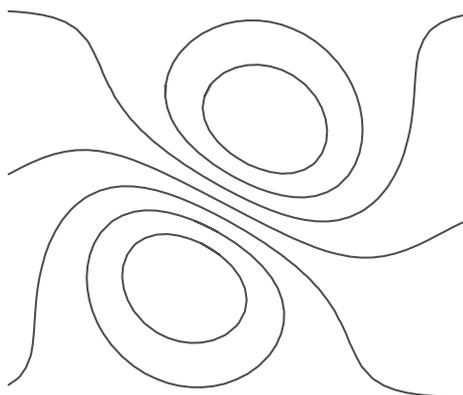


Calerga



**LME for Pocket PC 071208**  
User Manual

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**LME for Pocket PC User Manual, December 2007.**

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Most of the material in LME Pocket PC 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

LME for Pocket PC is LME, the programming language of Sysquake, compiled in a simple application for Windows Mobile ARM-based handheld devices. It has been tested on a Dell Axim X30 device, but should also run on other Axim and HP iPaq devices.

LME for Pocket PC is a *technology preview*. Its user interface is basic, and there is support only for static graphics, not for interactive graphics. It is based on LME, a Matlab-compatible programming language aimed at numerical computing used in Sysquake and other products of Calerga since 1997. You should install it only if you want to bring Matlab compatibility in your pocket. There is no expiration date. We do not guarantee that LME for Pocket PC will be a supported product in the long term, but we intend to continue providing solutions for Pocket PC based on LME in the future.

LME for Pocket PC is neither Sysquake nor LyME. It has neither the interactive graphics of Sysquake nor its SQ file support which makes developing user interfaces so easy. It has not the simple yet efficient user interface of LyME, with its integrated text editor. It is more a first step in the direction of a full Sysquake on Pocket PC: it supports files input/output and extensions, like Sysquake. Its graphical capabilities are borrowed from Sysquake Remote, i.e. it is fully compatible with Sysquake without the interactivity. Some of the most powerful extensions have been ported: audio input/output, web services with XML-RPC and SOAP, and TCP/IP sockets. And you can develop yours in C. It even supports multithreading. But as a technology preview on the way to something more finished, it has not received its own product name.

## 1.1 LME Installation

LME for Pocket PC is provided as a .cab file. One way to install it is to download it to your Pocket PC device, to locate it in the File Explorer and to open it. LME and all its support files are installed in \Program Files\Calerga, with a shortcut to the application in the Programs folder.

At startup, LME loads extensions in the subdirectory LMExt, executes the contents of lmstartup.txt if it exists, and restores the history of past commands from the file lmehistory.txt. All these directories and files must be located in the same directory as LME.exe. All of them are optional (lmehistory.txt is created when LME exits if it does not exist).

LME comes with libraries (files ending with the double suffix .lml.txt) which add useful functions in different domains, such as statistics, physical constants, signal processing or control. LME will find them only if they are in one of the directories enumerated in the Options. When you run LME for the first time, you should check in menu Tools>Options that the Paths of Libraries include \Program Files\Calerga\Lib .

## 1.2 User interface

When launched, LME presents a basic user interface with the following elements:

**Command input** The text field at the top of the screen lets you write commands with the LME syntax, such as  $1+2$  or  $v=1:5;fft(v)$ .

**Button Execute** Tapping the Execute button makes LME interpret the contents of the input field. You can interrupt computation by pressing the Scroll Down hardware button.

**History** To retrieve a previous command and display it in the command input, tap the button with the < sign. To go forward in the list of commands, tap the > button. You can edit the command before executing it.

**Output** The largest text field displays the result of commands. You cannot write directly to it. You can toggle between text output and graphics with the View menu, or by pressing the Select hardware button when the command field is empty.

**Menu** Menus at the bottom of the screen let you change between text and graphics, see the tree of code fragments (see below), change options, and open the user manual in the default HTML browser (usually Internet Explorer).

If the operating system supports it, you can rotate the screen; the layout will be optimized for the new orientation.

## 1.3 Code fragments

To discover the power of LME, you can execute small demonstrations, called *code fragments*. Code fragments are displayed as a tree with the menu Tools>Code fragments. To execute a fragment, double-tap its name. The command itself and the result are displayed. If the result is graphical, you can switch to text mode with the menu View>Text (or simply by pressing the Select hardware button when the command field is empty) to see the command.

All fragments are stored in the XML file `lme_code_fragments.xml`. To see or edit its contents, you can use a text editor or open it in a browser which supports CSS2 style sheets.

## 1.4 First steps

Here are some commands you can try.

```
1+2*3
```

Simple expressions follow the standard syntax found in many computing languages; if they are not followed by a semicolon, the result is displayed.

To permit you to chain computations easily, LME stores the anonymous result of expressions into the variable `ans`, which you can reuse in the next expression:

```
ans + sin(pi/3)
```

To keep multiple results, you can assign them to other variables:

```
x = 5
```

Operators and functions also support complex numbers and matrices. This will compute the complex eigenvalues of a 2-by-2 matrix:

```
eig([1,2;-1,1])
```

In addition to numerical results, LME can display graphics. `fplot` plots a function, inline in the example below, over some range:

```
fun = inline('(x+0.3)^2+7*exp(-3*x^2)');
fplot(fun, [-2,3]);
```

To clear the figure, use `clf`. You can switch manually between graphics and text output with the View menu.

LME also supports audio input and output (provided the hardware it runs on does!) We begin by recording one second of audio samples in an array. Say something after you tap the Execute button. The semicolon is important here; we do not want to see 88100 samples scrolling on the screen. If you forget it, interrupt the execution by pressing the selection button of your handheld.

```
snd = audiorecord(1);
```

The samples are stored as an array of two columns, one for the left channel, one for the right. They can be played back:

```
audioplay(snd)
```

There are probably simpler ways to record audio on a Pocket PC device. But with LME, you can manipulate the samples easily. For instance, you can play them backward by flipping the array upside-down:

```
audioplay(flipud(snd))
```

To learn more about LME, you can follow the LME tutorial.

## 1.5 User preferences

You can customize the way LME is working in the dialog displayed with the menu Tools>Options. Here is a description of the options:

**Memory** Amount of memory reserved for LME variables, programs, and data used for executing code.

**Evaluate assert** In LME programs, the `assert` function can help in reporting errors during development. If this option is off, the evaluation of `assert` is skipped, which can provide slightly improved performance. Usually, you should switch it on during development, and off when using functions whose you trust the quality.

**Load extensions** Extensions are files which provide additional functions to LME. Unlike functions written in the LME programming language, extensions are compiled to native machine code, so they are very fast and they have access to all the features of the operating system. LME is provided with several extensions; you can also develop your own.

Extensions are stored in the directory `LMEEExt`, itself located at the same place as `LME.exe`. To disable all extensions, you can switch off the Load extensions option.

**Hide messages** Extensions usually display a short status message when they are loaded, at LME startup. When Hide messages is switched on, no message is displayed.

**Path of libraries** Libraries contain additional functions written in the LME programming language (see below). The Path of libraries option specifies where LME looks for library files. Directories are separated with semicolons. For example, if you store your own libraries in \My Documents\LMELib, you could add this directory to the default path: \Program Files\Calerga\Lib;\My Documents\LMELib

The path of libraries can also be set with the LME function path.

**Startup commands** Startup commands are LME commands executed every time the application is launched. For instance, if you often use functions from libraries (such as stdlib for function which extend the built-in functions of LME or stat for advanced functions related to statistics), you can add use statements as startup commands to make them always available: use stdlib, stat

Startup commands can also be used to define variables. They are executed after the file lmestartup.txt, if it exists (see above).

**Format** Format specified in the Options dialog are an alternative to the LME function format. It defines how numbers are displayed.

**Line width** The default line width can be set to 1 or 2 pixels. In graphics commands, an uppercase color letter in the style argument increases the width to 2 or 3 pixels, respectively.

