 1.

If sizes and colors are provided, sizes must be a vector of n values and colors an n-by-3 matrix whose columns correspond respectively to the red, green, and blue components. The locus corresponding to  $dC*inv(P)*dC' < sizes(i)^2$  is displayed with colors(i,:). The vector sizes must be sorted from the smallest to the largest ellipsoid. The default values are sizes = [0.1;0.5;1;2] and colors = [0,0,0;0,0,1;0.4,0.4,1;0.8,0.8,0.8] (i.e. black, dark blue, light blue, and light gray).

**Warning:** depending on the size of the figure (in pixels) and the speed of the computer, the computation may be slow (several seconds). The number of sizes does not have a big impact.

# **Example**

Roots of the polynomial (z-0.8)(z-0.7-0.6j)(z-0.7+0.6j), where the coefficients, in  $R^3$ , have an uncertainty bounded by a unit sphere (see Fig. 6.14).

```
scale('equal', [-2,2,-2,2]);
erlocus(poly([0.8, 0.7+0.6j, 0.7-0.6j]), eye(3));
zgrid;
```

#### See also

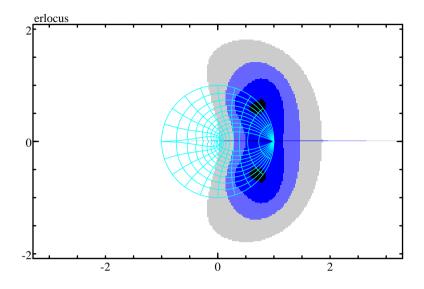
plotroots, rlocus

# hgrid

Hall chart grid.

# Syntax

```
hgrid
hgrid(style)
```



**Figure 6.14** erlocus(poly([.8,.7+.6j,.7-.6j]), eye(3))

## **Description**

hgrid plots a Hall chart in the complex plane of the Nyquist diagram. The Hall chart represents circles which correspond to the same magnitude or phase of the closed-loop frequency response. The optional argument specifies the style.

The whole grid is displayed only if the user selects it in the Grid menu, or after the command plotoption fullgrid. By default, only the unit circle and the real axis are displayed. The whole grid is made of the circles corresponding to a closed-loop magnitude of plus or minus 0, 2, 4, 6, 10, and 20 dB; and to a closed-loop phase of plus or minus 0, 10, 20, 30, 45, 60, and 75 degrees.

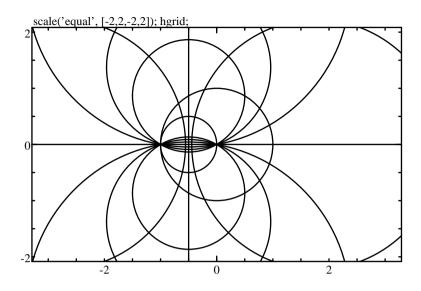
## **Example**

Hall chart grid with a Nyquist diagram (see Fig. 6.15):

```
scale('equal', [-1.5, 1.5, -1.5, 1.5]);
hgrid;
nyquist(20, poly([-1,-2+1j,-2-1j]))
```

#### See also

ngrid, nyquist, plotset, plotoption



**Figure 6.15** Result of hgrid when the whole grid is displayed.

# hstep

Step response plot of a discrete-time transfer function followed by a continuous-time transfer function.

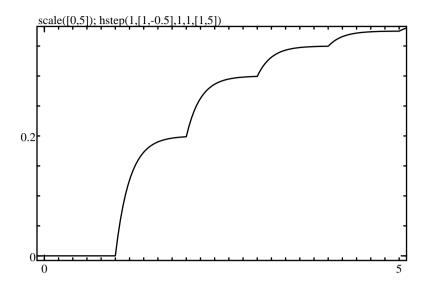
## Syntax

```
hstep(numd, dend, Ts, numc, denc)
hstep(numd, dend, Ts, numc, denc, style)
hstep(numd, dend, Ts, numc, denc, style, id)
```

# Description

A step is filtered first by numd/dend, a discrete-time transfer function with sampling period Ts; the resulting signal is converted to continuous-time with a zero-order hold, and filtered by the continuous-time transfer function numc/denc.

Most discrete-time controllers are used with a zero-order hold and a continuous-time system. hstep can display the simulated output of the system when a step is applied somewhere in the loop, e.g. as a reference signal or a disturbance. The transfer function numd/dend should correspond to the transfer function between the step and the system input; the transfer function numc/denc should be the model of the system.



**Figure 6.16** scale([0,5]); hstep(1,[1,-0.5],1,1,[1,5])

Note that the simulation is performed in open loop. If an unstable system is stabilized with a discrete-time feedback controller, all closed-loop transfer functions are stable; however, the simulation with hstep, which uses the unstable model of the system, may diverge if it is run over a long enough time period, because of round-off errors. But in most cases, this is not a problem.

## **Example**

Exact simulation of the output of a continuous-time system whose input comes from a zero-order hold converter (see Fig. 6.16):

```
% unstable system continuous-time transfer function
num = 1;
den = [1, -1];
% sampling at Ts = 1 (too slow, only for illustration)
Ts = 1;
[numd, dend] = c2dm(num, den, Ts);
% stabilizing proportional controller
kp = 1.5;
% transfer function between ref. signal and input
b = conv(kp, dend);
a = addpol(conv(kp, numd), dend);
% continuous-time output for a ref. signal step
scale([0,10]);
hstep(b, a, Ts, num, den);
```

```
% discrete-time output (exact) dstep(conv(b, numd), conv(a, dend), Ts, 'o');
```

step, dstep, plotset

# impulse

Impulse response plot of a continuous-time linear system.

## **Syntax**

```
impulse(numc, denc)
impulse(numc, denc, opt)
impulse(Ac, Bc, Cc, Dc)
impulse(Ac, Bc, Cc, Dc, opt)
impulse(..., style)
impulse(..., style, id)
(y, t) = impulse(...)
```

## **Description**

impulse(numc,denc) plots the impulse response of the continuoustime transfer function numc/denc.

Further options can be provided in a structure opt created with responseset; fields Delay and Range are utilized. The optional arguments style and id have their usual meaning.

impulse(Ac,Bc,Cc,Dc) plots the impulse response of the continuous-time state-space model (Ac,Bc,Cc,Dc) defined as

$$\frac{dx}{dt}(t) = A_c x(t) + B_c u(t)$$

$$y(t) = C_c x(t) + D_c u(t)$$

where u is a Dirac impulse. The state-space model must have a scalar input, and may have a scalar or vector output.

With output arguments, impulse gives the output and the time as column vectors. No display is produced.

## **Example**

```
impulse(1, 1:4, 'm');
```

dimpulse, step, lsim, initial, responseset, plotset

#### initial

Time response plot for a continuous-time state-space model with initial conditions.

## **Syntax**

```
initial(Ac, Bc, Cc, Dc, x0)
initial(Ac, Cc, x0)
initial(Ac, Bc, Cc, Dc, x0, opt)
initial(..., style)
initial(..., style, id)
(y, t) = initial(...)
```

## **Description**

initial(Ac,Bc,Cc,Dc,x0) plots the output(s) of the continuous-time state-space model (Ac,Bc,Cc,Dc) with null input and initial state x0. The model is defined as

$$\frac{dx}{dt}(t) = A_c x(t) + B_c u(t)$$

$$y(t) = C_c x(t) + D_c u(t)$$

where u(t) is null. The state-space model may have a scalar or vector output.

Since there is no system input, matrices Bd and Dd are not used. They can be omitted.

The simulation time range can be provided in a structure opt created with responseset. It is a vector of two elements, the start time and the end time. Such an explicit time range is required when the response is not displayed in a plot where the x axis represents the time.

The optional arguments style and id have their usual meaning.

With output arguments, initial gives the output and the time as column vectors. No display is produced.

## **Example**

Response of a continuous-time system whose initial state is [5;3] (see Fig. 6.17):

```
initial([-0.3,0.1;-0.8,-0.4],[2;3],[1,3;2,1],[2;1],[5;3])
```

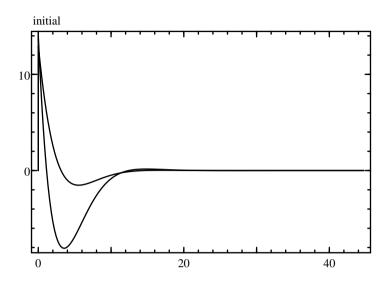


Figure 6.17 Example of initial

dinitial, impulse, responseset, plotset

#### Isim

Time response plot of a continuous-time linear system with piece-wise linear input.

# **Syntax**

```
lsim(numc, denc, u, t)
lsim(numc, denc, u, t, opt)
lsim(Ac, Bc, Cc, Dc, u, t)
lsim(Ac, Bc, Cc, Dc, u, t, opt)
lsim(Ac, Bc, Cc, Dc, u, t, x0)
lsim(Ac, Bc, Cc, Dc, u, t, x0, opt)
lsim(..., style)
lsim(..., style, id)
(y, t) = lsim(...)
```

# **Description**

lsim(numc,denc,u,t) plots the time response of the continuous-time transfer function numd/dend. The input is piece-wise linear; it is de-

fined by points in real vectors t and u, which must have the same length. Input before t(1) and after t(end) is 0. The input used for the simulation is interpolated to have a smooth response.

lsim(Ac,Bc,Cc,Dc,u,t) plots the time response of the continuous-time state-space model (Ac,Bc,Cc,Dc) defined as

$$\frac{dx}{dt}(t) = A_c x(t) + B_c u(t)$$

$$y(t) = C_c x(t) + D_c u(t)$$

where the system input at time sample t(i) is u(i,:)'. For single-input systems, u can also be a row vector.

lsim(Ac,Bc,Cc,Dc,u,t,x0) starts with initial state x0 at time t=0. The length of x0 must match the number of states. The default initial state is the zero vector.

Options can be provided in a structure opt created with responseset:

- 'Range' The range is a vector of two elements, the start time and the end time. Such an explicit time range is required when the response is not displayed in a plot where the x axis represents the time.
- 'tOnly' When opt.tOnly is true, lsim produces output only at the time instants defined in t. The logical value false gives the default interpolated values.

The optional arguments style and id have their usual meaning.

With output arguments, lsim gives the output and the time as column vectors (or an array for the output of a multiple-output statespace model, where each row represents a sample). No display is produced.

#### **Example**

Response of continuous-time system given by its transfer function with an input defined by linear segments (see Fig. 6.18):

```
t = [0, 10, 20, 30, 50];
u = [1, 1, 0, 1, 1];
lsim(1, [1, 2, 3, 4], u, t, 'b');
```

#### See also

step, impulse, initial, dlsim, plotset

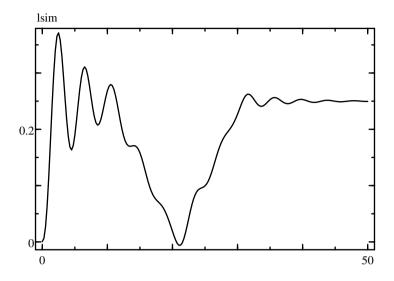


Figure 6.18 lsim(1, [1,2,3,4], u, t)

## ngrid

Nichols chart grid.

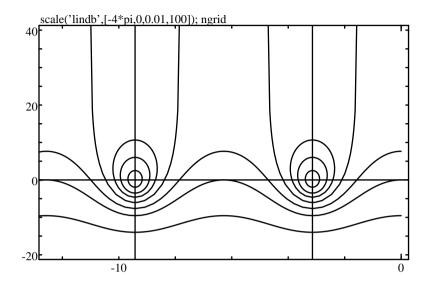
## **Syntax**

ngrid
ngrid(style)

## **Description**

ngrid plots a Nichols chart in the complex plane of the Nichols diagram (see Fig. 6.19). The Nichols chart is a set of lines which correspond to the same magnitude of the closed-loop frequency response. The optional argument specifies the style.

The whole grid is displayed only if the user selects it in the Grid menu, or after the command plotoption fullgrid. By default, only the lines corresponding to unit magnitude and to a phase equal to  $-\pi(1+2k)$ , with integer k, are displayed. The whole grid is made of the lines corresponding to a closed-loop magnitude of -12, -6, -3, 0, 3, 6 and 12 dB.



**Figure 6.19** Result of ngrid in dB when the whole grid is displayed.

## **Example**

```
ngrid;
nichols(7, 1:3);
```

#### See also

hgrid, nichols, plotset, plotoption

## nichols

Nichols diagram of a continuous-time system.

## **Syntax**

```
nichols(numc, denc)
nichols(numc, denc, w)
nichols(numc, denc, opt)
nichols(numc, denc, w, opt)
nichols(..., style)
nichols(..., style, id)
w = nichols(...)
(mag, phase) = nichols(...)
(mag, phase, w) = nichols(...)
```

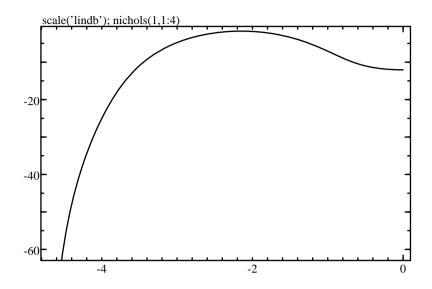


Figure 6.20 scale('lindb'); nichols(1,1:4)

#### **Description**

nichols (numc, denc) displays the Nichols diagram of the continuous-time transfer function given by polynomials numc and denc. In continuous time, the Nichols diagram is the locus of the complex values of the transfer function evaluated at  $j\omega$ , where  $\omega$  is real positive, displayed in the phase-magnitude plane. Usually, the magnitude is displayed with a logarithmic or dB scale; use scale('lindb') or scale('linlog/lindb') before nichols.

The range of frequencies is selected automatically or can be specified in an optional argument w, a vector of frequencies.

Further options can be provided in a structure opt created with responseset; fields Delay, NegFreq and Range are utilized. The optional arguments style and id have their usual meaning.

With output arguments, nichols gives the phase and magnitude of the frequency response and the frequency as column vectors. No display is produced.

In Sysquake, when the mouse is over a Nichols diagram, in addition to the magnitude and phase which can be retrieved with  $_{y0}$  and  $_{x0}$ , the frequency is obtained in  $_{q}$ .

## **Examples**

Nichols diagram of a third-order system (see Fig. 6.20):

```
scale('lindb');
ngrid;
nichols(20,poly([-1,-2+1j,-2-1j]));

Same plot with angles in degrees:

scale('lindb');
scalefactor([180/pi, 1]);
ngrid;
nichols(20,poly([-1,-2+1j,-2-1j]));
```

dnichols, ngrid, nyquist, responseset, plotset, scalefactor

## nyquist

Nyquist diagram of a continuous-time system.

## **Syntax**

```
nyquist(numc, denc)
nyquist(numc, denc, w)
nyquist(numc, denc, opt)
nyquist(numc, denc, w, opt)
nyquist(..., style)
nyquist(..., style, id)
w = nyquist(...)
(re, im) = nyquist(...)
(re, im, w) = nyquist(...)
```

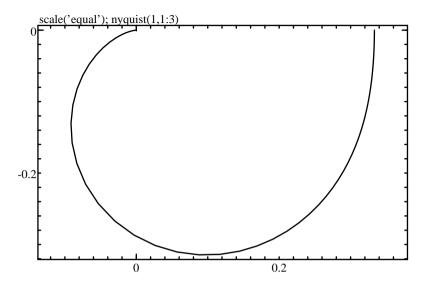
## **Description**

The Nyquist diagram of the continuous-time transfer function given by polynomials numc and denc is displayed in the complex plane. In continuous time, the Nyquist diagram is the locus of the complex values of the transfer function evaluated at  $j\omega$ , where  $\omega$  is real positive (other definitions include the real negative values, which gives a symmetric diagram with respect to the real axis).

The range of frequencies is selected automatically or can be specified in an optional argument w, a vector of frequencies.

Further options can be provided in a structure opt created with responseset; fields Delay, NegFreq and Range are utilized. The optional arguments style and id have their usual meaning.

With output arguments, nyquist gives the real and imaginary parts of the frequency response and the frequency as column vectors. No display is produced.



**Figure 6.21** scale equal; nyquist(1,[1,2,3])

In Sysquake, when the mouse is over a Nyquist diagram, in addition to the complex value which can be retrieved with  $_z0$  or  $_x0$  and  $_y0$ , the frequency is obtained in  $_q$ .

#### **Example**

Nyquist diagram of a third-order system (see Fig. 6.21):

```
scale equal;
hgrid;
nyquist(20, poly([-1,-2+1j,-2-1j]))
```

#### See also

dnyquist, hgrid, nichols, responseset, plotset

# plotroots

Roots plot.

## **Syntax**

```
plotroots(pol)
plotroots(pol, style)
plotroots(pol, style, id)
```

## **Description**

plotroots(pol) displays the roots of the polynomial pol in the complex plane. If this argument is a matrix, each line corresponds to a different polynomial. The default style is crosses; it can be changed with a second argument, or with named arguments.

#### **Example**

```
den = [1, 2, 3, 4];
num = [1, 2];
scale equal;
plotroots(den, 'x');
plotroots(num, 'o');
```

#### See also

rlocus, erlocus, sgrid, zgrid, plotset, movezero

## responseset

Options for frequency responses.

## Syntax

```
options = responseset
options = responseset(name1, value1, ...)
options = responseset(options0, name1, value1, ...)
```

# **Description**

responseset (name1, value1,...) creates the option argument used by functions which display frequency and time responses, such as nyquist and step. Options are specified with name/value pairs, where the name is a string which must match exactly the names in the table below. Case is significant. Options which are not specified have a default value. The result is a structure whose fields correspond to each option. Without any input argument, responseset creates a structure with all the default options. Note that functions such as nyquist and step also interpret the lack of an option argument as a request to use the default values. Contrary to other functions which accept options in structures, such as ode45, empty array [] cannot be used (it would be interpreted incorrectly as a numeric argument).

When its first input argument is a structure, responseset adds or changes fields which correspond to the name/value pairs which follow. Here is the list of permissible options:

Name	Default	Meaning
Delay	0	time delay
NegFreq	false	negative frequencies
Offset	0	offset
Range	[]	time or frequency range
t0nly	false	samples for specified time only (lsim)

Option Delay is used only by continuous-time frequency-response and time-response functions; for frequency responses, it subtracts a phase of delay\*w, where w is the angular frequency. Option Offset adds a a value to the step or impulse response.

Option NegFreq is used in Nyquist and Nichols diagrams, continuous-time or discrete-time; when true, the response is computed for negative frequencies instead of positive frequencies. Option Range should take into account the sampling period for discrete-time commands where it is specified.

## **Examples**

Default options:

```
responseset Delay: 0 NegFreq: false Nyquist diagram of e^{-s}/(s+1): nyquist(1, [1,1], responseset('Delay', 1));
```

Complete Nyquist diagram of  $1/(s^3 + 2s^2 + 2s + 1)$  with dashed line for negative frequencies:

```
nyquist(2, [1,2,2,1]);
nyquist(2, [1,2,2,1], responseset('NegFreq',true), '-');
```

#### See also

bodemag, bodephase, dbodemag, dbodephase, dlsim, dnichols, dnyquist, dsigma, impulse, lsim, nichols, nyquist, sigma, step

#### rlocus

Root locus.

## **Syntax**

```
rlocus(num, den)
rlocus(num, den, style)
rlocus(num, den, style, id)
branches = rlocus(num, den)
```

## **Description**

The root locus is the locus of the roots of the denominator of the closed-loop transfer function (characteristic polynomial) of the system whose open-loop transfer function is num/den when the gain is between 0 and +∞ inclusive. The characteristic polynomial is num +  $k \cdot \text{den.}$  with k > 0. rlocus requires a system with real coefficients. causal or not. Note that the rlocus is defined the same way in the domain of the Laplace transform, the z transform, and the delta transform. The root locus is made of length(den)-1 branches which start from each pole and end to each zero or to a real or complex point at infinity. The locus is symmetric with respect to the real axis, because the coefficients of the characteristic polynomial are real. By definition, closed-loop poles for the current gain (i.e. the roots of num+den) are on the root locus, and move on it when the gain change, rlocus plots only the root locus, not the particular values of the roots for the current gain, a null gain or an infinite gain. If necessary, these values should be plotted with plotroots.

The part of the root locus which is calculated and drawn depends on the scale. If no scale has been set before explicitly with scale or implicitly with plotroots or plot, the default scale is set such that the zeros of num and den are visible.

With an output argument, rlocus gives the list of root locus branches, i.e. a list of row vectors which contain the roots. Different branches do not always have the same numbers of values, because rlocus adapts the gain steps for each branch. Parts of the root locus outside the visible area of the complex plane, as defined by the current scale, have enough points to avoid any interference in the visible area when they are displayed with plot. The gains corresponding to roots are not available directly; they can be computed as real(polyval(den,r)/polyval(num,r)) for root r.