**Font** The default font used in figures can be set. The font used by text can be changed with fontset; currently the option Font can be 'serif', 'sansserif' or 'monospace', and Size, Italic and Underline are ignored.

## 1.6 Libraries

LME for Pocket PC does not implement its own text editor. Functions are stored in library files. To make their editing easier, library files end with .lml.txt (double suffix). Tapping their icon in File Explorer should open them with the standard application used for editing text files, typically Pocket Word; the .txt suffix is hidden. When you create a new library, make sure you add a .lml.txt suffix and save it as text only. When you transfer libraries from a desktop application, for instance Sysquake, you should also add this suffix if you want to edit

it on the Pocket PC device (LME also finds library files with a simple `.lml` suffix, but you cannot edit them as easily).

LME finds libraries in the directory `Lib` in the same directory as `LME.exe`. If you want to store libraries in other places, you should add directories with the Options dialog or the command path (see above).

## Chapter 2

# LME Tutorial

The remainder of this chapter introduces LME(TM) (Lightweight Matrix Engine), the interpreter for numerical 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.

### 2.1 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:

```
> 2^5
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
```

## 2.2 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 =  
2-3j
```

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

```

## 2.3 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 `*` 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
```

## 2.4 Polynomials

LME handles only numerical values. Therefore, it cannot differentiate functions like  $f(x) = \sin(e^x)$ . However, a class of functions has a paramount importance in numerical computing, the polynomials. Polynomials are weighted sums of powers of a variable, such as  $2x^2 + 3x - 5$ . LME, which handles only matrices, 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  $(x + 2)(2x^2) + 4x + 5 = 2x^3 + 8x^2 + 13x + 10$ .

## 2.5 Strings

You type strings by bracketing 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
```

## 2.6 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 =
  9
```

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
```

## 2.7 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
```

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

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

## 2.8 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
  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

```

## 2.9 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 initialize 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 already exists, it is discarded.

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



## Chapter 3

# LME Reference

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

### 3.1 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.

## 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 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
<code>line</code>	<code>n</code>	Set the current line number to <code>n</code>

`_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.

## 3.2 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...
[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.

In some cases, a simpler syntax can be used when the function has only literal character strings as input arguments. 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 str1 'str 2'  fun('str1','str 2')

```

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

### 3.3 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.

### 3.4 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-numerical types exist for character arrays and logical (boolean) arrays. Cell arrays, which contain composite types, are described in a section below.

<b>Type</b>	<b>Description</b>
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 numerical 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 > uint64 > int64 > uint32 > int32 > uint16 > int16 >
uint8 > int8
```

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

To convert arrays to numerical, char, or logical arrays, use functions `+` (unary operator), `char`, or `logical` respectively. To convert the numerical 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 numerical 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. `2:int16` 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:

<b>Expression</b>	<b>Value</b>
<code>uint64(123456789012345678)</code>	<code>123456789012345696</code>
<code>123456789012345678uint64</code>	<code>123456789012345678</code>

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.	Character code
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	\ooo	ooo
16-bit UTF-16	\uhhhh	unicode UTF-16 code

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 (multi-byte 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).

## 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 two ways to make structures: 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.

```
a = struct('name', 'Sysquake', ...
          'os', {'Windows', 'Mac OS X', 'Linux'});
```

```
b.x = 200;
b.y = 280;
b.radius = 90;
```

```
c.s = b;
```

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

## 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 numerical 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 numerical types in array concatenation. Numerical sets and list or cell arrays of strings cannot be mixed.

## 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. Classes 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 classes` imports the definitions of methods for class `polynom` and others.

```
use classes;
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
```

### 3.5 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.

## 3.6 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

```
> factor({2})

Wrong type (stdlib:primes:164) 'ones'
-> stdlib:factor:174
```

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 `stdlib` is the library name, `primes` is the function name, and 164 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, `primes` was called by `factor` at line 174 in library `stdlib`.

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 numerical 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.
- 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 numerical 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 structures** Not enough memory to compile that many nested programming structures 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.

**"catch" without "try"** The catch statement does not match a try 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.

### 3.7 List of Commands, Functions, and Operators

#### Programming keywords

break	error	repeat
case	for	rethrow
catch	function	return
clear	global	switch
continue	if	try
define	otherwise	until
endfunction	persistent	use
else	private	useifexists
elseif	public	while

## Programming operators and functions

assert	fun2str	nargout
Variable assignment	help	sandbox
Operator ()	info	sandboxtrust
Operator @	inline	str2fun
builtin	inmem	str2obj
deal	isglobal	subsasgn
dumpvar	iskeyword	subsref
eval	lasterr	variables
exist	lasterror	which
feval	lookfor	
fevalx	nargin	

## Debugging

dbclear	dbstack	dbtype
dbcont	dbstatus	echo
dbhalt	dbstep	profile
dbquit	dbstop	

## Arrays

[]	ind2sub	ones
,	interp	permute
;	ipermute	rand
:	iscell	randn
beginning	isempty	repmat
cat	length	reshape
cell	magic	rot90
cellfun	matrixcol	size
diag	matrixrow	sort
end	meshgrid	squeeze
eye	ndgrid	sub2ind
find	ndims	tril
flipdim	nnz	triu
fliplr	num2cell	unique
flipud	numel	zeros

**Strings**

base64decode	isspace	strcmpi
base64encode	length	strmatch
char	lower	strtok
deblank	md5	strtrim
findstr	setstr	upper
ischar	sha1	utf8decode
isdigit	sprintf	utf8encode
isempty	sscanf	
isletter	strcmp	

**Lists**

{}	islist	num2list
apply	length	replist
join	list2num	
isempty	map	

**Structures**

fieldnames	isstruct	setfield
getfield	orderfields	struct
isfield	rmfield	

**Objects**

class	isobject
isa	methods

## Logical operators

==	>	
===	<=	&&
~=	>=	
~=	~>=	?
<	&	

## Logical functions

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

## 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	intmax	pi
false	intmin	realmax
i	j	realmin
inf	nan	true

## Arithmetic functions

+	\	diff
-	.\	kron
*	^	mod
.*	.^	prod
/	cumprod	rem
./	cumsum	sum

## Trigonometric and hyperbolic functions

acos	asinh	csch
acosh	atan	sec
acot	atan2	sech
acoth	atanh	sin
acsc	cos	sinh
acsch	cosh	tan
asec	cot	tanh
asech	coth	
asin	csc	

## Other scalar math functions

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

## Type conversion functions

cast	fix	swapbytes
ceil	floor	typecast
complex	round	
double	single	

## Matrix math functions

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

## Geometry functions

cross      dot

## Statistic functions

cov	mean	skewness
kurtosis	min	std
max	moment	var

## Polynomial math functions

addpol	filter	polyint
conv	poly	polyval
deconv	polyder	roots

## Triangulation functions

delaunay	griddatan	voronoi
delaunayn	tsearch	voronoin
griddata	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	

## Quaternion other 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

## Non-linear numerical functions

fminbnd	ode23	optimset
fminsearch	ode45	quad
fzero	odeset	

## Dynamical systems functions

c2dm	dmargin	ss2tf
d2cm	margin	tf2ss

## Input/output

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

## Files

fopen

## Graphical functions specific to LME for Pocket PC

clf      drawnow

## Basic graphics

activeregion	fplot	polar
area	image	quiver
bar	label	scale
barh	legend	scalefactor
circle	line	subplot
colormap	pcolor	text
contour	plot	title
fontset	plotoption	

### 3D graphics

camdolly	camup	material
camorbit	camva	mesh
campan	camzoom	plot3
campos	contour3	plotpoly
camproj	daspect	sensor3
camroll	lightangle	surf
camtarget	line3	

### Graphics for dynamical systems

bodemag	dsigma	nichols
bodephase	dstep	nyquist
dbodemag	erlocus	plotroots
dbodephase	hgrid	rlocus
dimpulse	hstep	sgrid
dinitial	impulse	sigma
dlsim	initial	step
dnichols	lsim	zgrid
dnyquist	ngrid	

### Date and time

cal2julian	julian2cal	toc
clock	tic	

### Threads

semaphoredelete	semaphoreunlock	threadset
semaphorelock	threadkill	threadsleep
semaphorenew	threadnew	

**Extensions loaded on demand**

exteval    extload    extunload

**LME for Pocket PC functions**

lmepath    path

**3.8 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 user-defined classes. Its definition should 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 functions 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 functions 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 `[1,1,1;2,2,2]`.

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 user-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  $A(\text{expr})$ , `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  $[1,2,3;1,2,3]$ .

In an expression used as the first of multiple subscripts to access some elements of an array  $A(\text{expr}, \dots)$ , `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 user-defined classes. Its definition should have a header equivalent to `function e=C::matrixrow(obj,i,n)`, where  $C$  is the name of the class,  $\text{obj}$  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 = subsasgn(A, s, B)
```

**Description**

When an assignment is made to a subscripted part of an object in a statement like  $A(s_1, s_2, \dots) = B$ , LME executes  $A = \text{subsasgn}(A, s, B)$ , 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  $\{s_1, s_2, \dots\}$ . 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 when `s.type` is a list, and for structures when `s.type` is `'.'`.

**Examples**

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

**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 `suboref(A, s)`, where `suboref` 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 `suboref(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 `suboref` is to permit the use of subscripts with objects, a built-in implementation of `suboref` is provided for arrays when `s.type` is `'()''`, for lists when `s.type` is `'{}'`, and for structures when `s.type` is `'.'`.

## Examples

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

## See also

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

## 3.9 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 keyword is reserved for future use:

`goto`

### **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`

**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.

## Syntax

```
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 =
    1
    i =
    2
    i =
    3
for i = (1:3)'; i, end
    i =
    1
    2
    3
for i = 1:2:5; end; i
    i =
    5
for i = 1:3; break; end; i
    i =
    1
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
```

**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 never require `endfunction`). Like `function`, `endfunction` must be the first element of a line.

Not all of the input and output arguments are necessarily used. The caller fixes the number of input and output arguments, which can be retrieved by the called function with `nargin` and `nargout`, respectively. The unused 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. The unused output arguments (from `nargout+1` to the last one) do not have to be set, but may be.

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

```

    if nargin == 1    % product of elements of x
        Prod = prod(x);
    else            % product of x and y
        Prod = x .* y;
    end
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 nargin < 2
        b = a;
    end
    if nargin < 3
        ellipse = false;
    end
    S = ellipse ? pi * a * b / 4 : a * b;

```

## See also

return, nargin, nargout, define, inline, global, persistent

## 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 exist, it does nothing.

### **See also**

`include`, `useifexists`, `private`, `public`

## **otherwise**

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

### **See also**

`switch`

## **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` may 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
        so
        ...
end

switch string
    case str1
        s1
        ...
    case str2
        s2
        ...
    case {str3,str4,...}
        s34
        ...
    otherwise
        so
        ...
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

if

**try**

Error recovery.

**Syntax**

```
try
  ...
end
```

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

**Description**

The statements after `try` are executed. If an error occurs, execution is switched to the statements following `try`, if any, or to the statements following `end`. 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 Mac OS X, 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.

**Syntax**

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

**Description**

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

All `use` statements are parsed before execution begins. They may 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.

## Syntax

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

## 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
```

**See also**

repeat, for, break, continue, if

## 3.10 Debugging Commands

**dbclear**

Remove a breakpoint.

**Syntax**

```
dbclear fun
dbclear fun line
dbclear('fun', line)
dbclear
```

**Description**

`dbclear fun` removes all breakpoints in function `fun`. `dbclear fun line` or `dbclear('fun', line)` removes the breakpoint in function `fun` at line number `line`.

Without argument, `dbclear` removes all breakpoints.

**See also**

`dbstop`, `dbstatus`

**dbcont**

Resume execution.

**Syntax**

```
dbcont
```

**Description**

When execution has been suspended by a breakpoint or `dbhalt`, it can be resumed from the command-line interface with `dbcont`.

**See also**

`dbstop`, `dbhalt`, `dbstep`, `dbquit`

**dbhalt**

Suspend execution.

**Syntax**

```
dbhalt
```

**Description**

In a function, `dbhalt` suspends normal execution as if a breakpoint had been reached. Commands `dbstep`, `dbcont` and `dbquit` can then be used from the command line to resume or abort execution.

**See also**

`dbstop`, `dbcont`, `dbquit`

**dbquit**

Abort suspended execution.

**Syntax**

```
dbquit
```

**Description**

When execution has been suspended by a breakpoint or `dbhalt`, it can be aborted completely from the command-line interface with `dbquit`.

**See also**

`dbstop`, `dbcont`, `dbhalt`

## dbstack

Chain of function calls.

### Syntax

```
dbstack
s = dbstack
dbstack all
s = dbstack('all')
```

### Description

dbstack displays the chain of function calls which lead to the current execution point, with the line number where the call to the subfunction is made. It can be executed in a function or from the command-line interface when execution is suspended with a breakpoint or dbhalt.

dbstack all (or dbstack('all')) displays the whole stack of function calls. For instance, if two executions are successively suspended at breakpoints, dbstack displays only the second chain of function calls, while dbstack all displays all functions.

With an output argument, dbstack returns the result in a list of structures. Each structure contains the function name (or class and method names) in field name and the line number in field line. Note that you cannot assign the result of dbstack to a new variable in suspended mode.

### Examples

```
use stdlib
dbstop primes
isprime(113)
<primes:164> p = ones(1, n);
dbstack
  stdlib/primes;164
  stdlib/isprime;157
dumpvar('stack', dbstack)
  stack = {struct('name','stdlib/primes', ...
    'line',164), ...
    struct('name','stdlib/isprime', ...
    'line',157)};
```

### See also

dbstop, dbhalt

## **dbstatus**

Display list of breakpoints.

### **Syntax**

```
dbstatus  
dbstatus fun
```

### **Description**

`dbstatus` displays the list of all breakpoints. `dbstatus fun` displays the list of breakpoints in function `fun`.

### **See also**

`dbstop`, `dbclear`, `dbtype`

## **dbstep**

Execute a line of instructions.

### **Syntax**

```
dbstep  
dbstep in  
dbstep out
```

### **Description**

When normal execution is suspended after a breakpoint set with `dbstop` or the execution of function `dbhalt`, `dbstep`, issued from the command line, executes the next line of the suspended function. If the line is the last one of the function, execution resumes in the calling function.

`dbstep in` has the same effect as `dbstep`, except if a subfunction is called. In this case, execution is suspended at the beginning of the subfunction.

`dbstep out` resumes execution in the current function and suspends it in the calling function.

**Example**

Load library `stdlib` and put a breakpoint at the beginning of function `linspace`:

```
use stdlib
dbstop linspace
```

Start execution of function `linspace` until the breakpoint is reached (the next line to be executed is displayed):

```
v = linspace(1,2,5)
<linspace:8>   if nargin < 3
```

When the execution is suspended, any function can be called. Local variables of the function can be accessed and changed; but no new variable can be created. Here, the list of variables and the value of `x2` are displayed:

```
info v
  r (not defined)
  x1 (1x1)
  x2 (1x1)
  n (1x1)
x2
  x2 =
    2
```

Display the stack of function calls:

```
dbstack
  stdlib/linspace;8
```

Execute next line:

```
dbstep
<linspace:11>   r = x1 + (x2 - x1) * (0:n-1) / (n-1);
```

Execute last line; then normal execution is resumed:

```
dbstep
v =
  1  1.25  1.5  1.75  2
```

Display breakpoint and clear it:

```
dbstatus
  stdlib/linspace;0
dbclear
```

**See also**

dbstop, dbcont, dbquit

**dbstop**

Set a breakpoint.