As with other plots, the id is used for interactive manipulation. Manipulating a root locus means changing the gain of the controller, which keeps the locus at the same place but makes the closed-loop poles move on it. Other changes are done by dragging the open-loop poles and zeros, which are plotted by plotroots. To change the gain, you must also plot the current closed-loop poles with the plotroots function and use the same ID, so that the initial click identifies the nearest closed-loop pole and the mouse drag makes Sysquake use the root locus to calculate the change of gain, which can be retrieved in  $_{-}q$  (see the example below).

## **Examples**

Root locus of  $(s^2 + 3s + 2)/(s^3 + 2s^2 + 3s + 4)$  with open-loop poles and zeros added with plotroots (see Fig. 6.22):

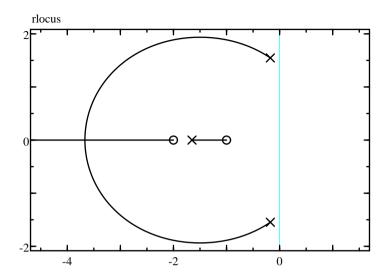


Figure 6.22 Example of rlocus

```
num = [1, 3, 2];
den = [1, 2, 3, 4];
scale('equal', [-4,1,-2,2]);
sgrid;
rlocus(num, den);
plotroots(num, 'o');
plotroots(den, 'x');
```

The second example shows how rlocus can be used interactively in Sysquake.

```
figure "Root Locus"
  draw myPlotRLocus(num, den);
  mousedrag num = myDragRLocus(num, _q);

function
{@
function myPlotRLocus(num, den)
  scale('equal', [-3, 1, -2, 2]);
  rlocus(num, den, '', 1);
  plotroots(addpol(num, den), '^', 1);

function num = myDragRLocus(num, q)
  if isempty(q)
    cancel;
  else
    num = q * num;
```

```
end
@}
```

#### Caveat

The Laguerre algorithm is used for fast evaluation (roots and plotroots are based on eig and have a better accuracy, but their evaluation for a single polynomial is typically 10 times slower). The price to pay is a suboptimal precision for multiple roots and/or high-order polynomials.

#### See also

plotroots, plotset, erlocus, sgrid, zgrid

# sgrid

Relative damping and natural frequency grid for the poles of a continuous-time system.

## **Syntax**

```
sgrid
sgrid(damping, freq)
sgrid(..., style)
```

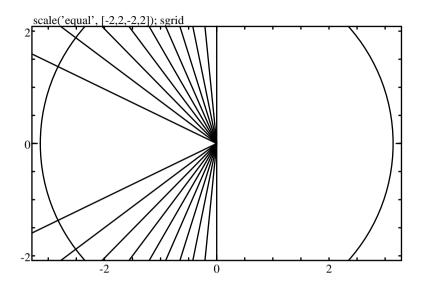
## **Description**

With no numeric argument, sgrid plots a grid of lines with constant relative damping and natural frequencies in the complex plane of s (see Fig. 6.23).

The whole grid is displayed only if the user selects it in the Grid menu, or after the command plotoption fullgrid. By default, only the imaginary axis (the stability limit for the poles of the Laplace transform) is displayed.

With one or two numeric arguments, sgrid plots only the lines for the specified values of damping and natural frequency. Let p and  $\overline{p}$  be the complex conjugate roots of the polynomial  $s^2+2\omega\zeta s+\omega^2$ , where  $\omega$  is the natural frequency and  $\zeta<1$  the damping. The locus of roots with a constant damping  $\zeta$  is generated by  $|\mathrm{Im}\,p|=\sqrt{1-\zeta^2}\mathrm{Re}\,p$  with  $\mathrm{Re}\,p<0$ . The locus of roots with a constant natural frequency  $\omega$  is a circle of radius  $\omega$ .

The style argument has its usual meaning.



**Figure 6.23** Result of sgrid when the whole grid is displayed.

#### **Example**

Typical use for poles or zeros displayed in the s plane:

```
scale equal;
sgrid;
plotroots(pol);
```

#### See also

zgrid, plotroots, hgrid, ngrid, plotset, plotoption

# sigma

Singular value plot for a continuous-time state-space model.

# Syntax

```
sigma(Ac, Bc, Cc, Dc)
sigma(Ac, Bc, Cc, Dc, w)
sigma(Ac, Bc, Cc, Dc, opt)
sigma(Ac, Bc, Cc, Dc, w, opt)
sigma(..., style)
sigma(..., style, id)
(sv, w) = sigma(...)
```

## **Description**

sigma(Ac,Bc,Cc,Dc) plots the singular values of the frequency response of the continuous-time state-space model (Ac,Bc,Cc,Dc) defined as

$$j\omega X(j\omega) = A_c X(j\omega) + B_c U(j\omega)$$
  
 $Y(j\omega) = C_c X(j\omega) + D_c U(j\omega)$ 

The range of frequencies is selected automatically or can be specified in an optional argument w, a vector of frequencies.

Further options can be provided in a structure opt created with responseset; field Range is utilized. The optional arguments style and id have their usual meaning.

sigma is the equivalent of bodemag for multiple-input systems. For single-input systems, it produces the same plot.

With output arguments, sigma gives the singular values and the frequency as column vectors. No display is produced.

#### See also

bodemag, bodephase, dsigma, responseset, plotset

## step

Step response plot of a continuous-time linear system.

## **Syntax**

```
step(numc, denc)
step(numc, denc, opt)
step(Ac, Bc, Cc, Dc)
step(Ac, Bc, Cc, Dc, opt)
step(..., style)
step(..., style, id)
(y, t) = step(...)
```

# Description

step(numc,denc) plots the step response of the continuous-time transfer function numc/denc.

Further options can be provided in a structure opt created with responseset; fields Delay and Range are utilized. The optional arguments style and id have their usual meaning.

step(Ac,Bc,Cc,Dc) plots the step response of the continuous-time state-space model (Ac,Bc,Cc,Dc) defined as

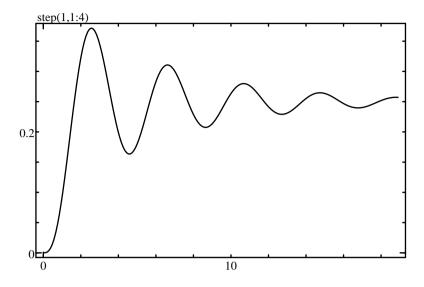


Figure 6.24 step(1, [1,2,3,4])

$$\frac{dx}{dt}(t) = A_c x(t) + B_c u(t)$$

$$y(t) = C_c x(t) + D_c u(t)$$

where u is a unit step. The state-space model must have a scalar input, and may have a scalar or vector output.

With output arguments, step gives the output and the time as column vectors. No display is produced.

## **Example**

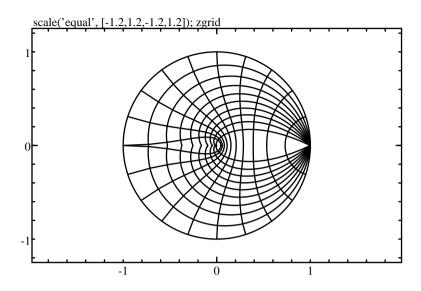
Step response of the continuous-time system  $1/(s^3 + 2s^2 + 3s + 4)$  (see Fig. 6.24):

#### See also

impulse, lsim, dstep, hstep, responseset, plotset

# zgrid

Relative damping and natural frequency grid for the poles of a discrete-time system.



**Figure 6.25** Result of zgrid when the whole grid is displayed.

#### Syntax

```
zgrid
zgrid(damping, freq)
zgrid(..., style)
```

#### **Description**

With no numeric argument, zgrid plots a grid of lines with constant relative damping and natural frequencies in the complex plane of z (see Fig. 6.25).

The whole grid is displayed only if the user selects it in the Grid menu, or after the command plotoption fullgrid. By default, only the unit circle (the stability limit for the poles of the z transform) is displayed.

With one or two numeric arguments, zgrid plots only the lines for the specified values of damping and natural frequency. The damping  $\zeta$  and the natural frequency  $\omega$  are defined the same way as for the sgrid function, with the mapping  $z=e^s$  (a normalized sampling frequency is assumed). With a damping  $\zeta$ , the line z and its complex conjugate  $\overline{z}$  are generated by  $z=e^{(-1+j\sqrt{1-\zeta^2}/\zeta)u}$ , with  $0 \le u \le u_{\text{max}}$  and  $u_{\text{max}}$  chosen such that the line has a positive imaginary part. With a natural frequency  $\omega$  (typically in the range 0 for a null frequency to  $\pi$  for the Nyquist frequency), the line is generated by  $e^{\omega e^{jv}}$ , where v is such that the curve is inside the unit circle.

The style argument has its usual meaning.

### **Example**

Typical use for poles or zeros displayed in the z plane:

```
scale('equal',[-1.2,1.2,-1.2,1.2]);
zgrid;
plotroots(pol);
```

#### See also

sgrid, plotroots, hgrid, ngrid, plotset, plotoption

## 6.43 Sysquake for LaTeX Functions

Functions described in this section are specific to Sysquake for LaTeX. Some of them are documented because they appear in .lme files generated when .tex files which include file sysquake.sty are typeset, but they should not be called directly.

## sqlxendfragment

End fragment of code.

## **Syntax**

```
sqlxendfragment(fragid)
```

## **Description**

sqlxendfragment(fragid) ends a fragment of code beginning with beginlmefragment. Argument fragid (a string) must match the first argument of the previous beginlmefragment.

sqlxendfragment is used by the Sysquake package; it should not be called from .tex files.

#### See also

sqlxstartfragment

## sqlxvalue

Format value for an equation.

```
sqlxvalue(v)
str = sqlxvalue(v)
```

### **Description**

sqlxvalue(v) inserts value v in the LaTeX output, enclosed in \ensuremath so that it can be used in math mode as well as in text mode.

With an output argument, sqlxvalue(v) returns its result in a string.

## sqlxsource

Insert or get source code of LME function.

### **Syntax**

```
sqlxsource fun
source = sqlxsource('fun');
```

## **Description**

sqlxsource inserts the source code of an LME function in the LaTeX document. The function is specified by its name; methods are specified as 'classname::methodname'. The source code is displayed with the typewriter font (\tt).

With an output argument, sqlxsource get the source code as a string and does not insert it in the LaTeX document.

## sqlxstartfragment

Start a new fragment of code.

## **Syntax**

```
sqlxstartfragment(fragid, width, height)
```

## **Description**

sqlxstartfragment(fragid,width,height) starts a new fragment of code which corresponds to command \sqexpr or environment sysquake in LaTeX. Argument fragid is a string made only of letters which is used as fragment identifier; it is typically the fragment index

as a lowercase roman number starting at 1. Arguments width and height are the dimensions of the figure reserved in LaTeX, or negative if the result is text without graphics.

sqlxstartfragment is used by the Sysquake package; it should not be called from .tex files.

#### See also

sqlxendfragment, beginfigure

## **Chapter 7**

# Libraries

Libraries are collections of functions which complement the set of built-in functions and operators of LME, the programming language of Sysquake. To use them, type (or add in the functions block of the SQ files which rely on them) a use command, such as

use stdlib

- bench bench implements a benchmark which can be used to compare the performance of LME on different platforms.
- bitfield bitfield implements constructors and methods for bit fields (binary numbers). Standard operators are redefined to enable the use of & and | for bitwise operations, and subscripts for bit extraction and assignment.
- colormaps colormaps defines functions which create color maps for command colormap.
- constants constants defines physical constants in SI units.
- date date implements functions for date and time manipulation and conversion to and from strings.
- filter filter implements functions for the design of analog and digital filters.
- lti lti implements constructors and methods for Linear Time-Invariant models, which may represent dynamical systems as continuous-time or discrete-time state-space models or transfer functions. With them, you can use standard operator notations such as + or \*, array building operators such as [A,B;C,D], connection functions such as parallel or feedback, and much more.

- lti\_filter lti\_filter implements functions for the design of analog and digital filters given as lti objects.
- lti\_gr lti\_gr, loaded automatically with lti, defines methods
  which provide for lti objects the same functionality as the native
  graphical functions of Sysquake for dynamical systems, such as
  bodemag for the magnitude of the Bode diagram or step for the
  step response.
- polyhedra polyhedra implements functions which create solid shapes with polygonal faces in 3D.
- polynom polynom implements constructors and methods for polynomial and rational functions. With them, you can use standard operator notations such as + or \*.
- probdist probdist defines classes for probability distributions.
- sigenc sigenc implements functions related to signal encoding to and decoding from a digital representation.
- solids solids implements functions which create solid shapes in 3D. Solids are generated with parametric equations and displayed with surf.
- stat stat provides more advanced statistical functions.
- stdlib stdlib is the standard library of general-purpose functions for LME. Functions span from array creation and manipulation to coordinates transform and basic statistics.
- wav wav implements functions for reading and writing WAV files, or encoding and decoding data encoded as WAV in memory.

## 7.1 stdlib

stdlib is a library which extends the native LME functions in the following areas:

- creation of matrices: blkdiag, compan, hankel, toeplitz
- geometry: subspace
- functions on integers: primes
- statistics: corrcoef, perms
- data processing: circshift, cumtrapz, fftshift, filter2, hist, ifftshift, polyfit, polyvalm, trapz

```
- other: isreal, sortrows
```

The following statement makes available functions defined in stdlib:

```
use stdlib
```

## **Functions**

### circshift

Shift the elements of a matrix in a circular way.

## **Syntax**

```
use stdlib
B = circshift(A, shift_vert)
B = circshift(A, [shift_vert, shift_hor])
```

## **Description**

circshift(A,sv) shifts the rows of matrix A downward by sv rows. The sv bottom rows of the input matrix become the sv top rows of the output matrix. sv may be negative to go the other way around.

circshift(A,[sv,sh]) shifts the rows of matrix A downward by sv rows, and its columns to the right by sh columns. The sv bottom rows of the input matrix become the sv top rows of the output matrix, and the sh rightmost columns become the sh leftmost columns.

#### See also

```
rot90, fliplr, flipud
```

## **blkdiag**

Block-diagonal matrix.

## Syntax

```
use stdlib
X = blkdiag(B1, B2, ...)
```

## **Description**

blkdiag(B1,B2,...) creates a block-diagonal matrix with matrix blocks B1, B2, etc. Its input arguments do not need to be square.

### **Example**

```
use stdlib
blkdiag([1,2;3,4], 5)
    1 2 0
    3 4 0
    0 0 5
blkdiag([1,2], [3;4])
    1 2 0
    0 0 3
    0 0 4
```

### See also

diag

### compan

Companion matrix.

## **Syntax**

```
use stdlib
X = compan(pol)
```

## **Description**

compan(pol) gives the companion matrix of polynomial pol, a square matrix whose eigenvalues are the roots of pol.

## **Example**

```
use stdlib
compan([2,3,4,5])
-1.5 -2.0 -2.5
1.0 0.0 0.0
0.0 1.0 0.0
```

### See also

poly, eig

#### corrcoef

Correlation coefficients.

```
use stdlib
S = corrcoef(X)
S = corrcoef(X1, X2)
```

### **Description**

corrcoef(X) calculates the correlation coefficients of the columns of the m-by-n matrix X. The result is a square n-by-n matrix whose diagonal is 1.

corrcoef(X1,X2) calculates the correlation coefficients of X1 and X2 and returns a 2-by-2 matrix. It is equivalent to corrcoef([X1(:),X2(:)]).

## **Example**

```
use stdlib
corrcoef([1, 3; 2, 5; 4, 4; 7, 10])
    1     0.8915
    0.8915    1
corrcoef(1:5, 5:-1:1)
    1 -1
    -1    1
```

#### See also

COV

## cumtrapz

Cumulative numeric integration with trapezoidal approximation.

## **Syntax**

```
use stdlib
S = cumtrapz(Y)
S = cumtrapz(X, Y)
S = cumtrapz(X, Y, dim)
```

## **Description**

cumtrapz(Y) calculates an approximation of the cumulative integral of a function given by the samples in Y with unit intervals. The trapezoidal approximation is used. If Y is neither a row nor a column vector, integration is performed along its columns. The result has the same size as Y. The first value(s) is (are) 0.

cumtrapz(X,Y) specifies the location of the samples. A third argument may be used to specify along which dimension the integration is performed.

### **Example**

```
use stdlib
cumtrapz([2, 3, 5])
    0    2.5    6.5
cumtrapz([1, 2, 5], [2, 3, 5])
    0    2.5    14.5
```

#### See also

cumsum, trapz

### fftshift

Shift DC frequency of FFT from beginning to center of spectrum.

## **Syntax**

```
use stdlib
Y = fftshift(X)
```

## **Description**

fftshift(X) shifts halves of vector (1-d) or matrix (2-d) X to move the DC component to the center. It should be used after fft or fft2.

#### See also

```
fft. ifftshift
```

#### filter2

Digital 2-d filtering of data.

```
use stdlib
Y = filter2(F, X)
Y = filter2(F, X, shape)
```

filter2(F,X) filters matrix X with kernel F with a 2-d correlation. The result has the same size as X.

An optional third argument is passed to conv2 to specify another method to handle the borders.

filter2 and conv2 have three differences: arguments F and X are permuted, filtering is performed with a correlation instead of a convolution (i.e. the kernel is rotated by 180 degrees), and the default method for handling the borders is 'same' instead of 'full'.

#### See also

filter, conv2

#### hankel

Hankel matrix.

### **Syntax**

```
use stdlib
X = hankel(c, r)
```

## **Description**

hankel (c,r) creates a Hankel matrix whose first column contains the elements of vector c and whose last row contains the elements of vector r. A Hankel matrix is a matrix whose antidiagonals have the same value. In case of conflict, the first element of r is ignored. The default value of r is a zero vector the same length as c.

### **Example**

```
use stdlib
hankel(1:3, 3:8)
1 2 3 4 5 6
2 3 4 5 6 7
3 4 5 6 7 8
```

#### See also

toeplitz, diag

#### hist

Histogram.

## **Syntax**

```
use stdlib
(N, X) = hist(Y)
(N, X) = hist(Y, m)
(N, X) = hist(Y, m, dim)
N = hist(Y, X)
N = hist(Y, X, dim)
```

## **Description**

hist(Y) gives the number of elements of vector Y in 10 equallyspaced intervals. A second input argument may be used to specify the number of intervals. The center of the intervals may be obtained in a second output argument.

If Y is an array, histograms are computed along the dimension specified by a third argument or the first non-singleton dimension; the result N has the same size except along that dimension.

When the second argument is a vector, it specifies the centers of the intervals.

#### **Example**

```
use stdlib
(N, X) = hist(logspace(0,1), 5)
N =
    45    21    14    11    9
X =
    1.9    3.7    5.5    7.3    9.1
```

## ifftshift

Shift DC frequency of FFT from center to beginning of spectrum.

```
use stdlib
Y = ifftshift(X)
```

ifftshift(X) shifts halves of vector (1-d) or matrix (2-d) X to move the DC component from the center. It should be used before ifft or ifft2. It reverses the effect of fftshift.

#### See also

```
ifft, fftshift
```

## isreal

Test for a real number.

### **Syntax**

```
use stdlib
b = isreal(x)
```

## **Description**

isreal(x) is true if x is a real scalar or a matrix whose entries are all real.

### **Examples**

```
use stdlib
isreal([2,5])
  true
isreal([2,3+2j])
  false
isreal(exp(pi*1j))
  true
```

#### See also

isnumeric, isfloat, isscalar

## perms

Array of permutations.

```
use stdlib
M = perms(v)
```

perm(v) gives an array whose rows are all the possible permutations of vector v.

#### **Example**

```
use stdlib
perms(1:3)
3 2 1
3 1 2
2 3 1
1 3 2
2 1 3
1 2 3
```

#### See also

sort

## polyfit

Polynomial fit.

## **Syntax**

```
use stdlib
pol = polyfit(x, y, n)
```

## **Description**

polyfit(x,y,n) calculates the polynomial (given as a vector of descending power coefficients) of order n which best fits the points given by vectors x and y. The least-square algorithm is used.

## **Example**

```
use stdlib
pol = polyfit(1:5, [2, 1, 4, 5, 2], 3)
  pol =
    -0.6667   5.5714 -12.7619   9.8000
polyval(pol, 1:5)
    1.9429   1.2286   3.6571   5.2286   1.9429
```

## polyvalm

Value of a polynomial with square matrix argument.

```
use stdlib
Y = polyvalm(pol, X)
```

## **Description**

polyvalm(pol,X) evaluates the polynomial given by the coefficients pol (in descending power order) with a square matrix argument.

## **Example**

```
use stdlib
polyvalm([1,2,8],[2,1;0,1])
   16   5
   0  11
```

#### See also

polyval

## primes

List of primes.