**Syntax**

```
dbstop fun
dbstop fun line
dbstop('fun', line)
```

**Description**

`dbstop fun` sets a breakpoint at the beginning of function `fun`. `dbstop fun line` or `dbstop('fun', line)` sets a breakpoint in function `fun` at line `line`.

When LME executes a line where a breakpoint has been set, it suspends execution and returns to the command-line interface. The user can inspect or change variables, executes expressions or other functions, continue execution with `dbstep` or `dbcont`, or abort execution with `dbquit`.

**Example**

```
use stdlib
dbstop linspace
dbstatus
    stdlib/linspace;0
dbclean linspace
```

**See also**

dbhalt, dbclear, dbstatus, dbstep, dbcont, dbquit, dbtype

**dbtype**

Display source code with line numbers, breakpoints, and current execution point.

**Syntax**

```
dbtype fun
dbtype
```

## Description

`dbtype fun` displays the source code of function `fun` with line numbers, breakpoints, and the position where execution is suspended (if it is in `fun`). Without argument, `dbtype` displays the function which is suspended.

`dbtype` can be used at any time to check the source code of any function known to LME.

## Example

```
use stdlib
dbstop linspace
linspace(1,2,5);
  <linspace:8>   if nargin < 3
dbstep
  <linspace:11>   r = x1 + (x2 - x1) * (0:n-1) / (n-1);
dbtype
#   6 function r = linspace(x1, x2, n)
#   7
#   8   if nargin < 3
#   9     n = 100;
#  10   end
>  11   r = x1 + (x2 - x1) * (0:n-1) / (n-1);
```

## See also

`dbstatus`, `dbstack`, `echo`

## echo

Echo of code before its execution.

## Syntax

```
echo on
echo off
echo fun on
echo(state)
echo(state, fd)
echo(fun, state)
echo(fun, state, fd)
```

## Description

`echo on` enables the display of an echo of each line of function code before execution. The display includes the function name and the line

number. `echo off` disables the echo.

The argument can also be passed as a boolean value with the functional form `echo(state)`: `echo on` is equivalent to `echo(true)`.

`echo fun on` enables echo for function named `fun` only. `echo fun off` disables echo (the function name is ignored); `echo off` has the same effect.

By default, the echo is output to the standard error channel (file descriptor 2). Another file descriptor can be specified as an additional numeric argument, with the functional form only.

## Example

Trace of a function:

```
use stdlib
echo on
C = compan([2,5,4]);
  compan 26 if min(size(v)) > 1
  compan 29 v = v(:).';
  compan 30 n = length(v);
  compan 31 M = [-v(2:end)/v(1); eye(n-2, n-1)];
```

Echo stored into a file 'log.txt':

```
fd = fopen('log.txt', 'w');
echo(true, fd);
...
echo off
fclose(fd);
```

## See also

`dbtype`

## 3.11 Profiler

### profile

Install, remove, or display a function profile.

### Syntax

```
profile fun
profile report
profile done
```

```
profile function fun
```

```
profile off
profile on
profile reset
profile('report', format)
```

## Description

The purpose of the profiler is to measure the amount of time spent executing each line of code of a function. This helps in evaluating where effort should be put in order to optimize the code. With LME, a single function can be profiled at any given time. Command `profile` manages all aspects related to profiling, from specifying which function is to be profiled to displaying the results and resuming normal operations.

The time measured for each line includes time spent executing sub-functions called from that line. Only the cumulative times are collected; lines of code in loops are likely to have a larger impact on the overall execution time.

The profile accuracy is limited mainly by two factors:

- The resolution of the timer, which is typically between 1e-9 and 1e-6 second. This has obviously a larger effect on lines executed quickly. Lines which contain scalar assignments or statements like `if` and `for` may completely escape from the timing.
- The time overhead to perform the timing and add the data. Here again, its effect is more dramatic with fast lines.

To profile a function, one usually proceeds in four steps:

**Setup** `profile fun` sets up profiling for function `fun`. Room in memory is allocated and initialized for collecting the cumulative time of execution for each line in `fun`.

**Function execution** Each execution of the function adds to the profile data. Since the relative execution times are usually what is really interesting, you may want to execute the function several times to reduce fluctuations due to rounding errors. Time spent outside the function (such as the time you spend typing the commands at the command-line interface) is not included.

**Profile report** `profile report` displays a report for the function being profiled. The default format is a listing of all lines with the line number, the cumulative time spent for the line in seconds, its percentage with respect to the time spent in the whole function, and the source code of the line. You can continue executing the function and creating new reports; times are cumulative (but see `profile reset` and `profile off` below).

**End** profile done releases the data structures set up with profile fun.

Other options are available. profile off suspends profiling, and profile on resumes it. When profiling is suspended, calls to the profiled function are not taken into account.

profile reset resets all the times and resumes profiling if it was suspended.

profile function fun is equivalent to profile fun, but it may also be used to profile functions with the same name as one of the options which have a special meaning for profile, like report or done.

profile('report', format) produces a report with a special format specified by the string format. This string is similar to the format argument of sprintf; it is reused for each line of the profiled function. Its characters are output literally, except for sequences which begin with a percent character, whose meaning is given in the table below.

**Char. Meaning**

%%	single %
%l	line number
%t	cumulative time
%p	percentage of the total time
%s	source code of the line

Like with sprintf, precision numbers may be inserted between the percent sign and the letter; for instance, %8.3t displays the cumulative time in a column of 8 characters with a fractional part of 3 digits. The percentage is displayed only if it is greater than 1 %; otherwise, it is replaced (together with the percent character which may follow it) with spaces. The default format is '%4l%9.3t%6.1p%% %s\n'.

**Example**

We shall profile function logspace from library stdlib (the source code of this function has been revised since the profiling was done).

```

use stdlib
profile logspace
x = logspace(1,10);
profile report
  13  0.000          function r = logspace(x1, x2, n)
  14  0.000
  15  0.000  14.8%    if nargin < 3
  16  0.000   5.8%      n = 100;
  17  0.000   2.2%    end
  18  0.000  77.1%    r = exp(log(x1)+log(x2/x1)*(0:n-1)/(n-1));

```

While the times spent for all lines are smaller than half a millisecond, the resolution is fine enough to permit relative timing of each line. The function header does not correspond to any code and is not timed. To improve the accuracy of the timing, we repeat the execution 10000 times.

```
for i=1:10000; x = logspace(1,10); end
profile report
13  0.000          function r = logspace(x1, x2, n)
14  0.000
15  0.055   8.9%   if nargin < 3
16  0.057   9.2%       n = 100;
17  0.047   7.6%   end
18  0.458  74.3%   r = exp(log(x1)+log(x2/x1)*(0:n-1)/(n-1));
```

Finally, here is a report with a different format: the first column is the percentage as an integer, a space and the percent sign, followed by spaces and the source code:

```
profile('report', '%3.0p %% %s\n')
      function r = logspace(x1, x2, n)

 9 %   if nargin < 3
 9 %       n = 100;
 8 %   end
74 %   r = exp(log(x1) + log(x2/x1) * (0:n-1) / (n-1));
```

## See also

tic, toc, sprintf

## 3.12 Miscellaneous Functions

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

### assert

Check that an assertion is true.