## Syntax

```
use stdlib
v = primes(n)
```

## **Description**

 $\operatorname{primes}(n)$  gives a row vector which contains the primes up to n.

## **Example**

```
use stdlib
primes(20)
2 3 5 7 11 13 17 19
```

#### sortrows

Sort matrix rows.

```
use stdlib
(S, index) = sortrows(M)
(S, index) = sortrows(M, sel)
(S, index) = sortrows(M, sel, dim)
```

### **Description**

sortrows (M) sort the rows of matrix M. The sort order is based on the first column of M, then on the second one for rows with the same value in the first column, and so on.

sortrows (M, sel) use the columns specified in sel for comparing the rows of M. A third argument dim can be used to specify the dimension of the sort: 1 for sorting the rows, or 2 for sorting the columns.

The second output argument of sortrows gives the new order of the rows or columns as a vector of indices.

#### **Example**

```
use stdlib
sortrows([3, 1, 2; 2, 2, 1; 2, 1, 2])
  2  1  2
  2  2  1
  3  1  2
```

#### See also

sort

## subspace

Angle between two subspaces.

## **Syntax**

```
use stdlib
theta = subspace(A, B)
```

## **Description**

subspace(A,B) gives the angle between the two subspaces spanned by the columns of A and B.

### **Examples**

Angle between two vectors in R<sup>2</sup>:

```
use stdlib
a = [3; 2];
b = [1; 5];
subspace(a, b)
0.7854
```

Angle between the vector [1;1;1] and the plane spanned by [2;5;3] and [7;1;0] in R<sup>3</sup>:

```
subspace([1;1;1], [2,7;5,1;3,0])
0.2226
```

## toeplitz

Toeplitz matrix.

## **Syntax**

```
use stdlib
X = toeplitz(c, r)
X = toeplitz(c)
```

## **Description**

toeplitz(c,r) creates a Toeplitz matrix whose first column contains the elements of vector c and whose first row contains the elements of vector r. A Toeplitz matrix is a matrix whose diagonals have the same value. In case of conflict, the first element of r is ignored. With one argument, toeplitz gives a symmetric square matrix.

## Example

```
use stdlib
toeplitz(1:3, 1:5)
1 2 3 4 5
2 1 2 3 4
3 2 1 2 3
```

#### See also

hankel, diag

## trapz

Numeric integration with trapezoidal approximation.

## **Syntax**

```
use stdlib
s = trapz(Y)
s = trapz(X, Y)
s = trapz(X, Y, dim)
```

### **Description**

trapz(Y) calculates an approximation of the integral of a function given by the samples in Y with unit intervals. The trapezoidal approximation is used. If Y is an array, integration is performed along the first non-singleton dimension.

trapz(X,Y) specifies the location of the samples. A third argument may be used to specify along which dimension the integration is performed.

## **Example**

```
use stdlib
trapz([2, 3, 5])
6.5
trapz([1, 2, 5], [2, 3, 5])
14.5
```

#### See also

sum, cumtrapz

## **7.2** stat

stat is a library which adds to LME advanced statistical functions.
The following statement makes available functions defined in stat:

```
use stat
```

## **Functions**

## **bootstrp**

Bootstrap estimate.

Libraries — stat 595

### **Syntax**

```
use stat
(stats, samples) = bootstrp(n, fun, D1, ...)
```

### **Description**

bootstrp(n, fun, D) picks random observations from the rows of matrix (or column vector) D to form n sets which have all the same size as D; then it applies function fun (a function name or reference or an inline function) to each set and returns the results in the columns of stats. Up to three different set of data can be provided.

bootstrp gives an idea of the robustness of the estimate with respect to the choice of the observations.

## **Example**

```
use stat
D = rand(1000, 1);
bootstrp(5, @std, D)
    0.2938
    0.2878
    0.2793
    0.2859
    0.2844
```

## geomean

Geometric mean of a set of values.

## **Syntax**

```
use stat
m = geomean(A)
m = geomean(A, dim)
```

## Description

geomean(A) gives the geometric mean of the columns of array A or of the row vector A. The dimension along which geomean proceeds may be specified with a second argument.

The geometric mean of vector v of length n is defined as  $(\prod_i v_i)^{1/n}$ .

#### **Example**

```
use stat
geomean(1:10)
   4.5287
mean(1:10)
   5.5
exp(mean(log(1:10)))
   4.5287
```

#### See also

harmmean, mean

#### harmmean

Harmonic mean of a set of values.

### **Syntax**

```
use stat
m = harmmean(A)
m = harmmean(A, dim)
```

## **Description**

harmmean(A) gives the harmonic mean of the columns of array A or of the row vector A. The dimension along which harmmean proceeds may be specified with a second argument.

The inverse of the harmonic mean is the arithmetic mean of the inverse of the observations.

## **Example**

```
use stat
harmmean(1:10)
   3.4142
mean(1:10)
   5.5
```

#### See also

geomean, mean

## iqr

Interquartile range.

Libraries — stat 597

### **Syntax**

```
use stat
m = iqr(A)
m = iqr(A, dim)
```

## **Description**

iqr(A) gives the interquartile range of the columns of array A or of the row vector A. The dimension along which iqr proceeds may be specified with a second argument.

The interquartile range is the difference between the 75th percentile and the 25th percentile.

### **Example**

```
use stat
iqr(rand(1,1000))
0.5158
```

#### See also

trimmean, prctile

#### mad

Mean absolute deviation.

## **Syntax**

```
use stat
m = mad(A)
m = mad(A, dim)
```

## **Description**

mad(A) gives the mean absolute deviation of the columns of array A or of the row vector A. The dimension along which mad proceeds may be specified with a second argument.

The mean absolute deviation is the mean of the absolute value of the deviation between each observation and the arithmetic mean.

## **Example**

```
use stat
mad(rand(1,1000))
0.2446
```

#### See also

trimmean, mean, igr

### nancorrcoef

Correlation coefficients after discarding NaNs.

#### **Syntax**

```
use stat
S = nancorrcoef(X)
S = nancorrcoef(X1, X2)
```

### **Description**

nancorrcoef(X) calculates the correlation coefficients of the columns of the m-by-n matrix X. NaN values are ignored. The result is a square n-by-n matrix whose diagonal is 1.

nancorrcoef(X1,X2) calculates the correlation coefficients of X1 and X2 and returns a 2-by-2 matrix, ignoring NaN values. It is equivalent to nancorrcoef([X1(:),X2(:)]).

#### See also

nanmean, nanstd, nancov, corrcoef

#### nancov

Covariance after discarding NaNs.

## **Syntax**

```
use stat
M = nancov(data)
M = nancov(data, 0)
M = nancov(data, 1)
```

## Description

nancov(data) returns the best unbiased estimate m-by-m covariance matrix of the n-by-m matrix data for a normal distribution. NaN values are ignored. Each row of data is an observation where n quantities were measured. nancov(data,0) is the same as nancov(data).

Libraries — stat 599

nancov(data,1) returns the m-by-m covariance matrix of the n-by-m matrix data which contains the whole population; NaN values are ignored.

#### See also

nanmean, nanstd, nancorrcoef, cov

#### nanmean

Mean after discarding NaNs.

### **Syntax**

```
use stat
y = nanmean(A)
y = nanmean(A, dim)
```

### **Description**

nanmean(v) returns the arithmetic mean of the elements of vector v. nanmean(A) returns a row vector whose elements are the means of the corresponding columns of array A. nanmean(A,dim) returns the mean of array A along dimension dim; the result is a row vector if dim is 1, or a column vector if dim is 2. In all cases, NaN values are ignored.

## **Examples**

```
use stat
nanmean([1,2,nan;nan,6,7])
    1 4 7
nanmean([1,2,nan;nan,6,7],2)
    1.5
    6.5
nanmean([nan,nan])
    nan
```

#### See also

nanmedian, nanstd, mean

## nanmedian

Median after discarding NaNs.

```
use stat
y = nanmedian(A)
y = nanmedian(A, dim)
```

### **Description**

nanmedian(v) gives the median of vector v, i.e. the value x such that half of the elements of v are smaller and half of the elements are larger. NaN values are ignored.

nanmedian(A) gives a row vector which contains the median of the columns of A. With a second argument, nanmedian(A,dim) operates along dimension dim.

#### See also

nanmean, median

#### nanstd

Standard deviation after discarding NaNs.

## **Syntax**

```
use stat
y = nanstd(A)
y = nanstd(A, p)
y = nanstd(A, p, dim)
```

## **Description**

nanstd(v) returns the standard deviation of vector v with NaN values ignored, normalized by one less than the number of non-NaN values. With a second argument, nanstd(v,p) normalizes by one less than the number of non-NaN values if p is true, or by the number of non-NaN values if p is false.

nanstd(M) gives a row vector which contains the standard deviation of the columns of M. With a third argument, nanstd(M,p,dim) operates along dimension dim. In all cases, NaN values are ignored.

## Example

```
use stat
nanstd([1,2,nan;nan,6,7;10,11,12])
6.3640 4.5092 3.5355
```

Libraries — stat 601

#### See also

nanmedian, nanstd, mean

#### nansum

Sum after discarding NaNs.

### **Syntax**

```
use stat
y = nansum(A)
v = nansum(A, dim)
```

### **Description**

nansum(v) returns the sum of the elements of vector v. NaN values are ignored. nansum(A) returns a row vector whose elements are the sums of the corresponding columns of array A. nansum(A,dim) returns the sum of array A along dimension dim; the result is a row vector if dim is 1. or a column vector if dim is 2.

#### See also

nanmean, sum

## pdist

Pairwise distance between observations.

## Syntax

```
use stat
d = pdist(M)
d = pdist(M, metric)
d = pdist(M, metric, p)
```

## **Description**

pdist calculates the distance between pairs of rows of the observation matrix M. The result is a column vector which contains the distances between rows i and j with i < j. It can be resized to a square matrix with squareform.

By default, the metric used to calculate the distance is the euclidean distance; but it can be specified with a second argument:

```
'euclid' euclidean distance
'seuclid' standardized euclidean distance
'mahal' Mahalanobis distance
'cityblock' sum of absolute values
'minkowski' Minkowski metric with parameter p
```

The standardized euclidean distance is the euclidean distance after each column of M has been divided by its standard deviation. The Minkowski metric is based on the p-norm of vector differences.

### **Examples**

#### See also

squareform

## prctile

Percentile.

## **Syntax**

```
use stat
m = prctile(A, prc)
m = prctile(A, prc, dim)
```

## **Description**

prctile(A,prc) gives the smallest values larger than prc percent of the elements of each column of array A or of the row vector A. The dimension along which prctile proceeds may be specified with a third argument.

Libraries — stat 603

#### **Example**

```
prctile(rand(1,1000),90)
   0.8966
```

#### See also

trimmean, igr

## range

Data range.

### **Syntax**

```
use stat
m = range(A)
m = range(A, dim)
```

### **Description**

range(A) gives the differences between the maximum and minimum values of the columns of array A or of the row vector A. The dimension along which range proceeds may be specified with a second argument.

### **Example**

```
range(rand(1,100))
0.9602
```

#### See also

iqr

## squareform

Resize the output of pdist to a square matrix.

```
use stat
D = squareform(d)
```

squareform(d) resize d, which should be the output of pdist, into a symmetric square matrix D, so that the distance between observations i and j is D(i,j).

#### See also

pdist

### trimmean

Trimmed mean of a set of values.

### **Syntax**

```
use stat
m = trimmean(A, prc)
m = trimmean(A, prc, dim)
```

## **Description**

trimmean(A,prc) gives the arithmetic mean of the columns of array A or of the row vector A once prc/2 percent of the values have been removed from each end. The dimension along which trimmean proceeds may be specified with a third argument.

trimmean is less sensitive to outliers than the regular arithmetic mean.

#### See also

prctile, geomean, median, mean

#### zscore

Z score (normalized deviation).

```
use stat
Y = zscore(X)
Y = zscore(X, dim)
```

zscore(X) normalizes the columns of array X or the row vector X by subtracting their mean and dividing by their standard deviation. The dimension along which zscore proceeds may be specified with a second argument.

## 7.3 probdist

probdist is a library which adds to LME classes related to probability distributions. They provide an alternative interface to the algorithms in functions pdf, cdf, icdf and random. In addition, they provide methods to compute their mean, their median, their variance and their standard deviation when an explicit formula is known.

Probability distribution objects, which bundle both the distribution type and parameters, should be created with function makedist.

The following statement makes available classes defined in probdist:

```
use probdist
```

## **Functions**

## distribution::cdf

Cumulative distribution function for a distribution.

## **Syntax**

```
s = cdf(pd, x)
```

## Description

cdf(pd,x) calculates the integral of a probability density function from -infinity to x. The distribution is specified by the distribution object pd, typically created by makedist.

## **Example**

```
use probdist
pd = makedist('normal', mu=1, sigma=0.5);
x = linspace(-1, 3);
p = pdf(pd, x);
c = cdf(pd, x);
```

```
plot(x, p, '-');
plot(x, c);
```

#### See also

distribution::pdf, distribution::icdf, distribution::random,
makedist, cdf

### distribution::icdf

Inverse cumulative distribution function for a distribution.

### **Syntax**

```
x = icdf(pd, p)
```

### **Description**

icdf(pd,p) calculates the value of x such that cdf(pd,x) is p. The distribution is specified by the distribution object pd, typically created by makedist.

icdf is defined for distributions beta, chi2, gamma, lognormal, normal, student, and uniform.

## **Example**

```
use probdist
pd = makedist('student', nu=3);
p = cdf(pd, 4)
    p =
        0.9860
x = icdf(pd, p)
x =
        4.0000
```

#### See also

distribution::cdf, distribution::pdf, distribution::random,
makedist,icdf

#### makedist

Make a distribution object.

```
use probdist
pd = makedist(name, param1=value1, ...)
```

## **Description**

makedist(name) creates a distribution object with the default parameters. Parameters can be specified with named arguments. The result is an object whose class is a subclass of distribution.

Here is a list of distributions with the default parameter values.

Name	<b>Default parameters</b>	Class
'beta'	a=1,b=1	betaDistribution
'chi'	nu=1	chiDistribution
'chi2'	nu=1	chi2Distribution
'exp'	mu=1	exponentialDistribution
'logn'	mu=1,sigma=1	lognormalDistribution
'nakagami'	mu=1,omega=1	nakagamiDistribution
'norm'	mu=0,sigma=1	normalDistribution
'rayl'	b=1	rayleighDistribution
't'	nu=1	studentDistribution
'unif'	Lower=0,Upper=1	uniformDistribution
'weib'	a=1,b=1	weibullDistribution

## **Example**

```
use probstat
pd = makedist('chi2', nu=3)
pd =
   Chi2 distribution
m_th = mean(pd)
m_th =
   3
m_data = mean(random(pd, [1, 10000]))
m_data =
   3.0027
```

## distribution::mean

Mean of a distribution.

```
m = mean(pd)
```

mean(pd) gives the arithmetic mean of a distribution.

### **Example**

```
use probdist
pd = makedist('normal', mu=3, sigma=2);
mean(pd)
   3
```

#### See also

```
distribution::var, distribution::sdev, distribution::median,
makedist, mean
```

#### distribution::median

Median of a distribution.

### **Syntax**

```
m = median(pd)
```

## **Description**

median(pd) gives the arithmetic median of a distribution, or NaN if it cannot be computed.

## Example

```
use probdist
pd = makedist('exp', mu=2);
median(pd)
3
```

#### See also

```
distribution::var, distribution::sdev, distribution::median,
makedist, median
```

## distribution::pdf

Probability density function of a distribution.

```
s = pdf(pd, x)
```

## **Description**

pdf(pd,x) gives the probability of a distribution. The distribution is specified by the distribution object pd, typically created by makedist.

### **Example**

```
use probdist
pd = makedist('lognormal', mu=2, sigma=1.5);
x = logspace(-2,1);
p = pdf(pd, x);
plot(x, p);
```

#### See also

```
distribution::cdf, distribution::icdf, distribution::random,
makedist, pdf
```

### distribution::random

Random generator for a distribution.

## **Syntax**

```
x = random(pd)
x = random(pd, size)
```

## **Description**

random(pd) calculates a pseudo-random number whose distribution function is specified by the distribution object pd, typically created by makedist.

Additional input arguments specify the size of the result, either as a vector (or a single scalar for a square matrix) or as scalar values. The result is an array of the specified size where each value is an independent pseudo-random variable. The default size is 1 (scalar).

## **Example**

```
use probdist
pd = makedist('exp');
dataSize = [10, 100];
data = random(pd, dataSize);
```

#### See also

```
distribution::pdf, makedist, random
```

#### distribution::std

Standard deviation of a distribution.

### **Syntax**

```
s = std(pd)
```

### **Description**

std(pd) gives the standard deviation of a distribution.

## **Example**

```
use probdist
pd = makedist('lognormal', mu=2, sigma=1.5);
std(pd)
   66.3080
std(random(pd,[1,100000]))
   68.0868
```

#### See also

```
distribution::var, distribution::mean, distribution::median,
makedist, std
```

#### distribution::var

Variance of a distribution.

## **Syntax**

```
s2 = var(pd)
```

## Description

var(pd) gives the variance of a distribution.

### **Example**

```
use probdist
pd = makedist('uniform', Lower=2, Upper=10);
var(pd)
   5.3333
var(random(pd,[1,100000]))
   5.3148
```

#### See also

```
distribution::mean,
distribution::sdev,
distribution::median, makedist, var
```

## 7.4 polynom

Library polynom implements the constructors and methods of two classes: polynom for polynomials, and ratfun for rational functions. Basic arithmetic operators and functions are overloaded to support expressions with the same syntax as for numbers and matrices.

The following statement makes available functions defined in polynom:

```
use polynom
```

Methods for conversion to MathML are defined in library polynom\_mathml. Both libraries can be loaded with a single statement:

```
use polynom, polynom_mathml
```

## **Functions**

## polynom::polynom

Polynom object constructor.

```
use polynom
p = polynom
p = polynom(coef)
```

polynom(coef) creates a polynom object initialized with the coefficients in vector coef, given in descending powers of the variable. Without argument, polynom returns a polynom object initialized to 0.

The following operators and functions may be used with polynom arguments, with results analog to the corresponding functions of LME. Function roots ignores leading zero coefficients.

```
- minus + plus
^ mpower rem
\ mldivide roots
/ mrdivide - uminus
* mtimes + uplus
```

### **Examples**

```
use polynom
p = polynom([3,0,1,-4,2])
p =
    3x^4+x^2-4x+2
q = 3 * p^2 + 8
q =
    27x^8+18x^6-72x^5+39x^4-24x^3+60x^2-48x+20
```

#### See also

```
polynom::disp, polynom::double, polynom::subst,
polynom::diff, polynom::int, polynom::inline, polynom::feval,
ratfun::ratfun
```

## polynom::disp

Display a polynom object.

## **Syntax**

```
use polynom
disp(p)
```

## **Description**

disp(p) displays polynomial p. It is also executed implicitly when LME displays the polynom result of an expression which does not end with a semicolon.

## **Example**

```
use polynom
p = polynom([3,0,1,-4,2])
p =
    3x^4+x^2-4x+2
```

### See also

```
polynom::polynom, disp
```

# polynom::double

Convert a polynom object to a vector of coefficients.

### **Syntax**

```
use polynom
coef = double(p)
```

## **Description**

double(p) converts polynomial p to a row vector of descending-power coefficients.

### **Example**

```
use polynom
p = polynom([3,0,1,-4,2]);
double(p)
3 0 1 -4 2
```

#### See also

```
polynom::polynom
```

# polynom::subst

Substitute the variable of a polynom object with another polynomial.

```
use polynom
subst(a, b)
```

## **Description**

subst(a,b) substitutes the variable of polynom a with polynom b.

## **Example**

```
use polynom
p = polynom([1,2,3])
p =
    x^2+3x+9
q = polynom([2,0])
    q =
     2x
r = subst(p, q)
r =
    4x^2+6x+9
```

### See also

```
polynom::polynom, polynom::feval
```

# polynom::diff

Polynom derivative.

## **Syntax**

```
use polynom
diff(p)
```

# **Description**

diff(p) differentiates polynomial p.

# **Example**

```
use polynom
p = polynom([3,0,1,-4,2]);
q = diff(p)
q =
    12x^3+2x-4
```

#### See also

```
polynom::polynom, polynom::int, polyder
```

# polynom::int

Polynom integral.

### **Syntax**

```
use polynom
int(p)
```

## **Description**

int(p) integrates polynomial p.

## **Example**

```
use polynom
p = polynom([3,0,1,-4,2]);
q = int(p)
q =
    0.6x^5+0.3333x^3-2x^2+2x
```

#### See also

```
polynom::polynom, polynom::diff, polyint
```

# polynom::inline

Conversion from polynom object to inline function.

# **Syntax**

```
use polynom
fun = inline(p)
```

# **Description**

inline(p) converts polynomial p to an inline function which can then be used with functions such as feval and ode45.

# **Example**

```
use polynom
p = polynom([3,0,1,-4,2]);
fun = inline(p)
  fun =
     <inline function>
```

```
dumpvar('fun', fun);
fun = inline('function y=f(x);y=polyval([3,0,1,-4,2],x);');
```

#### See also

```
polynom::polynom, polynom::feval, ode45
```

# polynom::feval

Evaluate a polynom object.

## **Syntax**

```
use polynom
y = feval(p, x)
```

## **Description**

feval(p,x) evaluates polynomial p for the value of x. If x is a vector or a matrix, the evaluation is performed separately on each element and the result has the same size as x.

## **Example**

```
use polynom
p = polynom([3,0,1,-4,2]);
y = feval(p, 1:5)
y =
    2    46    242    770    1882
```

#### See also

```
polynom::polynom, polynom::inline, feval
```

# polynom::mathml

Conversion to MathML.

```
use polynom, polynom_mathml
str = mathml(p)
str = mathml(p, false)
```

# **Description**

mathml(p) converts its argument p to MathML presentation, returned as a string.

By default, the MathML top-level element is <math>. If the result is to be used as a MathML subelement of a larger equation, a last input argument equal to the logical value false can be specified to suppress <math>.

## **Example**

```
use polynom, polynom_mathml
p = polynom([3,0,1,-4,2]);
m = mathml(p);
math(0, 0, m);
```

#### See also

mathmlpoly, mathml

### ratfun::ratfun

Ratfun object constructor.

# Syntax

```
use polynom
r = ratfun
r = ratfun(coefnum)
r = ratfun(coefnum, coefden)
```

# **Description**

ratfun(coefnum, coefden) creates a ratfun object initialized with the coefficients in vectors coefnum and coefden, given in descending powers of the variable. Without argument, ratfun returns a ratfun object initialized to 0. If omitted, coefden defaults to 1.

The following operators and functions may be used with ratfun arguments, with results analog to the corresponding functions of LME.

```
inv * mtimes
- minus + plus
\ mldivide - uminus
^ mpower + uplus
/ mrdivide
```

### **Example**

```
use polynom

r = ratfun([3,0,1,-4,2], [2,5,0,1])

r = (3x^4+x^2-4x+2)/(2x^3+5x^2+1)
```

#### See also

```
ratfun::disp, ratfun::inline, ratfun::feval, polynom::polynom
```

# ratfun::disp

Display a ratfun object.

## **Syntax**

```
use polynom
disp(r)
```

## **Description**

disp(r) displays rational function r. It is also executed implicitly when LME displays the ratfun result of an expression which does not end with a semicolon.

#### See also

```
ratfun::ratfun, disp
```

### ratfun::num

Get the numerator of a ratfun as a vector of coefficients.

## **Syntax**

```
use polynom
coef = num(r)
```

# **Description**

num(r) gets the numerator of r as a row vector of descending-power coefficients.

#### See also

```
ratfun::den, ratfun::ratfun
```

## ratfun::den

Get the denominator of a ratfun as a vector of coefficients.

## **Syntax**

```
use polynom
coef = den(a)
```

## **Description**

den(a) gets the denominator of a as a row vector of descending-power coefficients.

#### See also

```
ratfun::num, ratfun::ratfun
```

# ratfun::diff

Ratfun derivative.

## **Syntax**

```
use polynom
diff(r)
```

# **Description**

diff(r) differentiates ratfun r.

# **Example**

```
use polynom
r = ratfun([1,3,0,1],[2,5]);
q = diff(r)
q =
    (4x^3+21x^2+30x-2)/(4x^2+20x+25)
```

#### See also

ratfun::ratfun

# ratfun::inline

Conversion from ratfun to inline function.

### **Syntax**

```
use polynom
fun = inline(r)
```

### **Description**

inline(r) converts ratfun r to an inline function which can then be used with functions such as feval and ode45.

#### See also

```
ratfun::ratfun, ratfun::feval, ode45
```

### ratfun::feval

Evaluate a ratfun object.

## **Syntax**

```
use polynom
v = feval(r, x)
```

# **Description**

feval(r,x) evaluates ratfun r for the value of x. If x is a vector or a matrix, the evaluation is performed separately on each element and the result has the same size as x.

# **Example**

```
use polynom
r = ratfun([1,3,0,1],[2,5]);
y = feval(r, 1:5)
y =
    0.7143    2.3333    5.0000    8.6923    13.4000
```

#### See also

```
ratfun::ratfun, ratfun::inline, feval
```

Libraries — ratio 621

## ratfun::mathml

Conversion to MathML.

## **Syntax**

```
use polynom, polynom_mathml
str = mathml(r)
str = mathml(r, false)
```

## **Description**

mathml(r) converts its argument r to MathML presentation, returned as a string.

By default, the MathML top-level element is <math>. If the result is to be used as a MathML subelement of a larger equation, a last input argument equal to the logical value false can be specified to suppress <math>.

## **Example**

```
use polynom, polynom_mathml
r = ratfun([1,3,0,1],[2,5]);
m = mathml(r);
math(0, 0, m);
```

#### See also

mathml

# 7.5 ratio

Library ratio implements the constructors and methods of class ratio for rational numbers. It is based on long integers, so that the precision is limited only by available memory. Basic arithmetic operators and functions are overloaded to support expressions with the same syntax as for numbers.

The following statement makes available functions defined in ratio:

```
use ratio
```

# **Functions**

## ratio::ratio

Ratio object constructor.

## **Syntax**

```
use ratio
r = ratio
r = ratio(n)
r = ratio(num, den)
r = ratio(r)
```

### **Description**

ratio(num, den) creates a rational fraction object whose value is num/den. Arguments num and den may be double integer numbers or longint. Common factors are canceled out. With one numeric input argument, ratio(n) creates a rational fraction whose denominator is 1. Without input argument, ratio creates a rational number whose value is 0.

With one input argument which is already a ratio object, ratio returns it without change.

The following operators and functions may be used with ratio objects, with results analog to the corresponding functions of LME.

```
mldivide
    eq
>=
    ge
               mpower
    qt
               mrdivide
>
    inv
               mtimes
<=
    le
               ne
    lt
               plus
<
               uminus
    max
               uplus
    min
    minus
```

## **Examples**

```
use ratio
r = ratio(2, 3)
r =
    2/3
q = 5 * r - 1
q =
    7/3
```

Libraries — ratio 623

#### See also

```
ratio::disp, ratio::double, ratio::char
```

## ratio::char

Display a ratio object.

## **Syntax**

```
use ratio char(r)
```

## **Description**

char(r) converts ratio r to a character string.

#### See also

```
ratio::ratio, ratio::disp, char
```

# ratio::disp

Display a ratio object.

# Syntax

```
use ratio
disp(r)
```

# **Description**

disp(r) displays ratio r with the same format as char. It is also executed implicitly when LME displays the ratio result of an expression which does not end with a semicolon.

#### See also

```
ratio::ratio, ratio::char, disp
```

## ratio::double

Convert a ratio object to a floating-point number.

## **Syntax**

```
use ratio
x = double(r)
```

### **Description**

double(r) converts ratio r to a floating-point number of class double.

### **Example**

```
use ratio
r = ratio(2, 3);
double(r)
    0.6666
```

#### See also

ratio::ratio

# 7.6 bitfield

Library bitfield implements the constructor and methods of class bitfield for bit fields (binary numbers). Basic arithmetic operators and functions are overloaded to support expressions with the same syntax as for numbers and matrices. Contrary to integer numbers, bitfield objects have a length (between 1 and 32) and are displayed in binary.

The following statement makes available functions defined in bitfield:

use bitfield

# **Functions**

# bitfield::beginning

First bit position in a bitfield.

```
use bitfield
a(...beginning...)
```

### **Description**

In the index expression of a bitfield, beginning is the position of the least-significant bit, i.e. 0.

#### See also

bitfield::bitfield, bitfield::end

### bitfield::bitfield

Bitfield object constructor.

### **Syntax**

```
use bitfield
a = bitfield
a = bitfield(n)
a = bitfield(n, wordlength)
```

### **Description**

bitfield(n,wordlength) creates a bitfield object initialized with the wordlength least significant bits of the nonnegative integer number n. The default value of wordlength is 32 if n is a double, an int32 or a uint32 number; 16 is n is an int16 or uint16 number; or 8 if n is an int8 or uint8 number. Without argument, bitfield gives a bit field of 32 bits 0. Like any integer number in LME, n may be written in base 2, 8, 10, or 16: 0b1100, 014, 12, and 0xc all represent the same number.

The following operators and functions may be used with bitfield arguments, with results analog to the corresponding functions of LME. Logical functions operate bitwise.

```
& and ~ not
== eq | or
- minus + plus
\ mldivide - uminus
/ mrdivide + uplus
* mtimes xor
~= ne
```

Indexes into bit fields are non-negative integers: 0 represents the least-significant bit, and wordlength-1 the most-significant bit. Unlike arrays, bits are not selected with logical arrays, but with other bit fields where ones represent the bits to be selected; for example a(0b1011) selects bits 0, 1 and 3. This is consistent with the way bitfield::find is defined.

### **Examples**

## See also

bitfield::disp, bitfield::double

# bitfield::disp

Display a bitfield object.

### **Syntax**

```
use bitfield
disp(a)
```

## **Description**

disp(a) displays bitfield a. It is also executed implicitly when LME displays the bitfield result of an expression which does not end with a semicolon.

#### See also

```
bitfield::bitfield, disp
```

# bitfield::double

Convert a bitfield object to a double number.

```
use bitfield
n = double(a)
```

## **Description**

double(a) converts bitfield a to double number.

## **Example**

```
use bitfield
a = bitfield(123, 16);
double(a)
   123
```

#### See also

bitfield::bitfield

## bitfield::end

Last bit position in a bitfield.

### **Syntax**

```
use bitfield
a(...end...)
```

# **Description**

In the index expression of a bitfield, end is the position of the most-significant bit, i.e. 1 less than the word length.

#### See also

```
bitfield::bitfield, bitfield::beginning
```

# bitfield::find

Find the ones in a bitfield.

# **Syntax**

```
use bitfield
ix = find(a)
```

# **Description**

find(a) finds the bits equal to 1 in bitfield a. The result is a vector of bit positions in ascending order; the least-significant bit is number 0.

### **Example**

```
use bitfield
a = bitfield(123, 16)
a =
          0b0000000001111011
ix = find(a)
    ix =
          0 1 3 4 5 6
```

#### See also

bitfield::bitfield.find

# bitfield::int8 bitfield::int16 bitfield::int32

Convert a bitfield object to a signed integer number, with sign extension.

## **Syntax**

```
use bitfield
n = int8(a)
n = int16(a)
n = int32(a)
```

# **Description**

int8(a), int16(a), and int32(a) convert bitfield a to an int8, int16, or int32 number respectively. If a has less bits than the target integer and the most significant bit of a is 1, sign extension is performed; i.e. the most significant bits of the result are set to 1, so that it is negative. If a has more bits than the target integer, most significant bits are ignored.

## Example

#### See also

```
uint8, uint16, uint32, bitfield::int8, bitfield::int16,
bitfield::int32, bitfield::double, bitfield::bitfield
```

# bitfield::length

Word length of a bitfield.

## **Syntax**

```
use bitfield
wordlength = length(a)
```

## **Description**

length(a) gives the number of bits of bitfield a.

## **Example**

```
use bitfield
a = bitfield(123, 16);
length(a)
16
```

#### See also

bitfield::bitfield, length

# bitfield::sign

Get the sign of a bitfield.

# **Syntax**

```
use bitfield
s = sign(a)
```

# **Description**

sign(a) gets the sign of bitfield a. The result is -1 if the most-significant bit of a is 1, 0 if all bits of a are 0, or 1 otherwise.

### **Example**

```
use bitfield
a = bitfield(5, 3)
a =
    0b101
sign(a)
    -1
```

#### See also

bitfield::bitfield, sign

# bitfield::uint8 bitfield::uint16 bitfield::uint32

Convert a bitfield object to an unsigned integer number.

### **Syntax**

```
use bitfield
n = uint8(a)
n = uint16(a)
n = uint32(a)
```

# **Description**

uint8(a), uint16(a), and uint32(a) convert bitfield a to a uint8, uint16, or uint32 number respectively. If a has more bits than the target integer, most significant bits are ignored.

### **Example**

```
use bitfield
a = bitfield(1234, 16);
uint8(a)
210
```

#### See also

```
uint8, uint16, uint32, bitfield::int8, bitfield::int16,
bitfield::int32, bitfield::double, bitfield::bitfield
```

# 7.7 filter

filter is a library which adds to LME functions for designing analog (continuous-time) and digital (discrete-time) linear filters.

The following statement makes available functions defined in filter:

use filter

This library provides three kinds of functions:

- besselap, buttap, cheblap, cheblap, and ellipap, which compute the zeros, poles and gain of the prototype of analog low-pass filter with a cutoff frequency of 1 rad/s. They correspond respectively to Bessel, Butterworth, Chebyshev type 1, Chebyshev type 2, and elliptic filters.
- besself, butter, cheby1, cheby2, and ellip, which provide a higher-level interface to design filters of these different types. In addition to the filter parameters (degree, bandpass and bandstop ripples), one can specify the kind of filter (lowpass, highpass, bandpass or bandstop) and the cutoff frequency or frequencies. The result can be an analog or a digital filter, given as a rational transfer function or as zeros, poles and gain.
- lp2lp, lp2hp, lp2bp, and lp2bs, which convert analog lowpass filters respectively to lowpass, highpass, bandpass, and bandstop with specified cutoff frequency or frequencies.

Transfer functions are expressed as the coefficient vectors of their numerator num and denominator den in decreasing powers of s (Laplace transform for analog filters) or z (z transform for digital filters); or as the zeros z, poles p, and gain k.

# **Functions**

# besselap

Bessel analog filter prototype.

```
use filter
(z, p, k) = besselap(n)
```

### **Description**

besselap(n) calculates the zeros, the poles, and the gain of a Bessel analog filter of degree n with a cutoff angular frequency of 1 rad/s.

#### See also

besself, buttap, cheb1ap, cheb2ap, ellipap

### besself

Bessel filter.

### **Syntax**

```
use filter
(z, p, k) = besself(n, w0)
(num, den) = besself(n, w0)
(...) = besself(n, [wl, wh])
(...) = besself(n, w0, 'high')
(...) = besself(n, [wl, wh], 'stop')
(...) = besself(..., 's')
```

## **Description**

besself calculates a Bessel filter. The result is given as zeros, poles and gain if there are three output arguments, or as numerator and denominator coefficient vectors if there are two output arguments.

besself(n,w0), where w0 is a scalar, gives a digital lowpass filter of order n with a cutoff frequency of w0 relatively to half the sampling frequency.

besself(n, [wl,wh]), where the second input argument is a vector of two numbers, gives a digital bandpass filter of order 2\*n with passband between wl and wh relatively to half the sampling frequency.

besself(n,w0,'high') gives a digital highpass filter of order n with a cutoff frequency of w0 relatively to half the sampling frequency.

besself(n,[wl,wh],'stop'), where the second input argument is a vector of two numbers, gives a digital bandstop filter of order 2\*n with stopband between wl and wh relatively to half the sampling frequency.

With an additional input argument which is the string 's', besself gives an analog Bessel filter. Frequencies are given in rad/s.

#### See also

besselap, butter, cheby1, cheby2, ellip

### bilinear

Analog-to-digital conversion with bilinear transformation.

## **Syntax**

```
use filter
(zd, pd, kd) = bilinear(zc, pc, kc, fs)
(numd, dend) = bilinear(numc, denc, fs)
```

## **Description**

bilinear(zc,pc,kc,fs) converts the analog (continuous-time) transfer function given by its zeros zc, poles pc, and gain kc to a digital (discrete-time) transfer function given by its zeros, poles, and gain in the domain of the forward-shift operator q. The sampling frequency is fs. Conversion is performed with the bilinear transormation  $z_d = (1 + z_c/2f_s)/(1 - z_c/2f_s)$ . If the analog transfer function has less zeros than poles, additional digital zeros are added at -1 to avoid a delay.

With three input arguments, bilinear(numc,denc,fs) uses the coefficients of the numerators and denominators instead of their zeros, poles and gain.

# buttap

Butterworth analog filter prototype.

# Syntax

```
use filter
(z, p, k) = buttap(n)
```

# **Description**

buttap(n) calculates the zeros, the poles, and the gain of a Butterworth analog filter of degree n with a cutoff angular frequency of 1 rad/s.

#### See also

butter, besselap, cheb1ap, cheb2ap, ellipap

### butter

Butterworth filter.

### Syntax

```
use filter
(z, p, k) = butter(n, w0)
(num, den) = butter(n, w0)
(...) = butter(n, [wl, wh])
(...) = butter(n, w0, 'high')
(...) = butter(n, [wl, wh], 'stop')
(...) = butter(..., 's')
```

### **Description**

butter calculates a Butterworth filter. The result is given as zeros, poles and gain if there are three output arguments, or as numerator and denominator coefficient vectors if there are two output arguments.

butter(n,w0), where w0 is a scalar, gives a nth-order digital lowpass filter with a cutoff frequency of w0 relatively to half the sampling frequency.

butter(n, [wl,wh]), where the second input argument is a vector of two numbers, gives a 2nth-order digital bandpass filter with passband between wl and wh relatively to half the sampling frequency.

butter(n,w0,'high') gives a nth-order digital highpass filter with a cutoff frequency of w0 relatively to half the sampling frequency.

butter(n, [wl,wh], 'stop'), where the second input argument is a vector of two numbers, gives a 2nth-order digital bandstop filter with stopband between wl and wh relatively to half the sampling frequency.

With an additional input argument which is the string 's', butter gives an analog Butterworth filter. Frequencies are given in rad/s.

#### See also

buttap, besself, cheby1, cheby2, ellip

# cheb1ap

Chebyshev type 1 analog filter prototype.

```
use filter
(z, p, k) = cheblap(n, rp)
```

### **Description**

cheblap(n,rp) calculates the zeros, the poles, and the gain of a Chebyshev type 1 analog filter of degree n with a cutoff angular frequency of 1 rad/s. Ripples in the passband have a peak-to-peak magnitude of rp dB, i.e. the peak-to-peak ratio is  $10^{\circ}(rp/20)$ .

#### See also

cheby1, cheb2ap, ellipap, besselap, buttap

# cheb2ap

Chebyshev type 2 analog filter prototype.

## **Syntax**

```
use filter
(z, p, k) = cheb2ap(n, rs)
```

## **Description**

cheb2ap(n,rs) calculates the zeros, the poles, and the gain of a Chebyshev type 2 analog filter of degree n with a cutoff angular frequency of 1 rad/s. Ripples in the stopband have a peak-to-peak magnitude of rs dB, i.e. the peak-to-peak ratio is  $10^{\circ}(rs/20)$ .

#### See also

cheby1, cheb1ap, ellipap, besselap, buttap

# cheby1

Chebyshev type 1 filter.

```
use filter
(z, p, k) = cheby1(n, rp, w0)
(num, den) = cheby1(n, rp, w0)
(...) = cheby1(n, rp, [wl, wh])
(...) = cheby1(n, rp, w0, 'high')
(...) = cheby1(n, rp, [wl, wh], 'stop')
(...) = cheby1(..., 's')
```

### **Description**

cheby1 calculates a Chebyshev type 1 filter. The result is given as zeros, poles and gain if there are three output arguments, or as numerator and denominator coefficient vectors if there are two output arguments.

cheby1(n,rp,w0), where w0 is a scalar, gives a nth-order digital lowpass filter with a cutoff frequency of w0 relatively to half the sampling frequency. Ripples in the passband have a peak-to-peak magnitude of rp dB, i.e. the peak-to-peak ratio is  $10^(rp/20)$ .

cheby1(n,rp,[wl,wh]), where the second input argument is a vector of two numbers, gives a 2nth-order digital bandpass filter with passband between wl and wh relatively to half the sampling frequency.

cheby1(n,rp,w0,'high') gives a nth-order digital highpass filter with a cutoff frequency of w0 relatively to half the sampling frequency. cheby1(n,rp,[wl,wh],'stop'), where the second input argument is a vector of two numbers, gives a 2nth-order digital bandstop filter with stopband between wl and wh relatively to half the sampling frequency.

With an additional input argument which is the string 's', cheby1 gives an analog Chebyshev type 1 filter. Frequencies are given in rad/s.

#### See also

cheblap, besself, butter, cheby2, ellip

# cheby2

Chebyshev type 2 filter.