#### Syntax

```
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

**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 =
  2

```

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

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

Assign the elements of list l 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)

```

## 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.

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

## Examples

```
dumpvar(2+3)
5
a = 6; dumpvar('a', a)
a = 6;
s = 'abc'; dumpvar('string', s)
string = 'abc';
```

## 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, ...)
```

## 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 may 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.

## 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'

```

### See also

warning, try, lasterr, lasterror, assert, fprintf

## 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 variable ('variable').

**Examples**

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

**See also**

info

**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 inline function.

**Examples**

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

**See also**

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

## 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 by either by its name as a string, a function reference, or an inline function.

### Example

```
fevalx(@plus, 1:5, (10:10:30)')
    11    12    13    14    15
    21    22    23    24    25
    31    32    33    34    35
```

### See also

feval, meshgrid, repmat, 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 functions
info methods
info variables
info global
info persistent
info libraries
info usedlibraries
info threads
str = info
list = 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 list of structures which describe each built-in function, with the following fields:

<code>name</code>	function name
<code>module</code>	module name
<code>trusted</code>	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. 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 list of structures which describe each user-defined function, with the following fields:

```
library  library name
name     function name
loaded   true if loaded
```

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

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

`info variables` displays the list of variables with their type and size. With an output argument, `info('variables')` gives a list of structures which describe 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 persistent variable names.

`info libraries` displays the list of all loaded libraries. With an output argument, `info('libraries')` gives the list of the library names.

`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 threads` displays the ID of all threads. With an output argument, `info('threads')` gives a list of structures which describe 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`.

**Examples**

```

info
  LME 4.5
info b
  abs
  acos
  acosh
  (etc.)
info v
  ans (1x1 complex)
vars = info('v');
dumpvar(vars)
  {struct('name','ans', ...
    'defined',true), ...
  struct('name','vars', ...
    'defined',false)}

```

**See also**

`inmem`, `which`, `exist`

**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)

```

**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 list of arguments which must be supplied when the inline function is called; `paramstruct` is a structure whose fields define fixed parameters.

Anonymous functions 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):

```
fun = inline(['function (a,b)=f(v);', ...
            'a=mean(v);', ...
            '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
```

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})
```

## See also

function, operator @, feval, eval

## inmem

List of functions loaded in memory.

## Syntax

```
inmem
list = 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 a list of structures which describe 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

**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, 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', 'if',
```

```
'otherwise', 'persistent', 'private', 'public', 'repeat',  
'return', 'switch', 'try', 'until', 'use', 'useifexists',  
'while'}
```

**See also**

info, which

**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:indexOutOfRangeException
```

**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`

**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 may use the `nargin` function to know how many arguments were passed by the caller to avoid accessing the undefined variables.

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` may 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;
```

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

```
function fprintfstars(fd,n)
if nargin == 1 % fprintfstars(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 written "+"):

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

## See also

`nargout`, `varargin`, `function`

## nargout

Number of output arguments.

## Syntax

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

## Description

A function may 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` may 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)
```

## 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`.

**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

```

str2obj('1+2j')
1 + 2j
str = dumpvar({1, 'abc', 1:100})
str =
    {1, ...
      'abc', ...
      [1:100]}
str2obj(str)
{1, 'abc', real 1x100}
eval(str)
{1, 'abc', real 1x100}
str2obj('sin(2)')
Bad argument 'str2obj'
eval('sin(2)')
0.9093

```

### See also

`eval`, `dumpvar`

## varargin

Remaining input arguments.

### Syntax

```

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

```

**Description**

`varargin` is a special variable which may be used to collect input arguments. In the function declaration, it must be used as the last (or unique) input argument. 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 may 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    0    2    2    0
    0    0    0    2    2    0
    0    0    0    0    0    3
```

**See also**

`nargin`, `varargout`, `function`

**varargout**

Remaining output arguments.

**Syntax**

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

## Description

`varargout` is a special variable which may 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` may 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
v3 =
    -11    17   -13     5     9
v4 =
    28   -30    18     4
```

## See also

`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.');
```

**See also**

`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)`. If it is a variable, it is prefixed with `(var)`. If name is neither a function nor a variable, `which` returns the empty string.

## Examples

```
which logspace
  stdlib/logspace
which polynom::plus
  classes/polynom::plus
which sin
  (builtin)/sin
x = 2;
which x
  (var)/x
```

## See also

`info`

## 3.13 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, 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

### 3.14 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:

```

' . '
^ . ^
- (unary)
* .* / ./ \ .\
+ -
== ~= < > <= >= === ~==
~
&
|
&&
||
: ?
,
;
```

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 functions such as `feval` which take the name of a function as an argument.

Here is the correspondence between operators and functions:

[a;b]	vertcat(a,b)	a-b	minus(a,b)
[a,b]	horzcat(a,b)	a*b	mtimes(a,b)
a:b	colon(a,b)	a/b	mrdivide(a,b)
a:b:c	colon(a,b,c)	a\b	mldivide(a,b)
a b	or(a,b)	a.*b	times(a,b)
a&b	and(a,b)	a./b	rdivide(a,b)
a<=b	le(a,b)	a.\b	ldivide(a,b)
a<b	lt(a,b)	a^b	mpower(a,b)
a>=b	ge(a,b)	a.^b	power(a,b)
a>b	gt(a,b)	~a	not(a)
a==b	eq(a,b)	-a	uminus(a)
a~=b	ne(a,b)	+a	uplus(a)
a===b	same(a,b)	a'	ctranspose(a)
a~=b	unsame(a,b)	a.'	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 numerical 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 sometimes 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. 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 [].

## 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
```

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
6
```

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)) = {[]};
```

## See also

Operator {}, end, reshape, variable assignment, operator [], subref, 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; 2 5 , 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 4]
 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}
v{index}
v{index1, index2, ...}
v{index} = expr
fun(...,v{:},...)
```

**Description**

A pair of braces is used to define a list or a cell array given by its elements. 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.

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+3j
x =
  {1,2+3j,real 2x2}
x{3} = {2}
x =
  {1,2+3j,list}
x{end+1} = 123
x =
  {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

```

## 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 exist in an assignment); i.e. complicated assignments like `a{2}.f{3}(2,5)=3` are accepted.

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.

## Syntax

```
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 numerical 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]
  13
  29
[3 4] * 2
  6 8

```

**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.

If the second operand is 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

$x \setminus y$  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;  $M1 \setminus M2$  is equivalent to  $\text{inv}(M1) * M2$ .

`mldivide(x,y)` is equivalent to  $x \setminus y$ . 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.

`\divide(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.

**Syntax**

```
x ^ y
M ^ k
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 an integer;
- 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

```

**See also**

operator .^, expm

**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  $x.^y$ . It can be used to redefine this operator for objects.

**Examples**

```

[1,2;3,4].^2
1 4
9 16
[1,2,3].^[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.

**Syntax**

```
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 0===-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 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 >=, operator ==, operator ~==, strcmp

**Operator  $\sim=$** 

Inequality.

**Syntax**

```
x  $\sim=$  y
ne(x, y)
```

**Description**

$x \sim= y$  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  $x \sim= y$ . It can be used to redefine this operator for objects.

**Example**

```
1  $\sim=$  1
  false
inf  $\sim=$  inf
  false
nan  $\sim=$  nan
  true
[1,2,3]  $\sim=$  [1,3,3]
  F T F
```

**See also**

operator ==, operator <, operator <=, operator >, operator >=, operator ==, operator  $\sim=$ , strcmp

**Operator  $\sim==$** 

Object inequality.

**Syntax**

```
a  $\sim==$  b
unsame(a, b)
```

**Description**

a `~==` 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~==nan` is false and `0~==-0` is true. Contrary to the inequality operator, `~==` returns a single boolean even if its operands are arrays.

`unsame(a, b)` is equivalent to `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 `<=`, operator `>`, operator `>=`, `strcmp`

**Operator <**

Less than.

**Syntax**

```
x < y
lt(x, y)
```

**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**

```
[2,3,4] < [2,4,2]
  F T F
```

**See also**

operator ==, operator ~=, operator <=, operator >, operator >=

**Operator >**

Greater than.

**Syntax**

```
x > y
gt(x, y)
```

**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 >=

**Operator <=**

Less or equal to.

**Syntax**

```
x <= y
le(x, y)
```

**Description**

$x \leq y$  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 element-wise and the result has the same size.

$\text{le}(x, y)$  is equivalent to  $x \leq y$ . 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 >=

**Operator >=**

Greater or equal to.

**Syntax**

```
x >= y
ge(x, y)
```

**Description**

$x \geq y$  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 element-wise and the result has the same size.

$\text{ge}(x, y)$  is equivalent to  $x \geq y$ . 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 >

## 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.

### Examples

```
~true
false
~[1,0,3,false]
F T F T
```

### See also

operator ~=, bitcmp

## 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 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 `&`, 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 || 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 or matrices; their number of rows must be the same, unless one of them is an empty matrix.

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 re-define this operator for objects.

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 or matrices; their number of columns must be the same, unless one of them is an empty matrix.

Outside brackets, the comma is used to separate statements; they lose any meaning between parenthesis and give a syntax error.

vertcat(M1,M2) is equivalent to [M1;M2]. It can be used to re-define 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((z - x1)/(x2 - x1)) <= 1
  add z to the vector
  z = z + step
end
```

This algorithm is robust enough to stop even if `x2` is not on the complex straight line defined by `x1` and `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 `quad`, but it may also be stored in lists, cell arrays, or

structures. A reference cannot be cast to a double (unlike characters or logical values), nor can it be stored in an array.

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.

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:

```
quad(@sin, 0, pi)
    2
```

Anonymous function:

```
a = 2;
fun = @(x) sin(a * x);
fun(3)
    -0.2794
quad(fun, 0, 2)
    0.8268
```

Without anonymous function, parameter `a` should be passed as an additional argument after all the input arguments defined for `quad`, including those which are optional when parameters are missing:

```
quad(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)
```

**See also**

fun2str, str2fun, inline, feval, apply

## 3.15 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

**acosh**

Inverse hyperbolic cosine.

**Syntax**

$$y = \operatorname{acosh}(x)$$
**Description**

$\operatorname{acosh}(x)$  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**

$$y = \operatorname{acot}(x)$$
**Description**

$\operatorname{acot}(x)$  gives the inverse cotangent of  $x$ , which is complex if  $x$  is.

**See also**

cot, acoth, cos

**acoth**

Inverse hyperbolic cotangent.

**Syntax**

$$y = \operatorname{acoth}(x)$$

**Description**

$\operatorname{acoth}(x)$  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**

$$y = \operatorname{acsc}(x)$$

**Description**

$\operatorname{acsc}(x)$  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**

$$y = \operatorname{acsch}(x)$$

**Description**

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**
$$y = \operatorname{asech}(x)$$
**Description**

$\operatorname{asech}(x)$  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**
$$y = \operatorname{asin}(x)$$
**Description**

$\operatorname{asin}(x)$  gives the arc sine of  $x$ , which is complex if  $x$  is complex or if  $\operatorname{abs}(x) > 1$ .

**Examples**

```
asin(0.5)
0.5236
asin(2)
1.5708-1.317j
```

**See also**

sin, acos, asinh

## asinh

Inverse hyperbolic sine.

### Syntax

```
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.

**Syntax**

```
phi = atan2(y,x)
```

**Description**

$\text{atan2}(y, x)$  gives the direction of a point given by its Cartesian coordinates  $x$  and  $y$ . Imaginary component of complex numbers is ignored.  $\text{atan2}(y, x)$  is equivalent to  $\text{atan}(y/x)$  if  $x > 0$ .

**Examples**

```
atan2(1, 1)
0.7854
atan2(-1, -1)
-2.3562
atan2(0, 0)
0
```

**See also**

`atan`, `angle`

**atanh**

Inverse hyperbolic tangent.

**Syntax**

```
y = atanh(x)
```

**Description**

$\text{atan}(x)$  gives the inverse hyperbolic tangent of  $x$ , which is complex if  $x$  is complex or if  $\text{abs}(x) > 1$ .

**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**

```
beta(1,2)
0.5
```

**See also**

gamma, beta<sub>ln</sub>, beta<sub>inc</sub>

**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**

```
betainc(0.2,1,2)
0.36
```

**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

**cast**

Type conversion.

**Syntax**

```
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	Parameters
Beta	beta	a and b
Cauchy	cauchy	a and b
$\chi$	chi	deg. of freedom $\nu$
$\chi^2$	chi2	deg. of freedom $\nu$
$\gamma$	gamma	shape $\alpha$ and $\lambda$
exponential	exp	mean
F	f	deg. of freedom $\nu_1$ and $\nu_2$
half-normal	half-normal	$\theta$
Laplace	laplace	mean and scale
lognormal	logn	mean (0) and st. dev. (1)
normal	norm	mean (0) and st. dev. (1)
Rayleigh	rayl	b
Student's T	t	deg. of freedom $\nu$
uniform	unif	limits of the range (0 and 1)
Weibull	weib	a and b

### See also

pdf, icdf, erf

### ceil

Rounding towards +infinity.

### Syntax

```
y = ceil(x)
```

### Description

`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.

### Examples

```
ceil(2.3)
3
ceil(-2.3)
-2
ceil(2.3-4.5j)
3-4j
```

**See also**

floor, fix, round

**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

**cosh**

Hyperbolic cosine.

**Syntax**
$$y = \cosh(x)$$
**Description**

$\cosh(x)$  gives the hyperbolic cosine of  $x$ , which is complex if  $x$  is complex.

**Example**

```
cosh([0, 1+2j])  
1 -0.6421+1.0686j
```

**See also**

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**

$y = \text{csch}(x)$

**Description**

`csch(x)` gives the hyperbolic cosecant of  $x$ , which is complex if  $x$  is.

**See also**

`acsch`, `csc`, `sinh`

**diln**

Dilogarithm.

**Syntax**

$y = \text{diln}(x)$

**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

$$\text{diln}(x) = \int_1^x \frac{\log(t)}{t-1} dt$$

**Example**

```

diln([0.2, 0.7, 10])
-1.0748 -0.3261 3.9507

```

## 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`.

**Example**

```

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

$$E(\varphi|m) = \int_0^\varphi \sqrt{1 - m \sin^2 t} dt$$

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

$$F(\varphi|m) = \int_0^\varphi \frac{dt}{\sqrt{1 - m \sin^2 t}}$$

Complete elliptic integrals of first and second kinds, with  $\text{phi}=\text{pi}/2$ , can be obtained with `ellipke`.

**See also**

`ellipe`, `ellipke`, `ellipam`

**ellipj**

Jacobi elliptic functions.

**Syntax**

```
(sn, cn, dn) = ellipj(u, m)
(sn, cn, dn) = ellipj(u, m, tol)
```

**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`):

$$u = F(\varphi|m)$$

$$\text{sn}(u|m) = \sin(\varphi)$$

$$\text{cn}(u|m) = \cos(\varphi)$$

$$\text{dn}(u|m) = \sqrt{1 - m \sin^2 \varphi}$$

`ellipj(u,m,tol)` uses tolerance `tol`; the default tolerance is `eps`.

**Examples**

```
(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

$$F(m) = \int_0^{\pi/2} \frac{dt}{\sqrt{1 - m \sin^2 t}}$$

$$E(m) = \int_0^{\pi/2} \sqrt{1 - m \sin^2 t} dt$$

Parameter  $m$  must be in  $[0,1]$ .

ellipke(m, tol) uses tolerance tol; the default tolerance is eps.

**Example**

```
(K, E) = ellipke(0.3)
K =
  1.7139
E =
  1.4454
```

**See also**

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  $1+\text{eps}$  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  $1\text{single}+e$  can be distinguished from  $1\text{single}$ . `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
0
```

**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

$$\text{erf}(x) = \frac{2}{\sqrt{\pi}} \int_0^x e^{-t^2} dt$$

**Example**

```
erf(1)
0.8427
```

**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

$$\text{erfc}(x) = 1 - \text{erf}(x) = \frac{2}{\sqrt{\pi}} \int_x^\infty e^{-t^2} dt$$

**Example**

```
erfc(1)
0.1573
```

**See also**

erf, erfinv

**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, erfc

**exp**

Exponential.

**Syntax**

$$y = \exp(x)$$
**Description**

$\exp(x)$  is the exponential of  $x$ , i.e.  $2.7182818284590446\dots^x$ .

**Example**

```
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

## 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.

### Examples

```
fix(2.3)
    2
fix(-2.6)
   -2
```

**See also**

floor, ceil, round

**floor**

Rounding towards -infinity.