# **Syntax**

```
use filter
(z, p, k) = cheby2(n, rs, w0)
(num, den) = cheby2(n, rs, w0)
(...) = cheby2(n, rs, [wl, wh])
(...) = cheby2(n, rs, w0, 'high')
(...) = cheby2(n, rs, [wl, wh], 'stop')
(...) = cheby2(..., 's')
```

# **Description**

cheby2 calculates a Chebyshev type 2 filter. The result is given as zeros, poles and gain if there are three output arguments, or as nu-

merator and denominator coefficient vectors if there are two output arguments.

cheby2(n,rs,w0), where w0 is a scalar, gives a nth-order digital lowpass filter with a cutoff frequency of w0 relatively to half the sampling frequency. Ripples in the stopband have a peak-to-peak magnitude of rs dB, i.e. the peak-to-peak ratio is 10^(rs/20).

cheby2(n,rs,[wl,wh]), where the second input argument is a vector of two numbers, gives a 2nth-order digital bandpass filter with passband between wl and wh relatively to half the sampling frequency.

cheby2(n,rs,w0,'high') gives a nth-order digital highpass filter with a cutoff frequency of w0 relatively to half the sampling frequency. cheby2(n,rs,[wl,wh],'stop'), where the second input argument is a vector of two numbers, gives a 2nth-order digital bandstop filter with stopband between wl and wh relatively to half the sampling frequency.

With an additional input argument which is the string 's', cheby2 gives an analog Chebyshev type 2 filter. Frequencies are given in rad/s.

#### See also

cheb2ap, besself, butter, cheby1, ellip

# ellip

Elliptic filter.

# **Syntax**

```
use filter
(z, p, k) = ellip(n, rp, rs, w0)
(num, den) = ellip(n, rp, rs, w0)
(...) = ellip(n, rp, rs, [wl, wh])
(...) = ellip(n, rp, rs, w0, 'high')
(...) = ellip(n, rp, rs, [wl, wh], 'stop')
(...) = ellip(..., 's')
```

# **Description**

ellip calculates a elliptic filter, or Cauer filter. The result is given as zeros, poles and gain if there are three output arguments, or as numerator and denominator coefficient vectors if there are two output arguments.

ellip(n,rp,rs,w0), where w0 is a scalar, gives a nth-order digital lowpass filter with a cutoff frequency of w0 relatively to half the sampling frequency. Ripples have a peak-to-peak magnitude of rp dB in the passband and of rs dB in the stopband (peak-to-peak ratios are respectively  $10^{\circ}(rp/20)$  and  $10^{\circ}(rs/20)$ ).

ellip(n,rp,rs,[wl,wh]), where the second input argument is a vector of two numbers, gives a 2nth-order digital bandpass filter with passband between wl and wh relatively to half the sampling frequency.

ellip(n,rp,rs,w0,'high') gives a nth-order digital highpass filter with a cutoff frequency of w0 relatively to half the sampling frequency.

ellip(n,rp,rs,[wl,wh],'stop'), where the second input argument is a vector of two numbers, gives a 2nth-order digital bandstop filter with stopband between wl and wh relatively to half the sampling frequency.

With an additional input argument which is the string 's', ellip gives an analog elliptic filter. Frequencies are given in rad/s.

#### See also

ellipap, besself, butter, cheby1, cheby2

# ellipap

Elliptic analog filter prototype.

# Syntax

```
use filter
(z, p, k) = ellipap(n, rp, rs)
```

# **Description**

ellipap(n,rp,rs) calculates the zeros, the poles, and the gain of an elliptic analog filter of degree n with a cutoff angular frequency of 1 rad/s. Ripples have a peak-to-peak magnitude of rp dB in the passband and of rs dB in the stopband (peak-to-peak ratios are respectively  $10^{(rp/20)}$  and  $10^{(rs/20)}$ ).

### See also

ellip, cheblap, cheblap, besselap, buttap

# lp2bp

Lowpass prototype to bandpass filter conversion.

### **Syntax**

```
use filter

(z, p, k) = lp2bp(z0, p0, k0, wc, ww)

(num, den) = lp2bp(num0, den0, wc, ww)
```

## **Description**

lp2bp convert a lowpass analog filter prototype (with unit angular frequency) to a bandpass analog filter with the specified center angular frequency w0 and bandwidth ww. lp2bp(z0,p0,k0,wc,ww) converts a filter given by its zeros, poles, and gain; lp2bp(num0,den0,wc,ww) converts a filter given by its numerator and denominator coefficients in decreasing powers of s.

The new filter F(s) is

$$F(s) = F_0 \left( \frac{s^2 + \omega_c^2 - \omega_w^2 / 4}{\omega_w s} \right)$$

where  $F_0(s)$  is the filter prototype. The filter order is doubled.

#### See also

lp2lp, lp2hp, lp2bs

# lp2bs

Lowpass prototype to bandstop filter conversion.

# Syntax

```
use filter

(z, p, k) = lp2bs(z0, p0, k0, wc, ww)

(num, den) = lp2bs(num0, den0, wc, ww)
```

# **Description**

lp2bs convert a lowpass analog filter prototype (with unit angular frequency) to a bandstop analog filter with the specified center angular frequency w0 and bandwidth ww. lp2bs(z0,p0,k0,wc,ww) converts a filter given by its zeros, poles, and gain; lp2bs(num0,den0,wc,ww)

converts a filter given by its numerator and denominator coefficients in decreasing powers of s.

The new filter F(s) is

$$F(s) = F_0 \left( \frac{\omega_W s}{s^2 + \omega_C^2 - \omega_W^2 / 4} \right)$$

where  $F_0(s)$  is the filter prototype. The filter order is doubled.

#### See also

lp2lp, lp2hp, lp2bp

# lp2hp

Lowpass prototype to highpass filter conversion.

## **Syntax**

```
use filter

(z, p, k) = lp2hp(z0, p0, k0, w0)

(num, den) = lp2hp(num0, den0, w0)
```

# Description

lp2hp convert a lowpass analog filter prototype (with unit angular frequency) to a highpass analog filter with the specified cutoff angular frequency w0. lp2hp(z0,p0,k0,w0) converts a filter given by its zeros, poles, and gain; lp2hp(num0,den0,w0) converts a filter given by its numerator and denominator coefficients in decreasing powers of s.

The new filter F(s) is

$$F(s) = F_0(\frac{1}{\omega_0 s})$$

where  $F_0(s)$  is the filter prototype.

#### See also

lp2lp, lp2bp, lp2bs

# lp2lp

Lowpass prototype to lowpass filter conversion.

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### **Syntax**

```
use filter
(z, p, k) = lp2lp(z0, p0, k0, w0)
(num, den) = lp2lp(num0, den0, w0)
```

### **Description**

lp2lp convert a lowpass analog filter prototype (with unit angular frequency) to a lowpass analog filter with the specified cutoff angular frequency w0. lp2lp(z0,p0,k0,w0) converts a filter given by its zeros, poles, and gain; lp2lp(num0,den0,w0) converts a filter given by its numerator and denominator coefficients in decreasing powers of s.

The new filter F(s) is

$$F(s) = F_0 \left( \frac{s}{\omega_0} \right)$$

where  $F_0(s)$  is the filter prototype.

#### See also

lp2hp, lp2bp, lp2bs

# 7.8 Iti

Library lti defines methods related to objects which represent linear time-invariant dynamical systems. LTI systems may be used to model many different systems: electro-mechanical devices, robots, chemical processes, filters, etc. LTI systems map one or more inputs u to one or more outputs y. The mapping is defined as a state-space model or as a matrix of transfer functions, either in continuous time or in discrete time. Methods are provided to create, combine, and analyze LTI objects.

Graphical methods are based on the corresponding graphical functions; the numerator and denominator coefficient vectors or the statespace matrices are replaced with an LTI object. They accept the same optional arguments, such as a character string for the style.

The following statement makes available functions defined in lti:

```
use lti
```

Methods for conversion to MathML are defined in library lti\_mathml. Both libraries can be loaded with a single statement:

```
use lti, lti_mathml
```

### Class overview

The LTI library defines six classes. The three central ones correspond to the main model structures used for linear time-invariant systems in automatic control: ss for state-space models, tf for rational transfer functions given by the coefficients of the numerator and denominator polynomials, and zpk for rational transfer functions given by their zeros, poles and gain. State-space representation is restricted to causal systems, while transfer functions can be non-causal. Three additional classes are more specialized: frd (frequency response data) for systems described by a discrete set of frequency/complex response pairs, and pid or pidstd for PID controllers.

LTI classes share many properties and methods. They can represent systems with single or multiple inputs and/or outputs. Inputs, outputs and internal states are continuous in time (continuous-time systems) or defined at a fixed sampling frequency (discrete-time systems).

The variable of the Laplace transform can be 's' or 'p'. The variable of the z transform can be 'z' or 'q'. By multiplying the numerator and the denominator of a rational transfer function by a suitable power of  $q^{-1}$  (or  $z^{-1}$ ), polynomials in  $q^{-1}$  can be obtained, where  $q^{-1}$  is the delay operator; this yields directly a recurrence relation.

## Conversion

Conversion between ss, tf and zpk can be done simply by calling the target constructor. The only restriction is that systems to be converted to state-space models must be causal. For instance, a transfer function given by its zeros, poles and gain can be converted to a statespace model as follows:

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```
D = 0
```

Conversion from pid or pidstd objects is performed the same way. Conversion to pid or pidstd objects is possible only if the system to be converted has the structure of a P, Pl, PD, or PID controller, with or without filter on the derivative term.

Conversion to an frd object requires an array of frequency points where the frequency response is evaluated. Conversion of frd objects to other LTI objects is not possible.

Conversion between continuous-time and discrete-time objects of the same class is performed with c2d and d2c.

# **Building large systems**

Simple systems can be combined to create larger ones. All systems can be seen as matrices mapping inputs to outputs via a matrix product. Larger systems can be created by matrix concatenation, addition or multiplication. More specialized connections can be obtained with methods connect and feedback.

Mixing objects of different classes is possible for all classes except for frd (where a frequency array must be provided explicitly, which can only be done with a call of the frd constructor). Continuous-time objects cannot be connected with discrete-time objects, and discrete-time objects must have the same sampling period.

# **Functions**

# frd::frd

LTI frequency response data constructor.

# Syntax

```
use lti
a = frd
a = frd(resp, freq)
a = frd(resp, freq, Ts)
```

# **Description**

frd(response, frequency, Ts) creates an LTI object which represents a discrete set of frequency response data. Argument response is an array of complex frequency responses corresping to frequency array freq. A single-input single-output (SISO) PID controller has scalar parameters. If the parameters are matrices, they must all have the same size (scalar values are replicated as required), and the resulting controller has as many inputs as parameters have columns and as many outputs as parameters have rows; mapping from each input to each output is and independent SISO PID controller.

### **Examples**

Simple continuous-time frd object:

```
use lti
freq = 0:100;
resp = 3 ./ (1 + 0.1 * freq * 1j) + 0.1 * randn(size(freq));
r = frd(resp, freq)
r =
    continuous-time frequency response, units=rad/s
    1 input, 1 output
    101 frequencies
```

Conversion from a transfer function object:

```
freq = 0:100;
G = tf(1, [1, 2, 3, 4]);
r = frd(G, freq)
r =
    continuous-time frequency response, units=rad/s
    1 input, 1 output
    101 frequencies
```

#### See also

frd::frdata

# pid::pid

LTI PID controller constructor.

```
use lti
a = pid
a = pid(Kp, Ki, Kd, Tf)
a = pid(Kp, Ki, Kd, Tf, Ts)
a = pid(Kp, Ki, Kd, Tf, Ts, var)
a = pid(..., IFormula=f1, DFormula=f2)
```

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### **Description**

pid(Kp, Ki, Kd, Tf) creates an LTI object which represents the continuous-time PID controller  $K_p + K_i/s + K_ds/(T_fs + 1)$ , where s is the variable of the Laplace transform. Kp is the proportional gain, Ki is the integral gain, Kd is the derivative gain, and Tf is the time constant of the first-order filter of the derivative term. Missing Ki, Kd or Tf default to 0; without any input argument, Kp defaults to 1. If Tf=0 and Kd!=0, the derivative term is not filtered and the controller is not causal.

A single-input single-output (SISO) PID controller has scalar parameters. If the parameters are matrices, they must all have the same size (scalar values are replicated as required), and the resulting controller has as many inputs as parameters have columns and as many outputs as parameters have rows; mapping from each input to each output is an independent SISO PID controller.

pid(Kp, Ki, Kd, Tf, Ts) creates an LTI object which represents the discrete-time PID controller  $K_p + K_i I_i(z) + K_d / (T_f + I_d(z))$ , where  $I_i(z)$  is the integration formula used for the integral term,  $I_d(z)$  is the integration formula used for the derivative term, and z is the variable of the z transform. The formulae can be specified by named arguments IFormula and DFormula, strings with the following values:

```
Name Value 'ForwardEuler' T_s/(z-1) 'BackwardEuler' T_sz/(z-1) 'Trapezoidal' T_s/2(z+1)/(z-1)
```

The default formula for both the integral and the derivative terms is 'ForwardEuler'.

An additional argument var may be used to specify the variable of the Laplace ('s' (default) or 'p') or z transform ('z' (default) or 'q' for forward time shift, ' $z^-1$ ' or ' $q^-1$ ' for backward time shift).

For PID controllers based on the standard parameters Kp, Ti and Td, where Ki=Kp/Ti and Kd=Kp\*Td, pidstd objects should be used instead.

## **Examples**

Simple continuous-time PID controller:

```
use lti
C = pid(5,2,1)
C =
    continuous-time PID controller
    Kp + Ki/s + Kd s/(Tf s + 1)
    Kp = 5 Ki = 2 Kd = 1 Tf = 0
```

Discrete-time PD controller where the derivative term, filtered with a time constant of 20ms, is approximated with the Backward Euler formula, with a sampling period of 1ms. The controller is displayed with the backward-shift operator q^-1.

```
 C = pid(5,0,1,20e-3,1e-3,'q^-1',DFormula='BackwardEuler') \\ C = \\ discrete-time PD controller, Ts=1e-3 \\ Kp + Kd/(Tf + Id(q^-1)) \\ Id(q^-1) = Ts/(1-q^-1) (BackwardEuler) \\ Kp = 5 Kd = 1 Tf = 2e-2
```

Conversion of a first-order continuous-time transfer function with pole at 0 (integrator effect) to a continuous-time PI controller:

```
G = tf([1, 2], [1, 0])
G =
    continuous-time transfer function
    (s+2)/s
C = pid(G)
C =
    continuous-time PI controller
    Kp + Ki/s
    Kp = 1 Ki = 2
```

Conversion of a discrete-time PID controller with the Backward Euler formula for the integral term and the Trapezoidal formula for the derivative term to a transfer function, and back to a PID controller:

```
C1 = pid(5, 2, 3, 0.1, 0.01,
        IFormula='BackwardEuler', DFormula='Trapezoidal')
    discrete-time PID controller, Ts=1e-2
    Kp + Ki Ii(z) + Kd/(Tf + Id(z))
   Ii(z) = Ts z/(z-1) (BackwardEuler)
    Id(z) = Ts/2 (z+1)/(z-1) (Trapezoidal)
Kp = 5 Ki = 2 Kd = 3 Tf = 0.1
G = tf(C1)
    discrete-time transfer function, Ts=1e-2
    (3.5271z^2-7.0019z+3.475)/(0.105z^2-0.2z+9.5e-2)
C2 = pid(G, IFormula='BackwardEuler', DFormula='Trapezoidal')
 C2 =
    discrete-time PID controller, Ts=1e-2
   Kp + Ki Ii(z) + Kd/(Tf + Id(z))
   Ii(z) = Ts z/(z-1) (BackwardEuler)
    Id(z) = Ts/2 (z+1)/(z-1) (Trapezoidal)
   Kp = 5 Ki = 2 Kd = 3 Tf = 10e-2
```

#### See also

```
pidstd::pidstd, tf::tf
```

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# pidstd::pidstd

LTI standard PID controller constructor.

### **Syntax**

```
use lti
a = pidstd
a = pidstd(Kp, Ti, Td, N)
a = pidstd(Kp, Ti, Td, N, Ts)
a = pidstd(Kp, Ti, Td, N, Ts, var)
a = pidstd(..., IFormula=f1, DFormula=f2)
```

## **Description**

pidstd(Kp,Ti,Td,N) creates an LTI object which represents the standard continuous-time PID controller  $K_p(1/T_is + T_ds/(T_ds/N + 1))$ , where s is the variable of the Laplace transform. Kp is the proportional gain, Ti is the integral time, Td is the derivative time, and N is the relative frequency of the first-order filter of the derivative term. Missing Ti defaults to infinity (no integral term), missing Td to zero (no derivative term), and missing N to infinity (no filter on the derivative term, which means that the controller is noncausal if Td is nonzero).

A single-input single-output (SISO) PID controller has scalar parameters. If the parameters are matrices, they must all have the same size (scalar values are replicated as required), and the resulting controller has as many inputs as parameters have columns and as many outputs as parameters have rows; mapping from each input to each output is and independent SISO PID controller.

pid(Kp,Ti,Td,N,Ts) creates an LTI object which represents the standard discrete-time PID controller  $K_p(I_i(z)/T_i + T_d/(T_d/N + I_d(z)))$ , where  $I_i(z)$  is the integration formula used for the integral term,  $I_d(z)$  is the integration formula used for the derivative term, and z is the variable of the z transform. The formulae can be specified by named arguments IFormula and DFormula, strings with the following values:

```
Name Value 'ForwardEuler' T_s/(z-1) 'BackwardEuler' T_sz/(z-1) 'Trapezoidal' T_s/2(z+1)/(z-1)
```

The default formula for both the integral and the derivative terms is 'ForwardEuler'.

An additional argument var may be used to specify the variable of the Laplace ('s' (default) or 'p') or z transform ('z' (default) or 'q' for forward time shift, ' $z^-1$ ' or ' $q^-1$ ' for backward time shift).

For PID controllers based on the gain parameters Kp, Ki=Kp/Ti, Kd=Kp\*Td, and Tf=Td/N, pid objects should be used instead. Class

pidstd is a subclass of pid. The only differences are the arguments of their constructors and the way their objects are displayed by char, disp and mathml.

## **Examples**

Simple standard continuous-time PID controller:

```
use lti
C = pidstd(5,4,1)
C =
    continuous-time PID controller
    Kp (1 + 1/(Ti s) + Td s/(Td/N s + 1))
    Kp = 5 Ti = 4 Td = 1 N = inf
```

Conversion to a pid object:

```
C1 = pid(C)
C1 =
    continuous-time PID controller
    Kp + Ki/s + Kd s/(Tf s + 1)
    Kp = 5 Ki = 1.25 Kd = 5 Tf = 0
```

Standard discrete-time PD controller where the derivative term, filtered with a time constant 20 times smaller than the derivator time, is approximated with the Backward Euler formula, with a sampling period of 1ms. The controller is displayed with the backward-shift operator  $q^-1$ .

### See also

```
pid::pid, tf::tf
```

### SS::SS

LTI state-space constructor.

### **Syntax**

```
use lti
a = ss
a = ss(A, B, C, D)
a = ss(A, B, C, D, Ts)
a = ss(A, B, C, D, Ts, var)
a = ss(A, B, C, D, b)
a = ss(b)
```

## **Description**

ss(A,B,C,D) creates an LTI object which represents the continuoustime state-space model

```
x'(t) = A x(t) + B u(t)
y(t) = C x(t) + D u(t)
```

ss(A,B,C,D,Ts) creates an LTI object which represents the discretetime state-space model with sampling period Ts

```
x(k+1) = A x(k) + B u(k)

y(k) = C x(k) + D u(k)
```

In both cases, if D is 0, it is resized to match the size of B and C if necessary. An additional argument var may be used to specify the variable of the Laplace ('s' (default) or 'p') or z transform ('z' (default) or 'q').

ss(A,B,C,D,b), where b is an LTI object, creates a state-space model of the same kind (continuous/discrete time, sampling time and variable) as b.

ss(b) converts the LTI object b to a state-space model.

## **Examples**

```
use lti
sc = ss(-1, [1,2], [2;5], 0)
    continuous-time LTI state-space system
    A =
        - 1
    B =
         1
                2
    C =
         2
         5
    D =
         0
                0
                0
         0
```

```
sd = ss(tf(1,[1,2,3,4],0.1))
  sd =
    discrete-time LTI state-space system, Ts=0.1
             -3
      -2
                    - 4
       1
              0
                     0
              1
                     0
    B =
       1
       0
       0
    C =
              0
                     1
    D =
```

#### See also

tf::tf

### tf::tf

LTI transfer function constructor.

### **Syntax**

```
use lti
a = tf
a = tf(num, den)
a = tf(numlist, denlist)
a = tf(..., Ts)
a = tf(..., Ts, var)
a = tf(..., b)
a = tf(gain)
a = tf(b)
```

# Description

tf(num,den) creates an LTI object which represents the continuoustime transfer function specified by descending-power coefficient vectors num and den. tf(num,den,Ts) creates an LTI object which represents a discrete-time transfer function with sampling period Ts.

In both cases, num and den can be replaced with cell arrays of coefficients whose elements are the descending-power coefficient vectors. The number of rows is the number of system outputs, and the number of columns is the number of system inputs.

An additional argument var may be used to specify the variable of the Laplace ('s' (default) or 'p') or z transform ('z' (default) or 'q').

tf(...,b), where b is an LTI object, creates a transfer function of the same kind (continuous/discrete time, sampling time and variable) as b.