**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

**gamma**

Gamma function.

**Syntax**

`y = gamma(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

$$\Gamma(x) = \int_0^{\infty} t^{x-1} e^{-t} dt$$

For positive integer values,  $\Gamma(n) = (n - 1)!$ .

**Examples**

```
gamma(5)
24
gamma(-3)
inf
gamma(-3.5)
0.2701
```

**See also**

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

$$\text{gammainc}(x, a) = \frac{1}{\Gamma(a)} \int_0^x t^{a-1} e^{-t} dt$$

**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.

**Examples**

```
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.

## Example

```
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	Parameters
Beta	beta	a and b
$\chi^2$	chi2	deg. of freedom $\nu$
$\gamma$	gamma	shape $\alpha$ 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 $\nu$
uniform	unif	limits of the range (0 and 1)

**See also**

`cdf`, `pdf`

**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

```

**See also**

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

## 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
```

### See also

`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
```

### **See also**

`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`

**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, isvector

**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.

**Examples**

```
isscalar(2)
    true
isscalar([1, 2, 5])
    false
```

**See also**

isnumeric, isvector, 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, size, ndims, length

**lcm**

Least common multiple.

**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**

$\log_{10}$ ,  $\log_2$ ,  $\log_{1p}$ ,  $\text{reallog}$ ,  $\text{exp}$

**log10**

Decimal logarithm.

**Syntax**

$$y = \log_{10}(x)$$
**Description**

$\log_{10}(x)$  gives the decimal logarithm of  $x$ , defined by  $\log_{10}(x) = \log(x)/\log(10)$ . 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**

$\log$ ,  $\log_2$

**log1p**

Logarithm of  $x$  plus one.

**Syntax**

$$y = \log_{1p}(x)$$

**Description**

$\log_{1p}(x)$  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**

$$y = \log_2(x)$$
**Description**

$\log_2(x)$  gives the base 2 logarithm of  $x$ , defined as  $\log_2(x) = \log(x) / \log(2)$ . 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**

$$m = \text{mod}(x, y)$$

## Description

$\text{mod}(x, y)$  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
```

## See also

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,  $0/0$ ,  $\text{inf-inf}$  and  $0*\text{inf}$  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

**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`

**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

**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

## 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 `.`<sup>^</sup>, `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

`sqrt`, `reallog`, `realpow`

## 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)
 -3
```

**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

**sign**

Sign of a real number or direction of a complex number.

**Syntax**

```
s = sign(x)
z2 = sign(z1)
```

**Description**

With a real argument,  $\text{sign}(x)$  is 1 if  $x > 0$ , 0 if  $x == 0$ , or -1 if  $x < 0$ . With a complex argument,  $\text{sign}(z1)$  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**

$\text{sec}(x)$  gives the secant of  $x$ , which is complex if  $x$  is.

**See also**

asec, sech, cos

**sech**

Hyperbolic secant.

**Syntax**

$$y = \operatorname{sech}(x)$$

**Description**

$\operatorname{acot}(x)$  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**

$$\begin{array}{l} \sin(2) \\ 0.9093 \end{array}$$

**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  $x \neq 0$  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

**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**

$Y = \text{swapbytes}(X)$

**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 numerical type. `swapbytes` is its own inverse for real numbers.

**Example**

```
swapbytes(uint32)
16777216uint32
```

**See also**

typecast, cast

**tan**

Tangent.

**Syntax**

$y = \text{tan}(x)$

**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 is 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(uint32, 'uint8')
  1x4 uint8 array
   0   0   0   1
typecast(pi, 'uint8')
  1x8 uint8 array
  64   9  33 251  84  68  45  24

```

**See also**

swapbytes, bwrite, sread, cast

## 3.16 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  $T \backslash A * T = B$ .

### 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:

$$A'X + XA - XBB'X + Q = 0$$

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

$$L = \text{eig}(A - BK)$$

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)
X =
    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 + Q
    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:

$$M_1 = M_2' M_2$$

where 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)
```

**Description**

`conv(v1,v2)` convolves the vectors `v1` and `v2`, giving a vector whose length is `length(v1)+length(v2)-1`. 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.

**Example**

```
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
```

**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 (nl1,nc1)=size(M1) and (nl2,nc2)=size(M2). With kind='full' (default value), the result M has nl1+nl2-1 lines and nc1+nc2-1 columns. With kind='same', the result M has nl1 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 nl1-nl2+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, 0)
M = cov(data, 1)

```

**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 real and symmetric, even if `data` is complex. The diagonal is the variance of each column of `data`. `cov(data,0)` is the same as `cov(data)`.

`cov(data,1)` returns the  $m$ -by- $m$  covariance matrix of the  $n$ -by- $m$  matrix `data` which contains the whole population.

**Example**

```

cov([1,2;2,4;3,5])
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 column 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`

## cumprod

Cumulative products.

### Syntax

```
M2 = cumprod(M1)
M2 = cumprod(M1,dim)
```

### 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  $k \leq i$ . `cumprod(M1,dim)` operates along the dimension `dim` (column-wise if `dim` is 1, row-wise if `dim` is 2).

### 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`

**cumsum**

Cumulative sums.

**Syntax**

```
M2 = cumsum(M1)
M2 = cumsum(M1,dim)
```

**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  $k \leq i$ . `cumsum(M1,dim)` operates along the dimension `dim` (column-wise if `dim` is 1, row-wise if `dim` is 2).

**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
```

**See also**

`sum`, `diff`, `cumprod`

**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 discrete-time algebraic Riccati equation:

$$X = A'XA - A'XB(B'XB + I)^{-1}B'XA + Q$$

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:

$$X = A'XA - A'XB(B'XB + R)^{-1}B'XA + Q$$

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

$$L = \text{eig}(A - BK)$$

### Example

```
A = [-4,2;1,2];
B = [0;1];
C = [2,-1];
Q = C' * C;
R = 5;
(X, L, K) = dare(A, B, Q, R)
X =
  2327.9552  -1047.113
 -1047.113   496.0624
L =
  -0.2315
   0.431
K =
   9.3492  -2.1995
-X + A'*X*A - A'*X*B/(B'*X*B+R)*B'*X*A + Q
  1.0332e-9  -4.6384e-10
 -4.8931e-10  2.2101e-10
```

### See also

`care`

### deconv

Deconvolution or polynomial division.

### Syntax

```
q = deconv(a,b)
(q,r) = deconv(a,b)
```

**Description**

$(q, r) = \text{deconv}(a, b)$  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  $b$ .  $\text{deconv}$  is the inverse of  $\text{conv}$ :  $a = \text{addpol}(\text{conv}(b, q), r)$ .

**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**

$\text{det}(M)$  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])
    0
```

**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.

**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

```

**See also**

cross, operator \*, det

**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**

```
eig([1,2;3,4])
-0.3723
 5.3723
(V,D) = eig([1,2;2,1])
V =
 0.7071 0.7071
-0.7071 0.7071
D =
-1 0
 0 3
[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 `fft2(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:

```
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.

## Syntax

```

Y = funm(X, fun)
(Y, err) = funm(X, fun)

```

## Description

`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 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, inline('(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, logm, sqrtm, schur

**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,  $\text{ifft}(d) = \text{conj}(\text{fft}(d))/\text{length}(d)$ . 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`.

**See also**

`fftn`, `ifft`, `ifft2`

**hess**

Hessenberg reduction.