```
tf(b) converts the LTI object b to a transfer function.
tf(gain), where gain is a matrix, creates a matrix of gains.
```

## **Examples**

Simple continuous-time system with variable p (p is used only for display):

```
use lti
sc = tf(1,[1,2,3,4],'p')
sc =
   continuous-time transfer function
   1/(p^3+2p^2+3p+4)
```

Matrix of discrete-time transfer functions for one input and two outputs, with a sampling period of 1ms:

#### See also

```
zpk::zpk, pid::pid, pidstd::pidstd, ss::ss
```

# zpk::zpk

LTI zero-pole-gain constructor.

```
use lti
a = zpk(z, p, k)
a = zpk(Z, P, K)
a = zpk(..., Ts)
a = zpk(..., Ts, var)
a = zpk(..., b)
a = zpk(b)
```

zpk creates a zero-pole-gain LTI object. It accepts a vector of zeros, a vector of poles, and a scalar gain for a simple-input simple-output (SISO) system; or a cell array of zeros, a cell array of poles, and a real array of gains for multiple-input multiple-output (MIMO) systems. zpk(z,p,k,Ts) creates an LTI object which represents a discrete-time transfer function with sampling period Ts.

In both cases, z and p can be replaced with cell arrays of coefficients whose elements are the zeros and poles vectors, and k with a matrix of the same size. The number of rows is the number of system outputs, and the number of columns is the number of system inputs.

An additional argument var may be used to specify the variable of the Laplace ('s' (default) or 'p') or z transform ('z' (default) or 'q'). zpk(...,b), where b is an LTI object, creates a zero-pole-gain transfer function of the same kind (continuous/discrete time, sampling time and variable) as b.

zpk(b) converts the LTI object b to a zero-pole-gain transfer function.

### **Example**

```
use lti

sd = zpk(0.3, [0.8+0.5j; 0.8-0.5j], 10, 0.1)

discrete-time\ zero-pole-gain\ transfer\ function,\ Ts=0.1

10(z-0.3)/(z-(0.8+0.5j)(z-(0.8-0.5j))
```

#### See also

```
tf::tf, pid::pid, pidstd::pidstd, ss::ss
```

# Iti::append

Append the inputs and outputs of systems.

## **Syntax**

```
use lti
b = append(a1, a2, ...)
```

# **Description**

append(a1,a2) builds a system with inputs [u1;u2] and outputs [y1;y2], where u1 and u2 are the inputs of a1 and y1 and y2 their outputs, respectively. append accepts any number of input arguments.

#### See also

```
lti::connect, ss::augstate
```

# ss::augstate

Extend the output of a system with its states.

### **Syntax**

```
use lti
b = augstate(a)
```

## **Description**

augstate(a) extends the ss object a by adding its states to its outputs. The new output is [y;x], where y is the output of a and x is its states.

#### See also

```
lti::append
```

# lti::beginning

First index.

## **Syntax**

```
use lti
var(...beginning...)
```

# **Description**

In an expression used as an index between parenthesis, beginning (a) gives the first valid value for an index. It is always 1.

#### See also

```
lti::end, lti::subsasgn, lti::subsref
```

## lti::c2d

Conversion from continuous time to discrete time.

### **Syntax**

```
use lti
b = c2d(a, Ts)
b = c2d(a, Ts, method)
```

### **Description**

c2d(a,Ts) converts the continuous-time system a to a discrete-time system with sampling period Ts.

c2d(a,Ts,method) uses the specified conversion method. method is one of the methods supported by c2dm for classes ss, tf and zpk, and 'ForwardEuler', 'BackwardEuler' or 'Trapezoidal' for classes pid and pidstd.

#### See also

```
lti::d2c, c2dm
```

## lti::connect

Arbitrary feedback connections.

## **Syntax**

```
use lti
b = connect(a, links, in, out)
```

# **Description**

connect(a,links,in,out) modifies lti object a by connecting some of the outputs to some of the inputs and by keeping some of the inputs and some of the outputs. Connections are specified by the rows of matrix link. In each row, the first element is the index of the system input where the connection ends; other elements are indices to system outputs which are summed. The sign of the indices to outputs gives the sign of the unit weight in the sum. Zeros are ignored. Arguments in and out specify which input and output to keep.

#### See also

lti::feedback

# Iti::ctranspose

Conjugate transpose.

### **Syntax**

```
use lti
b = a'
b = ctranspose(a)
```

### **Description**

a' or ctranspose(a) gives the conjugate transpose of a.

The conjugate of the single-input single-output (SISO) continuous-time transfer function G(s) is defined as G(-s), and the conjugate of the SISO discrete-time transfer function G(z) is defined as G(1/z); the conjugate transpose is the conjugate of the transpose of the original system.

#### See also

```
lti::transpose, operator '
```

### ss::ctrb

Controllability matrix.

### **Syntax**

```
use lti
C = crtb(a)
```

# Description

ctrb(a) gives the controllability matrix of system a, which is full-rank if and only if a is controllable.

#### See also

```
ss::obsv
```

## lti::d2c

Conversion from discrete time to continuous time.

```
use lti
b = d2c(a)
b = d2c(a, method)
```

d2c(a) converts the discrete-time system a to a continuous-time system.

d2c(a,method) uses the specified conversion method. method is one of the methods supported by d2cm for classes ss, tf and zpk, and is ignored for class pid and pidstd.

#### See also

```
lti::c2d, d2cm
```

# lti::dcgain

Steady-state gain.

### **Syntax**

```
use lti
g = dcgain(a)
```

# **Description**

dcgain(a) gives the steady-state gain of system a.

#### See also

```
lti::norm
```

### lti::end

Last index.

### **Syntax**

```
use lti
var(...end...)
```

# Description

In an expression used as an index between parenthesis, end gives the last valid value for that index. It is size(var,1) or size(var,2).

### **Example**

Time response when the last input is a step:

```
use lti
P = ss([1,2;-3,-4],[1,0;0,1],[3,5]);
P1 = P(:, end)
  continuous-time LTI state-space system
  A =
     1
         2
    - 3
       - 4
  B =
     0
     1
  C =
     3
         5
  D =
step(P1);
```

#### See also

lti::beginning, lti::subsasgn, lti::subsref

## lti::evalfr

Frequency value.

## **Syntax**

```
use lti
y = evalfr(a, x)
```

# **Description**

evalfr(a,x) evaluates system a at complex value or values x. If x is a vector of values, results are stacked along the third dimension.

# **Example**

```
use lti
sys = [tf(1, [1,2,3]), tf(2, [1,2,3,4])];
evalfr(sys, 0:1j:3j)
  ans =
    1x2x4 array
  (:,:,1) =
    0.3333
  (:,:,2) =
```

```
0.25 - 0.25j 0.5 - 0.5j

(:,:,3) =
-5.8824e-2-0.2353j -0.4 + 0.2j

(:,:,4) =
-8.3333e-2-8.3333e-2j -5.3846e-2+6.9231e-2j
```

### See also

polyval

## frd::fcat

Frequency concatenation.

### **Syntax**

```
use lti
c = fcat(a, b)
```

## **Description**

fcat(a,b) concatenates the frequency response data of frd objects a and b along the frequency axis, and sort data by increasing frequency. The size of a and b must be the same (same numbers of inputs and outputs).

# Example

```
use lti
G = tf(1, [1, 2, 3, 4]);
a = frd(G, 0:5);
b = frd(G, 6:20);
c = fcat(a, b);
d = frd(G, 0:20); // same as c
```

### See also

frd::frd

# lti::feedback

Feedback connection.

### **Syntax**

```
use lti
c = feedback(a, b)
c = feedback(a, b, sign)
c = feedback(a, b, ina, outa)
c = feedback(a, b, ina, outa, sign)
```

## **Description**

feedback(a,b) connects all the outputs of lti object a to all its inputs via the negative feedback lti object b.

feedback(a,b,sign) applies positive feedback with weight sign; the default value of sign is -1.

feedback(a,b,ina,outa) specifies which inputs and outputs of a to use for feedback. The inputs and outputs of the result always correspond to the ones of a.

### See also

lti::connect

## frd::frdata

Get frequency response data.

## **Syntax**

```
use lti
(resp, freq) = frdata(f)
(resp, freq, Ts) = frdata(f)
```

# **Description**

frdata(f), where f is an frd object, gives the complex frequency response, the corresponding frequencies, and optionally the sampling period or the empty array [] for continuous-time systems.

#### See also

frd::frd

## frd::fselect

Frequency selection.

### **Syntax**

```
use lti
b = fselect(a, ix)
b = fselect(a, sel)
b = fselect(a, freqmin, freqmax)
```

### **Description**

fselect(a,ix) selects frequencies of frd object a whose index are in array ix. The frequencies of the result are a.freq(ix).

fselect(a,sel) selects frequencies of frd object a corresponding to true values in logical array sel. The frequencies of the result are a.freq(sel).

fselect(a,freqmin,freqmax) selects frequencies of frd object a which are greater than or equal to freqmin and less than or equal to freqmax. The frequencies of the result are a.freq(a.freq>=freqmin&a.freq<=freqmax).

#### See also

```
frd::frd, operator ()
```

# frd::interp

Frequency interpolation.

## **Syntax**

```
use lti
b = interp(a, freq)
b = interp(a, freq, method)
```

# **Description**

interp(a, freq) interpolates response data of frd object a at the frequencies in array freq. The frequencies of the result are freq. The interpolation method is linear. Interpolation for frequencies outside the frequency range of a yields nan (not a number).

interp(a,freq,method) use the specified method for interpolation. Method is one of the strings accepted by interp1 ('0' or 'nearest', '<', '>', '1' or 'linear', '3' or 'cubic', 'p' or 'pchip').

#### See also

```
frd::frd, interp1
```

## lti::inv

System inverse.

## **Syntax**

```
use lti
b = inv(a)
```

## **Description**

inv(a) gives the inverse of system a.

### See also

```
lti::mldivide, lti::mrdivide
```

## isct

Test for a continous-time LTI.

## **Syntax**

```
use lti
b = isct(a)
```

# **Description**

isct(a) is true if system a is continuous-time or static, and false otherwise.

### See also

isdt

### isdt

Test for a discrete-time LTI.

## **Syntax**

```
use lti
b = isdt(a)
```

### **Description**

isdt(a) is true if system a is discrete-time or static, and false otherwise.

#### See also

isct

# **Iti::isempty**

Test for an LTI without input/output.

## **Syntax**

```
use lti
b = isempty(a)
```

## **Description**

isempty(a) is true if system a has no input and/or no output, and false otherwise.

## See also

```
lti::size, lti::issiso
```

# lti::isproper

Test for a proper (causal) LTI.

```
use lti
b = isproper(a)
```

### **Description**

isproper(a) is true if lti object a is causal, or false otherwise. An ss object is always causal. A tf object is causal if all the transfer functions are proper, i.e. if the degrees of the denominators are at least as large as the degrees of the numerators.

### lti::issiso

Test for a single-input single-output LTI.

### **Syntax**

```
use lti
b = issiso(a)
```

## **Description**

issiso(a) is true if lti object a has one input and one output (single-input single-output system, or SISO), or false otherwise.

```
lti::size, lti::isempty
```

# tf::mathml zpk::mathml pid::mathml pidstd::mathml

Conversion to MathML.

# Syntax

```
use lti, lti_mathml
str = mathml(G)
str = mathml(G, false)
str = mathml(..., Format=f, NPrec=n)
```

# **Description**

mathml(x) converts its argument x to MathML presentation, returned as a string.

By default, the MathML top-level element is <math>. If the result is to be used as a MathML subelement of a larger equation, a last input argument equal to the logical value false can be specified to suppress <math>.

By default, mathml converts numbers like format '%g' of sprintf. Named arguments can override them: format is a single letter format recognized by sprintf and NPrec is the precision (number of decimals).

### **Example**

```
use lti, lti_mathml
G = zpk(-1, [1, 2+j, 2-j], 2);
m = mathml(G);
math(0, 0, m);
```

#### See also

mathml.sprintf

## lti::minreal

Minimum realization.

### **Syntax**

```
use lti
b = minreal(a)
b = minreal(a, tol)
```

## **Description**

minreal(a) modifies lti object a in order to remove states which are not controllable and/or not observable. For tf objects, identical zeros and poles are canceled out.

minreal(a,tol) uses tolerance tol to decide whether to discard a state or a pair of pole/zero.

## Iti::minus

System difference.

## **Syntax**

```
use lti
c = a - b
c = minus(a, b)
```

# **Description**

a-b computes the system whose inputs are fed to both a and b and whose outputs are the difference between outputs of a and b. If a and b are transfer functions or matrices of transfer functions, this is equivalent to a difference of matrices.

#### See also

```
lti::parallel, lti::plus, lti::uminus
```

## lti::mldivide

System left division.

## **Syntax**

```
use lti
c = a \ b
c = mldivide(a, b)
```

## **Description**

a/b is equivalent to inv(a)\*b.

#### See also

```
lti::mrdivide, lti::times, lti::inv
```

# Iti::mrdivide

System right division.

## **Syntax**

```
use lti
c = a / b
c = mrdivide(a, b)
```

# **Description**

a/b is equivalent to a\*inv(b).

### See also

```
lti::mldivide, lti::times, lti::inv
```

## lti::mtimes

System product.

### **Syntax**

```
use lti
c = a * b
c = mtimes(a, b)
```

### **Description**

a\*b connects the outputs of lti object b to the inputs of lti object a. If a and b are transfer functions or matrices of transfer functions, this is equivalent to a product of matrices.

#### See also

```
lti::series
```

### lti::norm

H2 norm.

## **Syntax**

```
use lti
h2 = norm(a)
```

## **Description**

norm(a) gives the H2 norm of the system a.

### See also

```
lti::dcgain
```

## ss::obsv

Observability matrix.

## **Syntax**

```
use lti
0 = obsv(a)
```

## **Description**

obsv(a) gives the observability matrix of system a, which is full-rank if and only if a is observable.

#### See also

```
ss::ctrb
```

# lti::parallel

Parallel connection.

## **Syntax**

```
use lti
c = parallel(a, b)
c = parallel(a, b, ina, inb, outa, outb)
```

## **Description**

parallel(a,b) connects lti objects a and b in such a way that the inputs of the result is applied to both a and b, and the outputs of the result is their sum.

parallel(a,b,ina,inb,outa,outb) specifies which inputs are shared between a and b, and which outputs are summed. The inputs of the result are partitioned as [ua,uab,ub] and the outputs as [ya,yab,yb]. Inputs uab are fed to inputs ina of a and inb of b; inputs ua are fed to the remaining inputs of a, and ub to the remaining inputs of b. Similarly, outputs yab are the sum of outputs outa of a and outputs outb of b, and ya and yb are the remaining outputs of a and b, respectively.

#### See also

```
lti::series
```

# lti::piddata

Get PID parameters.

```
use lti
(Kp, Ki, Kd, Tf) = piddata(a)
(Kp, Ki, Kd, Tf, Ts) = piddata(a)
```

piddata(a), where a is any kind of LTI object which has the structure of a PID controller except for frd, gives the PID parameters Kp, Ki, Kd and Tf, and optionally the sampling period or the empty array [] for continuous-time systems. The parameters are given as matrices; the rows correspond to the outputs, and their columns to the inputs.

#### See also

```
pid::pid, lti::pidstddata, lti::tfdata
```

# lti::pidstddata

Get standard PID parameters.

## **Syntax**

```
use lti
(Kp, Ti, Td, N) = pidstddata(a)
(Kp, Ti, Td, N, Ts) = pidstddata(a)
```

## Description

pidstddata(a), where a is any kind of LTI object which has the structure of a PID controller except for frd, gives the standard PID parameters Kp, Ti, Td and N, and optionally the sampling period or the empty array [] for continuous-time systems. The parameters are given as matrices; the rows correspond to the outputs, and their columns to the inputs.

#### See also

```
pidstd::pidstd, lti::piddata, lti::tfdata
```

# lti::plus

System sum.

```
use lti
c = a + b
c = plus(a, b)
```

### **Description**

a+b computes the system whose inputs are fed to both a and b and whose outputs are the sum of the outputs of a and b. If a and b are transfer functions or matrices of transfer functions, this is equivalent to a sum of matrices.

#### See also

```
lti::parallel, lti::minus
```

# lti::repmat

Replicate a system.

### **Syntax**

```
use lti
b = repmat(a, n)
b = repmat(a, [m,n])
b = repmat(a, m, n)
```

## **Description**

repmat(a,m,n), when a is an lti object and m and n are positive integers, creates a new system of the same class with m times as many outputs and n times as many inputs. If a is a matrix of transfer functions, it is replicated m times vertically and n horizontally, as if a were a numeric matrix. If a is a state-space system, matrices B, C, and D are replicated to obtain the same effect.

repmat(a,[m,n]) gives the same result as repmat(a,m,n); repmat(a,n) gives the same result as repmat(a,n,n).

#### See also

```
lti::append
```

### lti::series

Series connection.

```
use lti
c = series(a, b)
c = series(a, b, outa, inb)
```

series(a,b) connects the outputs of lti object a to the inputs of lti object b.

series(a,b,outa,inb) connects outputs outa of a to inputs inb of b. Unconnected outputs of a and inputs of b are discarded.

#### See also

```
lti::mtimes, lti::parallel
```

### lti::size

Number of outputs and inputs.

### **Syntax**

```
use lti
s = size(a)
(nout, nin) = size(a)
n = size(a, dim)
```

## **Description**

With one output argument, size(a) gives the row vector [nout,nin], where nout is the number of outputs of system a and nin its number of inputs. With two output arguments, size(a) returns these results separately as scalars.

size(a,1) gives only the number of outputs, and size(a,2) only the number of inputs.

### See also

```
lti::isempty, lti::issiso
```

## lti::ssdata

Get state-space matrices.

```
use lti
(A, B, C, D) = ssdata(a)
(A, B, C, D, Ts) = ssdata(a)
```

### **Description**

ssdata(a), where a is any kind of LTI object except for frd, gives the four matrices of the state-space model, and optionally the sampling period or the empty array [] for continuous-time systems.

#### See also

```
ss::ss.lti::tfdata
```

# lti::subsasgn

Assignment to a part of an LTI system.

## **Syntax**

```
use lti
var(i,j) = a
var(ix) = a
var(select) = a
var.field = value
a = subsasgn(a, s, b)
```

## **Description**

The method subsasgn(a) permits the use of all kinds of assignments to a part of an LTI system. If the variable is a matrix of transfer functions, subsasgn produces the expected result, converting the righthand side of the assignment to a matrix of transfer function if required. If the variable is a state-space model, the result is equivalent; the result remains a state-space model. For state-space models, changing all the inputs or all the outputs with the syntax var(expr,:)=sys or var(:,expr)=sys is much more efficient than specifying both subscripts or a single index.

The syntax for field assignment, var.field=value, is defined for the following fields: for state-space models, A, B, C, and D (matrices of the state-space model); for transfer functions, num and den (cell arrays of coefficients); for zero-pole-gain transfer functions, z and p (cell arrays of zero or pole vectors), and k (gain matrix); for PID controllers, Kp, Ki, Kd, Tf, Ti and Td (controller parameter matrices); for all LTI objects, var (string) and Ts (scalar, or empty array for continuous-time systems). Field assignment must preserve the size of matrices and arrays.

The syntax with braces (var{i}=value) is not supported.

#### See also

```
lti::subsref, operator (), subsasgn
```

### lti::subsref

Extraction of a part of an LTI system.

### **Syntax**

```
use lti
var(i,j)
var(ix)
var(select)
var.field
b = subsref(a, s)
```

### **Description**

The method subsref(a) permits the use of all kinds of extraction of a part of an LTI system. If the variable is a matrix of transfer functions, subsref produces the expected result. If the variable is a state-space model, the result is equivalent; the result remains a state-space model, with the same state vector (the same matrix A) as the original system. For state-space models, extracting all the inputs or all the outputs with the syntax var(expr,:) or var(:,expr) is much more efficient than specifying both subscripts or a single index.

If the variable is an frd object, var('freq',i) produces a new frd object where the frequency vector is var.frequency(i) amd the response array contains the corresponding reponse. i can be a scalar index, a vector of indices or a logical array with the same size as var.frequency.

The syntax for field access, var.field, is defined for the following fields: for state-space models, A, B, C, and D (matrices of the state-space model); for transfer functions, num and den (cell arrays of coefficients); for zero-pole-gain transfer functions, z and p (cell arrays of zero or pole vectors), and k (gain matrix); for PID controllers, Kp, Ki, Kd, Tf, Ti and Td (controller parameter matrices); for all LTI objects, var (string) and Ts (scalar, or empty array for continuous-time systems).

The syntax with braces (var{i}) is not supported.

#### See also

```
lti::subsasgn, operator (), subsasgn
```

### lti::tfdata

Get transfer functions.

### **Syntax**

```
use lti
(num, den) = tfdata(a)
(num, den, Ts) = tfdata(a)
```

### **Description**

tfdata(a), where a is any kind of LTI object except for frd, gives the numerator and denominator of the transfer function model, and optionally the sampling period or the empty array [] for continuous-time systems. The numerators and denominators are given as a cell array of power-descending coefficient vectors; the rows of the cell arrays correspond to the outputs, and their columns to the inputs.

#### See also

```
tf::tf, lti::zpkdata, lti::ssdata
```

# Iti::transpose

Transpose.

## **Syntax**

```
use lti
b = a.'
b = transpose(a)
```

# **Description**

```
a.' or transpose(a) gives the transpose of a, i.e. a.'(i,j)=a(j,i).
```

#### See also

```
lti::ctranspose, operator .'
```

## Iti::uminus

Negative.

## **Syntax**

```
use lti
b = -a
b = uminus(a)
```

## **Description**

-a multiplies all the outputs (or all the inputs) of system a by -1. If a is a transfer functions or a matrix of transfer functions, this is equivalent to the unary minus.

### See also

```
lti::minus, lti::uplus
```

# Iti::uplus

Positive.

## **Syntax**

```
use lti
b = +a
b = uplus(a)
```

# **Description**

```
+a gives a.
```

### See also

```
lti::uminus, lti::plus
```

# lti::zpkdata

Get zeros, poles and gains.

```
use lti
(z, p, k) = zpkdata(a)
(z, p, k, Ts) = zpkdata(a)
```

zpkdata(a), where a is any kind of LTI object except for frd, gives the zeros, poles and gains of the transfer function model, and optionally the sampling period or the empty array [] for continuous-time systems. The zeros and poles are given as a cell array of vectors; the rows of the cell arrays correspond to the outputs, and their columns to the inputs.

#### See also

zpk::zpk, lti::tfdata

# 7.9 Iti (graphics)

In addition to the class definitions and the computational methods, library lti includes methods which provide for lti objects the same functionality as the native graphical functions of Sysquake for dynamical systems, such as bodemag for the magnitude of the Bode diagram or step for the step response. The system is provided as a single lti object instead of separate vectors for the numerator and denominator or four matrices for state-space models. For discrete-time systems, the sampling time is also obtained from the object, and the method name is the same as its continuous-time equivalent, without an initial d (e.g. step(G) is the discrete-time step response of G if G is a discrete-time tf, zpk or ss object).

The method definitions are stored in a separate file which is referenced in lti with includeifexists; this means that only lti must be loaded, with

use lti

# **Functions**

# lti::bodemag

Magnitude of the Bode plot.

```
use lti
bodemag(a)
bodemag(a, style, id)
(mag, w) = bodemag(a)
```

bodemag(a) plots the magnitude of the Bode diagram of system a, which can be any lti object with a single input (size(a,2) must be 1), continuous-time or discrete-time.

The optional arguments style and id have their usual meaning.

With output arguments, bodemag gives the magnitude and the frequency as column vectors. No display is produced.

## **Examples**

```
Green plot for |1/(s^3 + 2s^2 + 3s + 4)| with s = j\omega (see Fig. 6.9):

G = tf(1, [1, 2, 3, 4]); bodemag(G, 'g');
```

The same plot, between  $\omega=0$  and  $\omega=10$ , with a named argument for the color:

```
scale([0,10]);
bodemag(G, Color='green');
```

Frequency response of the discrete-time system 1/(z-0.9)(z-0.7-0.6j)(z-0.7+0.6j) with unit sampling period:

```
H = zpk([], [0.9,0.7+0.6j,0.7-0.6j], 1, 1);
bodemag(H);
```

#### See also

```
lti::bodephase, lti::nichols, lti::nyquist, plotset, bodemag
```

# lti::bodephase

Phase of the Bode plot.

```
use lti
bodephase(a)
bodephase(a, style, id)
(phase, w) = bodephase(a)
```

bodephase(a) plots the phase of the Bode diagram of system a, which can be any lti object with a single input (size(a,2) must be 1), continuous-time or discrete-time.

The optional arguments style and id have their usual meaning.

With output arguments, bodephase gives the phase and the frequency as column vectors. No display is produced.

#### See also

```
lti::bodemag, lti::nichols, lti::nyquist, plotset, bodephase
```

# Iti::impulse

Impulse response.

## Syntax

```
use lti
impulse(a)
impulse(a, style, id)
(y, t) = impulse(a)
```

# **Description**

impulse(a) plots the impulse response of system a, which can be any lti object with a single input (size(a,2) must be 1), continuous-time or discrete-time.

The optional arguments style and id have their usual meaning.

With output arguments, impulse gives the output and the time as column vectors. No display is produced.

# **Example**

Impulse response of the first order transfer function 1/(s/2 + 1):

```
G = tf(1, [1/2, 1]);
impulse(G);
```

#### See also

```
lti::step, lti::lsim, ss::initial, plotset, impulse
```

### ss::initial

Time response with initial conditions.

## **Syntax**

```
use lti
initial(a, x0)
initial(a, x0, style, id)
(y, t) = initial(a, x0)
```

## **Description**

initial(a,x0) plots the time response of state-space system a with initial state x0 and null input. System a can be continuous-time or discrete-time.

The optional arguments style and id have their usual meaning.

With output arguments, initial gives the output and the time as column vectors. No display is produced.

### **Example**

Response of a continuous-time system whose initial state is [5;3]:

```
a = ss([-0.3,0.1;-0.8,-0.4], [2;3], [1,3;2,1], [2;1]);
initial(a, [5;3])
```

#### See also

```
lti::impulse, lti::step, lti::lsim, plotset, initial
```

### lti::lsim

Time response.

```
use lti
lsim(a, u, t)
lsim(a, u, t, style, id)
(y, t) = lsim(a, u, t)
```

lsim(a,u,t) plots the time response of system a. For continuous-time systems, the input is piece-wise linear; it is defined by points in real vectors t and u, which must have the same length. Input before t(1) and after t(end) is 0. For discrete-time systems, u is sampled at the rate given by the system, and t is ignored or can be omitted.

The optional arguments style and id have their usual meaning.

With output arguments, lsim gives the output and the time as column vectors. No display is produced.

### **Example**

Response of continuous-time system given by its transfer function with an input defined by linear segments, displayed as a solid blue line:

```
G = tf(1, [1, 2, 3, 4]);
t = [0, 10, 20, 30, 50];
u = [1, 1, 0, 1, 1];
lsim(G, u, t, Color = 'blue');
```

### See also

```
lti::impulse, lti::step, ss::initial, plotset, lsim
```

### Iti::nichols

Nichols plot.

# **Syntax**

```
use lti
nichols(a, ...)
(mag, phase, w) = nichols(a, ...)
```

# Description

nichols(a) plots the Nichols diagram of system a, which can be any lti object with a single input (size(a,2) must be 1), continuous-time or discrete-time.

The optional arguments style and id have their usual meaning.

With output arguments, nichols gives the magnitude, the phase and the corresponding frequency as column vectors. No display is produced.

#### See also

```
lti::nyquist, lti::bodemag, lti::bodephase, plotset, nichols
```

# lti::nyquist

Nyquist plot.

## **Syntax**

```
use lti
nyquist(a, ...)
(re, im, w) = nyquist(a, ...)
```

## **Description**

nyquist(a) plots the Nyquist diagram of system a, which can be any lti object with a single input (size(a,2) must be 1), continuous-time or discrete-time.

The optional arguments style and id have their usual meaning.

With output arguments, nyquist gives the real part, the imaginary part and the corresponding frequency as column vectors. No display is produced.

#### See also

```
lti::nichols, lti::bodemag, lti::bodephase, plotset, nyquist
```

# lti::pzmap

Pole/zero map.

# **Syntax**

```
use lti
pzmap(a)
pzmap(a, style)
```

# Description

pzmap(a) plots the poles and the zeros of system a in the complex plane. Poles are represented with crosses and zeros with circles. The system must be SISO (single-input, single-output).

With a second input argument, pzmap(a, style) uses the specified style for the poles and zeros. Typically, style is a structure array of

two elements: the first element contains style options for the poles, and the second element, for the zeros. An empty structure (0 element) stands for the default style, and a simple structure uses the same style for the poles and the zeros.

### **Examples**

Pole/zero map of a transfer function:

```
use lti
G = tf([2, 3, 4], [1, 2, 3, 4]);
pzmap(G);
```

Pole/zero map with the same scale along x and y axes, a grid showing relative damping and natural frequencies, and explicit style:

```
use lti
G = tf([2, 3, 4], [1, 2, 3, 4]);
scale equal;
sgrid;
plotoption fullgrid;
style = {
   Marker='x', MarkerEdgeColor='red';
   Marker='o', MarkerEdgeColor='navy', MarkerFaceColor='yellow'
}
pzmap(G, style);
```

### See also

lti::rlocus, plotset, plotroots

### lti::rlocus

Root locus.

## **Syntax**

```
use lti
rlocus(a)
rlocus(a, style, id)
```

# **Description**

rlocus(a) plots the root locus of system a, i.e. the locus of the poles of the system obtained by adding a feedback loop with a positive real gain. Only the root locus itself is displayed, as a solid line by default.

Open-loop poles and zeros (the extremities of the root locus), which are typically displayed with special markers, can be added with pzmap. The optional arguments style and id have their usual meaning.

### **Example**

Root locus of a transfer function with open-loop poles and zeros displayed with pzmap. The scale is the same along x and y axes thanks to a call to scale, and a grid shows relative damping and natural frequencies.

```
use lti
G = tf([2, 3, 1], [1, 2, 3, 4]);
scale equal;
sgrid;
plotoption fullgrid;
rlocus(G);
pzmap(G);
```

#### See also

```
lti::pzmap, plotset, rlocus
```

# lti::step

Step response.

## **Syntax**

```
use lti
step(a)
step(a, style, id)
(y, t) = step(a)
```

## **Description**

step(a) plots the step response of system a, which can be any lti object with a single input (size(a,2) must be 1), continuous-time or discrete-time.

The optional arguments style and id have their usual meaning. With output arguments, step gives the output and the time as column vectors. No display is produced.

#### See also

```
lti::impulse, lti::lsim, ss::initial, plotset, step
```

# 7.10 sigenc

sigenc is a library which adds to LME functions for encoding and decoding scalar signals. It implements quantization, DPCM (differential pulse code modulation), and companders used in telephony.

The following statement makes available functions defined in sigenc:

use sigenc

## **Functions**

## alawcompress

A-law compressor.

## **Syntax**

```
use sigenc
output = alawcompress(input)
output = alawcompress(input, a)
```

## **Description**

alawcompress(input,a) compresses signal input with A-law method using parameter a. The signal is assumed to be in [-1,1]; values outside this range are clipped. input can be a real array of any size and dimension. The default value of a is 87.6.

The compressor and its inverse, the expander, are static, nonlinear filters used to improve the signal-noise ratio of quantized signals. The compressor should be used before quantization (or on a signal represented with a higher precision).

#### See also

alawexpand, ulawcompress

# alawexpand

A-law expander.

```
use sigenc
output = alawexpand(input)
output = alawexpand(input, a)
```

alawexpand(input,a) expands signal input with A-law method using parameter a. input can be a real array of any size and dimension. The default value of a is 87.6.

### See also

alawcompress, ulawexpand

# dpcmdeco

Differential pulse code modulation decoding.

### **Syntax**

```
use sigenc
output = dpcmdeco(i, codebook, predictor)
```

## **Description**

dpcmdeco(i,codebook,predictor) reconstructs a signal encoded with differential pulse code modulation. It performs the opposite of dpcmenco.

#### See also

dpcmenco, dpcmopt

# dpcmenco

Differential pulse code modulation encoding.

# Syntax

```
use sigenc
i = dpcmenco(input, codebook, partition, predictor)
```

# Description

dpcmenco(input,codebook,partition,predictor) quantizes the signal in vector input with differential pulse code modulation. It predicts the future response with the finite-impulse response filter given by polynomial predictor, and it quantizes the residual error

with codebook and partition like quantiz. The output i is an array of codes with the same size and dimension as input.

The prediction  $y^*(k)$  for sample k s

$$y^*(k) = \sum_{i=1}^{\text{degpredictor}} \text{predictor}_i \cdot y_q(k-i)$$

where  $y_q(k)$  is the quantized (reconstructed) signal. The predictor must be strictly causal: predictor(0) must be zero. To encode the difference between in(k) and yq(k-1), predictor=[0,1]. Note that there is no drift between the reconstructed signal and the input  $^1$ , contrary to the case where the input is differentiated, quantized, and integrated.

### **Example**

```
use sigenc
t = 0:0.1:10;
x = sin(t);
codebook = -.1:.01:.1;
partition = -.0:.01:.09;
predictor = [0, 1];
i = dpcmenco(x, codebook, partition, predictor);
y = dpcmdeco(i, codebook, predictor);
```

#### See also

quantiz, dpcmdeco, dpcmopt

## dpcmopt

Differential pulse code modulation decoding.

## Syntax

```
use sigenc
(predictor, codebook, partition) = dpcmopt(in, order, n)
(predictor, codebook, partition) = dpcmopt(in, order, codebook0)
(predictor, codebook, partition) = dpcmopt(in, predictor, ...)
(predictor, codebook, partition) = dpcmopt(..., tol)
predictor = dpcmopt(in, order)
```

<sup>&</sup>lt;sup>1</sup>Actually, there may be a drift if the arithmetic units used for encoding and decoding do not produce exactly the same results.

### **Description**

dpcmopt(in,order,n) gives the optimal predictor of order order, codebook of size n and partition to encode the signal in vector in with differential pulse code modulation. The result can be used with dpcmenco to encode signals with similar properties. If the second input argument is a vector, it is used as the predictor and not optimized further; its first element must be zero. If the third input argument is a vector, it is used as an initial guess for the codebook, which has the same length. An optional fourth input argument provides the tolerance (the default is 1e-7).

If only the predictor is required, only the input and the predictor order must be supplied as input arguments.

#### See also

dpcmenco, dpcmdeco, lloyds

## lloyds

Optimal quantization.

## Syntax

```
use sigenc
(partition, codebook) = lloyds(input, n)
(partition, codebook) = lloyds(input, codebook0)
(partition, codebook) = lloyds(..., tol)
```

## **Description**

lloyds(input,n) computes the optimal partition and codebook for quantizing signal input with n codes, using the Lloyds algorithm.

If the second input argument is a vector, lloyds(input,codebook0) uses codebook0 as an initial guess for the codebook. The result has the same length.

A third argument can be used to specify the tolerance used as the stopping criterion of the optimization loop. The default is 1e-7.

## Example

We start from a suboptimal partition and compute the distortion:

```
use sigenc
partition = [-1, 0, 1];
codebook = [-2, -0.5, 0.5, 2];
```

```
in = -5:0.6:3;
(i, out, dist) = quantiz(in, partition, codebook);
dist
  2.1421
```

The partition is optimized with lloyds, and the same signal is quantized again. The distortion is reduced.

#### See also

quantiz, dpcmopt

## quantiz

Table-based signal quantization.

## Syntax

```
use sigenc
i = quantiz(input, partition)
(i, output, distortion) = quantiz(input, partition, codebook)
```

## **Description**

quantiz(input,partition) quantizes signal input using partition as boundaries between different ranges. Range from  $-\infty$  to partition(1) corresponds to code 0, range from partition(1) to partition(2) corresponds to code 1, and so on. input may be a real array of any size and dimension; partition must be a sorted vector. The output i is an array of codes with the same size and dimension as input.

quantiz(input,partition,codebook) uses codebook as a lookup table to convert back from codes to signal. It should be a vector with one more element than partition. With a second output argument, quantiz gives codebook(i).

With a third output argument, quantiz computes the distortion between input and codebook(i), i.e. the mean of the squared error.

### **Example**

```
use sigenc
partition = [-1, 0, 1];
codebook = [-2, -0.5, 0.5, 2];
in = randn(1, 5)
  in =
    0.1799 -9.7676e-2 -1.1431
                                -0.4986
                                             1.0445
(i, out, dist) = quantiz(in, partition, codebook)
    2
         1
            0
                 1
                        2
  out =
    0.5 - 0.5 - 2 - 0.5 0.5
  dist =
    0.259
```

#### See also

lloyds, dpcmenco

## ulawcompress

mu-law compressor.

## **Syntax**

```
use sigenc
output = ulawcompress(input)
output = ulawcompress(input, mu)
```

## Description

ulawcompress(input,mu) compresses signal input with mu-law method using parameter mu. input can be a real array of any size and dimension. The default value of mu is 255.

The compressor and its inverse, the expander, are static, nonlinear filters used to improve the signal-noise ratio of quantized signals. The compressor should be used before quantization (or on a signal represented with a higher precision).

#### See also

ulawexpand, alawcompress

## ulawexpand

mu-law expander.

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### **Syntax**

```
use sigenc
output = ulawexpand(input)
output = ulawexpand(input, mu)
```

### **Description**

ulawexpand(input,mu) expands signal input with mu-law method using parameter a. input can be a real array of any size and dimension. The default value of mu is 255.

#### See also

ulawcompress, alawexpand

## 7.11 wav

wav is a library which adds to LME functions for encoding and decoding WAV files. WAV files contain digital sound. The wav library supports uncompressed, 8-bit and 16-bit, monophonic and polyphonic WAV files. It can also encode and decode WAV data in memory without files.

The following statement makes available functions defined in wav:

use wav

## **Functions**

### wavread

WAV decoding.

## **Syntax**

```
use wav
(samples, samplerate, nbits) = wavread(filename)
(samples, samplerate, nbits) = wavread(filename, n)
(samples, samplerate, nbits) = wavread(filename, [n1,n2])
(samples, samplerate, nbits) = wavread(data, ...)
```

### **Description**

wavread(filename) reads the WAV file filename. The result is a 2-d array, where each row corresponds to a sample and each column to a channel. Its class is the same as the native type of the WAV file, i.e. int8 or int16.

wavread(filename,n), where n is a scalar integer, reads the first n samples of the file. wavread(filename,[n1,n2]), where the second input argument is a vector of two integers, reads samples from n1 to n2 (the first sample corresponds to 1).

Instead of a file name string, the first input argument can be a vector of bytes, of class int8 or uint8, which represents directly the contents of the WAV file.

In addition to the samples, wavread can return the sample rate in Hz (such as 8000 for phone-quality speech or 44100 for CD-quality music), and the number of bits per sample and channel.

### See also

wavwrite

### wavwrite

WAV encoding.

## **Syntax**

```
use wav
wavwrite(samples, samplerate, nbits, filename)
data = wavwrite(samples, samplerate, nbits)
data = wavwrite(samples, samplerate)
```

## **Description**

wavwrite(samples, samplerate, nbits, filename) writes a WAV file filename with samples in array samples, sample rate samplerate (in Hz), and nbits bits per sample and channel. Rows of samples corresponds to samples and columns to channels. nbits can be 8 or 16.

With 2 or 3 input arguments, wavwrite returns the contents of the WAV file as a vector of class uint8. The default word size is 16 bits per sample and channel.

## **Example**

```
use wav
sr = 44100;
```

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```
t = (0:sr)' / sr;
s = sin(2 * pi * 740 * t);
wavwrite(map2int(s, -1, 1, 'int16'), sr, 16, 'beep.wav');
```

#### See also

wavread

## 7.12 date

date is a library which adds to LME functions to convert date and time between numbers and strings.

The following statement makes available functions defined in date:

use date

## **Functions**

#### datestr

Date to string conversion.

## **Syntax**

```
use date
str = datestr(datetime)
str = datestr(date, format)
```

## **Description**

datestr(datetime) converts the date and time to a string. The input argument can be a vector of 3 to 6 elements for the year, month, day, hour, minute, and second; a julian date as a scalar number; or a string, which is converted by datevec. The result has the following format:

```
jj-mmm-yyyy HH:MM:SS
```

where jj is the two-digit day, mmm the beginning of the month name, yyyy the four-digit year, HH the two-digit hour, MM the two-digit minute, and SS the two-digit second.

The format can be specified with a second input argument. When datestr scans the format string, it replaces the following sequences of characters and keeps the other ones unchanged:

Sequence	Replaced with
dd	day (2 digits)
ddd	day of week (3 char)
HH	hour (2 digits, 01-12 or 00-23)
MM	minute (2 digits)
mm	month (2 digits)
mmm	month (3 char)
PM	AM or PM
QQ	quarter (Q1 to Q4)
SS	second (2 digits)
SSS	fraction of second (1-12 digits)
уу	year (2 digits)
уууу	year (4 digits)

If the sequence PM is found, the hour is between 1 and 12; otherwise, between 0 and 23. Second fraction has as many digits as there are 's' characters in the format string.

### **Examples**

```
use date
datestr(clock)
   18-Apr-2005 16:21:55
datestr(clock, 'ddd mm/dd/yyyy HH:MM PM')
   Mon 04/18/2005 04:23 PM
datestr(clock, 'yyyy-mm-ddTHH:MM:SS,sss')
   2008-08-23T02:41:37,515
```

#### See also

datevec, julian2cal, clock

#### datevec

String to date and time conversion.

### **Syntax**

```
use date
datetime = datevec(str)
```

## **Description**

datevec(str) converts the string str representing the date and/or the time to a row vector of 6 elements for the year, month, day, hour, minute, and second. The following formats are recognized:

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Example	Value
20050418T162603	ISO 8601 date and time
2005-04-18	year, month and day
2005-Apr-18	year, month and day
18-Apr-2005	day, month and year
04/18/2005	month, day and year
04/18/00	month, day and year
18.04.2005	day, month and year
18.04.05	day, month and year
16:26:03	hour, minute and second
16:26	hour and minute
PM	afternoon

Unrecognized characters are ignored. If the year is given as two digits, it is assumed to be between 1951 and 2050.

### **Examples**

#### See also

datestr

## weekday

Week day of a given date.

## **Syntax**

```
use date
(num, str) = weekday(year, month, day)
(num, str) = weekday(datetime)
(num, str) = weekday(jd)
```

## **Description**

weekday finds the week day of the date given as input. The date can be given with three input arguments for the year, the month and the day, or with one input argument for the date or date and time vector, or julian date.

The first output argument is the number of the day, from 1 for Sunday to 7 for Saturday; and the second output argument is its name as a string of 3 characters, such as 'Mon' for Monday.

### **Example**

Day of week of today:

```
use date
(num, str) = weekday(clock)
num =
   2
str =
   Mon
```

#### See also

cal2julian

## 7.13 constants

constants is a library which defines physical constants in SI units (meter, kilogram, second, ampere, kelvin, candela, mole).

The following statement makes available constants defined in constants:

```
use constants;
```

The following constants are defined:

Name	Value	Unit
avogadro_number	6.0221367e23	1/mole
boltzmann_constant	1.380658e-23	J/K
earth_mass	5.97370e24	kg
$earth_{-}radius$	6.378140e6	m
electron_charge	1.60217733e-19	С
electron_mass	9.1093897e-31	kg
faraday_constant	9.6485309e4	C/mole
gravitational_constant	6.672659e-11	N m^2/kg^2
${\sf gravity\_acceleration}$	9.80655	m/s^2
hubble_constant	3.2e-18	1/s
ice_point	273.15	K
induction_constant	1.256e-6	V s/A m
molar_gaz_constant	8.314510	J/K mole
molar_volume	22.41410e-3	m^3/mole
${\sf muon\_mass}$	1.8835327e-28	kg
${\sf neutron\_mass}$	1.6749286e-27	kg
${\sf plank\_constant}$	6.6260755e-34	Js
plank_constant_reduced	1.0545727e-34	Js
plank_mass	2.17671e-8	kg
proton_mass	1.6726231e-27	kg
solar_radius	6.9599e8	m
speed_of_light	299792458	m/s
speed_of_sound	340.29205	m/s
stefan_boltzmann_constant		W/m^2 K^-4
$vacuum_{-}permittivity$	8.854187817e-12	A s/V m

## 7.14 colormaps

colormaps is a library containing functions related to color maps. Color maps are tables of colors which can be used with the colormap function; they are used by functions such as image and surf to map values to colors.

All functions accept at least the number of colors n as input argument, and produce an n-by-3 real double array which can be used directly as the argument of colormap. The default value of n is 256.

colormaps defines the following functions:

Function	Description
black2orangecm	color shades from black to orange
black2red2whitecm	color shades from black to red and white
blue2greencm	color shades from blue to green
blue2yellow2redcm	color shades from blue to yellow and red
cyan2magentacm	color shades from cyan to magenta
graycm	gray shades from black to white
green2yellowcm	color shades from green to yellow
huecm	color shades from red to red through green and blue
interprgbcm	colormap created with linear interpolation
magenta2yellowcm	color shades from magenta to yellow
red2yellowcm	color shades from red to yellow
sepiacm	sepia shades
whitecm	plain white

The following statement makes available functions defined in colormaps:

```
use colormaps
```

Functions are typically used directly as the argument of colormap:

```
colormap(blue2yellow2red);
```

## **Functions**

## black2orangecm

Colormap with shades from black to orange.

### **Syntax**

```
use colormaps
cm = black2orangecm
cm = black2orangecm(n)
```

## **Description**

black2orangecm(n) creates a color map with n entries corresponding to color shades from black to orange. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

### See also

colormap, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

### black2red2whitecm

Colormap with shades from black to red and white.

### **Syntax**

```
use colormaps
cm = black2red2whitecm
cm = black2red2whitecm(n)
```

### **Description**

black2red2whitecm(n) creates a color map with n entries corresponding to color shades from black to red and white. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

## blue2greencm

Colormap with shades from blue to green.

## Syntax

```
use colormaps
cm = blue2greencm
cm = blue2greencm(n)
```

### **Description**

blue2greencm(n) creates a color map with n entries corresponding to color shades from blue to green. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

## blue2yellow2redcm

Colormap with shades from blue to yellow and red.

### **Syntax**

```
use colormaps
cm = blue2yellow2redcm
cm = blue2yellow2redcm(n)
```

## **Description**

blue2yellow2redcm(n) creates a color map with n entries corresponding to color shades from blue to yellow and red. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, cyan2magentacm, graycm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

## cyan2magentacm

Colormap with shades from cyan to magenta.

### **Syntax**

```
use colormaps
cm = cyan2magentacm
cm = cyan2magentacm(n)
```

### **Description**

cyan2magentacm(n) creates a color map with n entries corresponding to color shades from cyan to magenta. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, graycm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

### graycm

Colormap with shades of gray.

## **Syntax**

```
use colormaps
cm = graycm
cm = graycm(n)
```

## **Description**

graycm(n) creates a color map with n entries corresponding to gray shades from black to white. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

## green2yellowcm

Colormap with shades from green to yellow.

### **Syntax**

```
use colormaps
cm = green2yellowcm
cm = green2yellowcm(n)
```

### **Description**

green2yellowcm(n) creates a color map with n entries corresponding to color shades from green to yellow. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

#### huecm

Colormap with hue from red to red through green and blue.

### **Syntax**

```
use colormaps
cm = huecm
cm = huecm(n)
```

## Description

huecm(n) creates a color map with n entries corresponding to color shades with hue varying linearly from red back to red through green and blue. In HSV (hue-saturation-value) space, saturation and value are 1 (maximum). The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

## interprgbcm

Colormap with entries obtained by linear interpolation.

## **Syntax**

```
use colormaps
cm = interprgbcm(i, r, g, b)
cm = interprgbcm(i, r, q, b, n)
```

### **Description**

interprgbcm(i,r,b,g,n) creates a color map with n entries. Color shades are interpolated between colors defined in RGB color space by corresponding elements of r, g and b, defined for input in i. These four arguments must be vectors of the same length larger or equal to 2 with elements between 0 and 1. Argument i must have monotonous entries with i(1)=0 and i(end)=1. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, huecm, magenta2yellowcm, red2yellowcm, sepiacm, whitecm

## magenta2yellowcm

Colormap with shades from magenta to yellow.

## **Syntax**

```
use colormaps
cm = magenta2yellowcm
cm = magenta2yellowcm(n)
```

### **Description**

magenta2yellowcm(n) creates a color map with n entries corresponding to color shades from magenta to yellow. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, huecm, interprgbcm, red2yellowcm, sepiacm, whitecm

## red2yellowcm

Colormap with shades from red to yellow.

### **Syntax**

```
use colormaps
cm = red2yellowcm
cm = red2yellowcm(n)
```

## **Description**

red2yellowcm(n) creates a color map with n entries corresponding to color shades from red to yellow. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, sepiacm, whitecm

## sepiacm

Colormap with shades of sepia.

### **Syntax**

```
use colormaps
cm = sepiacm
cm = sepiacm(n)
```

### **Description**

sepiacm(n) creates a color map with n entries corresponding to shades of sepia. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, whitecm

### whitecm

Colormap with plain white.

## **Syntax**

```
use colormaps
cm = whitecm
cm = whitecm(n)
```

## **Description**

whitecm(n) creates a color map with n identical entries corresponding to plain white. The color map is an n-by-3 array with one color per row; columns correspond to red, green, and blue components as real numbers between 0 to 1 (maximum intensity). The default value of n is 256.

The color map is suitable as the input argument of colormap.

#### See also

colormap, black2orangecm, black2red2whitecm, blue2greencm, blue2yellow2redcm, cyan2magentacm, graycm, green2yellowcm, huecm, interprgbcm, magenta2yellowcm, red2yellowcm, sepiacm

## 7.15 polyhedra

Library polyhedra implements functions which create solid shapes with polygonal facesin 3D. Solids are displayed with plotpoly. They are defined by the coordinates of their vertices and by the list of vertex indices for each face. Other solids, such as cylinder and sphere, are generated with parametric equations and displayed with surf. Some solids have parameters, e.g. for the number of discrete values used for parameters. When called without output argument, with an optional trailing string argument for the edge style, the solid is displayed with the current scaling and color map. With output arguments, arrays X, Y, Z expected by surf, mesh and plotpoly, and index array expected by plotpoly, are produced. They can be modified to move, scale or stretch the solids.

The following statement makes available functions defined in polyhedra:

use polyhedra

## **Functions**

### cube

Create a cube.

## **Syntax**

```
use polyhedra
cube;
cube(style);
(X, Y, Z, ind) = cube
```

## **Description**

Without output argument, cube displays a cube, i.e. a convex solid whose six faces are squares. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, cube produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

tetrahedron, octahedron, dodecahedron, icosahedron, plotpoly

### dodecahedron

Create a regular dodecahedron.

### **Syntax**

```
use polyhedra
dodecahedron;
dodecahedron(style);
(X, Y, Z, ind) = dodecahedron
```

### **Description**

Without output argument, dodecahedron displays a regular convex dodecahedron, i.e. a convex solid whose twelve faces are regular pentagons. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, dodecahedron produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

tetrahedron, cube, octahedron, icosahedron, greatdodecahedron, greatstellateddodecahedron, smallstellateddodecahedron, plotpoly

## greatdodecahedron

Create a great dodecahedron.

## Syntax

```
use polyhedra
greatdodecahedron;
greatdodecahedron(style);
(X, Y, Z, ind) = greatdodecahedron
```

## **Description**

Without output argument, greatdodecahedron displays a great dodecahedron, i.e. a regular nonconvex solid whose twelve faces are regular pentagons. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, greatdodecahedron produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

dodecahedron, greatstellateddodecahedron, greaticosahedron, plotpoly

## greaticosahedron

Create a great dodecahedron.

### **Syntax**

```
use polyhedra
greaticosahedron;
greaticosahedron(style);
(X, Y, Z, ind) = greaticosahedron
```

### **Description**

Without output argument, greaticosahedron displays a great icosahedron, i.e. a regular nonconvex solid whose twenty faces are equilateral triangles. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, greaticosahedron produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

icosahedron, greatdodecahedron, plotpoly

## greatstellateddodecahedron

Create a great stellated dodecahedron.

## **Syntax**

```
use polyhedra
greatstellateddodecahedron;
greatstellateddodecahedron(style);
(X, Y, Z, ind) = greatstellateddodecahedron
```

### **Description**

Without output argument, greatstellateddodecahedron displays a great stellated dodecahedron, i.e. a regular nonconvex solid whose twelve faces are regular star pentagons and where each vertex is common to three faces. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, greatstellateddodecahedron produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

dodecahedron, greatdodecahedron, smallstellateddodecahedron, plotpoly

### icosahedron

Create a regular icosahedron.

### **Syntax**

```
use polyhedra
icosahedron;
icosahedron(style);
(X, Y, Z, ind) = icosahedron
```

## **Description**

Without output argument, icosahedron displays a regular convex icosahedron, i.e. a convex solid whose twenty faces are equilateral triangles. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, icosahedron produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

tetrahedron, cube, octahedron, dodecahedron, plotpoly

### octahedron

Create a regular octahedron.

### **Syntax**

```
use polyhedra
octahedron;
octahedron(style);
(X, Y, Z, ind) = octahedron
```

### **Description**

Without output argument, octahedron displays a regular octahedron, i.e. a convex solid whose eight faces are equilateral triangles. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, octahedron produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

tetrahedron, cube, dodecahedron, icosahedron, plotpoly

### smallstellateddodecahedron

Create a small stellated dodecahedron.

## **Syntax**

```
use polyhedra
smallstellateddodecahedron;
smallstellateddodecahedron(style);
(X, Y, Z, ind) = smallstellateddodecahedron
```

## **Description**

Without output argument, smallstellateddodecahedron displays a small stellated dodecahedron, i.e. a regular nonconvex solid whose twelve faces are regular star pentagons and where each vertex is common to five faces. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, smallstellateddodecahedron produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

dodecahedron, greatdodecahedron, greatstellateddodecahedron, plotpoly

### tetrahedron

Create a regular tetrahedron.

### **Syntax**

```
use polyhedra
tetrahedron;
tetrahedron(style);
(X, Y, Z, ind) = tetrahedron
```

### **Description**

Without output argument, tetrahedron displays a regular tetrahedron, i.e. a solid whose four faces are equilateral triangles. By default, edges are not drawn. An optional string input argument specifies the edge style.

With four output arguments, tetrahedron produces the X, Y, Z and ind arrays expected by plotpoly, and it does not display anything.

#### See also

cube, octahedron, dodecahedron, icosahedron, plotpoly

## 7.16 solids

Library solids implements functions which create solid shapes in 3D. Solids are generated with parametric equations and displayed with surf. When called without output argument, with an optional trailing string argument for the edge style, the solid is displayed with the current scaling and color map. With output arguments, arrays X, Y, Z expected by surf or mesh are produced. They can be modified to move, scale or stretch the solids.

The following statement makes available functions defined in solids:

use solids

## **Functions**

#### cone

Cone.

### Syntax

```
use solids
cone
cone(cap)
cone(cap, n)
cone(cap, n, style)
(X, Y, Z) = cone
(X, Y, Z) = cone(n)
```

### **Description**

Without output argument, cone draws a cone approximated by a polyhedron. The optional first input argument, a logical value which is true by default, specifies if the cap is included. The optional second input argument, an integer, specifies the number of discrete values for the parameter which describes its surface.

By default, edges are not drawn. An optional third input argument, a string, specifies the edge style; it corresponds to the style argument of surf.

With three output arguments, cone produces the X, Y, and Z arrays expected by surf or mesh, and it does not display anything.

#### See also

cylinder, sphere, cube, surf

### crosscap

Cross-cap.

### **Syntax**

```
use solids
crosscap
crosscap(n)
crosscap(n, style)
(X, Y, Z) = crosscap
(X, Y, Z) = crosscap(n)
```

## Description

Without output argument, crosscap draws a cross-cap (a self-intersecting surface) approximated by a polyhedron. With an input argument, crosscap(n) draws a cross-cap where the two parameters which describe its surface are sampled with n discrete values.

By default, edges are not drawn. An optional second input argument, a string, specifies the edge style; it corresponds to the style argument of surf.

With three output arguments, crosscap produces the X, Y, and Z arrays expected by surf or mesh, and it does not display anything.

#### See also

klein, klein8, sphere, sphericon, surf

## cylinder

Cylinder.

### **Syntax**

```
use solids
cylinder
cylinder(cap)
cylinder(cap, n)
cylinder(cap, n, style)
(X, Y, Z) = cylinder
(X, Y, Z) = cylinder(n)
```

## **Description**

Without output argument, cylinder draws a cylinder approximated by a polyhedron. The optional first input argument, a logical value which is true by default, specifies if caps are included. The optional second input argument, an integer, specifies the number of discrete values for the parameter which describes its surface.

By default, edges are not drawn. An optional third input argument, a string, specifies the edge style; it corresponds to the style argument of surf.

With three output arguments, cylinder produces the X, Y, and Z arrays expected by surf or mesh, and it does not display anything.

#### See also

cone, sphere, torus, cube, surf

#### klein

Klein bottle.

### **Syntax**

```
use solids
klein
klein(p)
klein(p, n)
klein(p, n, style)
(X, Y, Z) = ...
```

### **Description**

Without output argument, klein draws a Klein bottle approximated by a polyhedron. With an input argument, klein(p) uses parameters stored in structure p. The following fields are used:

Field	Description	Default value
r0	average tube radius	0.7
d	tube variation	0.5
h	half height	3

With two input arguments, klein(p,n) draws a Klein bottle where the two parameters which describe its surface are sampled with n discrete values.

By default, edges are not drawn. An optional third input argument, a string, specifies the edge style; it corresponds to the style argument of surf.

With three output arguments, klein produces the X, Y, and Z arrays expected by surf or mesh, and it does not display anything.

#### See also

klein8, crosscap, surf

## klein8

Figure 8 Klein bottle immersion.

## **Syntax**

```
use solids
klein8
klein8(r)
klein8(r, n)
klein8(r, n, style)
(X, Y, Z) = ...
```

### **Description**

Without output argument, klein8 draws a figure 8 Klein bottle immersion (a closed, self-intersecting surface with one face) approximated by a polyhedron. With an input argument, klein8(r) draws the surface with a main radius of r (the default value is 1).

With two input arguments, klein8(r,n) samples the two parameters which describe its surface with n discrete values.

By default, edges are not drawn. An optional third input argument, a string, specifies the edge style; it corresponds to the style argument of surf.

With three output arguments, klein8 produces the X, Y, and Z arrays expected by surf or mesh, and it does not display anything.

#### See also

klein, crosscap, surf

## sphere

Sphere.

## Syntax

```
use solids
sphere
sphere(n)
sphere(n, style)
(X, Y, Z) = sphere
(X, Y, Z) = sphere(n)
```

## **Description**

Without output argument, sphere draws a sphere approximated by a polyhedron. With an input argument, sphere(n) draws a sphere where the two parameters which describe its surface are sampled with n discrete values.

By default, edges are not drawn. An optional second input argument, a string, specifies the edge style; it corresponds to the style argument of surf.

With three output arguments, sphere produces the X, Y, and Z arrays expected by surf or mesh, and it does not display anything.

#### See also

cylinder, cone, torus, cube, surf

## sphericon

Sphericon.

### **Syntax**

```
use solids
sphericon
sphericon(n)
sphericon(n, style)
(X, Y, Z) = sphericon
(X, Y, Z) = sphericon(n)
```

### **Description**

Without output argument, sphericon draws a sphericon (a 3D shape made from a bicone with a 90-degree apex, cut by a plane containing both apices, where one half is rotated by 90 degrees) approximated by a polyhedron. With an input argument, sphericon(n) draws a sphericon where the two parameters which describe its surface are sampled with n discrete values.

By default, edges are not drawn. An optional second input argument, a string, specifies the edge style; it corresponds to the style argument of surf.

With three output arguments, sphericon produces the X, Y, and Z arrays expected by surf or mesh, and it does not display anything.

#### See also

```
sphere, crosscap, surf
```

#### torus

Torus.

## **Syntax**

```
use solids
torus
torus(r)
torus(r, n)
torus(r, n, style)
(X, Y, Z) = ...
```

### **Description**

Without output argument, torus draws a torus approximated by a polyhedron with a main radius of 1 and a tube radius of 0.5. With an input argument, torus(r) draws a torus with tube radius r. With two input arguments, torus(r,n) draws a torus where the two parameters which describe its surface are sampled with n discrete values.

By default, edges are not drawn. An optional third input argument, a string, specifies the edge style; it corresponds to the style argument of surf.

With three output arguments, torus produces the X, Y, and Z arrays expected by surf or mesh, and it does not display anything.

#### See also

sphere, cylinder, surf

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