**Syntax**

```
(P,H) = hess(A)
H = hess(A)
```

**Description**

`hess(A)` reduces the square matrix `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])
P =
  1          0          0
  0      -0.4961  -0.8682
  0      -0.8682   0.4961
H =
  1      -3.597  -0.2481
 -8.0623 14.0462   2.8308
  0       0.8308 -4.6154e-2
P*H*P'
ans =
  1      2      3
  4      5      6
  7      8      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  $M2 * M1 = M1 * M2 = \text{eye}(\text{size}(M1))$ . `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
```

**See also**

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.

## Syntax

```
x = linprog(c, A, b)
x = linprog(c, A, b, xlb, xub)
```

## Description

`linprog(c,A,b)` solves the following linear programming problem:

$$\begin{aligned} \min & cx \\ \text{s.t.} & Ax \leq b \end{aligned}$$

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:

$$\begin{aligned} \min & cx \\ \text{s.t.} & Ax \leq b \\ & x \geq x_{lb} \\ & x \leq x_{ub} \end{aligned}$$

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  $x + y \leq 9$ ,  $3x + y \leq 18$ ,  $x \leq 7$ , and  $y \leq 6$ :

```
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`, `sqrtn`, `funm`, `schur`, `log`

## lu

LU decomposition.

## Syntax

$(L, U, P) = \text{lu}(A)$

$(L2, U) = \text{lu}(A)$

$Y = \text{lu}(A)$

## Description

With three output arguments,  $\text{lu}(A)$  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  $A$  is an  $m$ -by- $n$  matrix,  $L$  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,  $\text{lu}(A)$  permutes the lower triangular matrix and gives  $L2=P'*L$ , such that  $A=L2*U$ .

With a single output argument,  $\text{lu}$  gives  $Y=L+U-\text{eye}(n)$ .

## Example

```
X = [1,2,3;4,5,6;7,8,8];
```

```
(L,U,P) = lu(X)
```

```
L =
```

```
 1     0     0
 0.143 1     0
 0.571 0.5   1
```

```
U =
```

```
 7     8     8
 0     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
6

```

**See also**

`cov`, `std`, `var`, `sum`, `prod`

**min**

Minimum value of a vector or of two arguments.

**Syntax**

```

x = min(v)
(v,ind) = min(v)
v = min(M,[],dim)
(v,ind) = min(M,[],dim)
M3 = min(M1,M2)

```

**Description**

`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

```
(mx,ix) = min([1,3,2,5,8,7])
mx =
  1
ix =
  1
min([1,3;5,nan], [], 2)
  1
  5
min([1,3;5,nan], 2)
  1 2
  2 2
```

### 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	$\sqrt{\text{sum}(\text{abs}(v).^2)}$	largest singular value (induced 2-norm)
1	$\text{sum}(\text{abs}(V))$	largest column sum of abs
inf or 'inf'	$\text{max}(\text{abs}(v))$	largest row sum of abs
-inf	$\text{min}(\text{abs}(v))$	largest row sum of abs
p	$\text{sum}(\text{abs}(V).^p)^{1/p}$	invalid
'fro'	$\sqrt{\text{sum}(\text{abs}(v).^2)}$	$\sqrt{\text{sum}(\text{diag}(M'*M))}$

**Examples**

```
norm([3,4])
5
norm([2,5;9,3])
10.2194
norm([2,5;9,3],1)
11
```

**See also**

abs, hypot, svd

**null**

Null space.

**Syntax**

```
Z = null(A)
```

**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  $n - \text{rank}(A)$  columns, which are the last right singular values of  $A$  (that is, those corresponding to the negligible singular values).

**Example**

```
null([1,2,3;1,2,4;1,2,5])
-0.8944
 0.4472
 8.0581e-17
```

**See also**

`svd`, `orth`

**orth**

Orthogonalization.

**Syntax**

```
Q = orth(A)
```

**Description**

`orth(A)` returns a matrix  $Q$  whose columns are an orthonormal basis for the range of those of matrix  $A$ .  $Q$  has  $\text{rank}(A)$  columns, which are the first left singular vectors of  $A$  (that is, those corresponding to the largest singular values).

**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,e)
```

### 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. To 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.

If M1 is a full-rank matrix with more rows than columns, pinv returns the least-square solution  $pinv(M1) * y = (M1' * M1) \backslash M1' * y$  of the over-determined system  $M1 * x = y$ .

### Examples

```
pinv([1,2;3,4])
    -2    1
     1.5 -0.5
M2 = pinv([1;2])
M2 =
    0.2 0.4
[1;2] * M2 * [1;2]
     1
     2
M2 * [1;2] * M2
    0.2 0.4
```

### 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  $\det(x \cdot \text{eye}(\text{size}(M)) - M)$  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
```

**See also**

`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 their 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
```

**See also**

`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 their 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     4     5
Y = polyint([1, 2, 3, 4, 5], 10)
Y =
    0.2    0.5    1     2     5    10

```

**See also**

polyder, polyval, poly, addpol, conv

**polyval**

Numerical value of a polynomial evaluated at some point.

**Syntax**

```
y = polyval(pol, x)
```

**Description**

polyval(pol,x) evaluates the polynomial pol at x, which can be a scalar or a matrix of arbitrary size. 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
```

**See also**

`sum`, `mean`, operator `*`

**qr**

QR decomposition.

**Syntax**

```
(Q, R, E) = qr(A)
(Q, R) = qr(A)
(Qe, Re, e) = qr(A, false)
(Qe, 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  $A \cdot E = Q \cdot R$ . With two output arguments, `qr(A)` computes the QR decomposition without pivoting, such that  $A = Q \cdot R$ .

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:  $A(:, e) = Q \cdot R$ .

## Example

```
(Q,R) = qr([1,2;3,4;5,6])
Q =
-0.169      0.8971    0.4082
-0.5071     0.276   -0.8165
-0.8452    -0.345    0.4082
R =
-5.9161    -7.4374
      0      0.8281
      0      0
(Q,R) = qr([1,2;3,4;5,6],false)
Q =
 0.169      0.8971
 0.5071     0.276
 0.8452    -0.345
R =
 5.9161     7.4374
 0         0.8281
```

## See also

`lu`, `schur`, `hess`, `svd`

## rank

Rank of a matrix.

## Syntax

```
x = rank(M)
x = rank(M,e)
```

## 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.

## 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.

**Syntax**

```

(U,T) = schur(A)
T = schur(A)
(U,T) = schur(A, 'c')
T = schur(A, 'c')

```

**Description**

`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'`.

**Example**

```

(U,T) = schur([1,2;3,4])
U =
-0.8246 -0.5658
0.5658 -0.8246
T =
-0.3723 -1
0 5.3723
eig([1,2;3,4])
ans =
-0.3723
5.3723

```

```

T = schur([1,0,0;0,1,2;0,-3,1])
T =
    1     0     0
    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 - 2.4495j

```

**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  $\text{sqrtm}(X)^2=X$ .  $X$  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  $\text{norm}(\text{sqrtm}(X)^2-X)/\text{norm}(X)$ .

**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`, `sqrt`

**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  $\text{length}(v)-1$ . With a second argument, `std(v,p)` normalizes by  $\text{length}(v)-1$  if  $p$  is true, or by  $\text{length}(v)$  if  $p$  is false.

`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  $\text{dim}$ .

**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.

### Syntax

```
s = svd(M)
(U,S,V) = svd(M)
(U,S,V) = svd(M, false)
```

### Description

The singular value decomposition  $(U, S, V) = \text{svd}(M)$  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.  $(U, S, V) = \text{svd}(M, \text{false})$  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
$m$ by $n, m \leq n$	$m$ by $m$	$m$ by $m$	$n$ by $m$
$m$ by $n, m > n$	$m$ by $n$	$n$ by $n$	$n$ by $n$

$\text{svd}(M, \text{true})$  produces the same result as  $\text{svd}(M)$ .

With one output argument,  $s = \text{svd}(M)$  returns the vector of singular values  $s = \text{diag}(S)$ .

The singular values of  $M$  can also be computed with  $s = \text{sqrt}(\text{eig}(M'*M))$ , but  $\text{svd}$  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

**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

## 3.17 Array Functions

**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.

### Syntax

```
C = cell(n)
C = cell(n1,n2,...)
C = cell([n1,n2,...])
```

### Description

`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 = cell(fun, C, ...)
```

### 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`.

`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`

## 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 0 3
```

```
diag(1:3,1)
 0 1 0 0
 0 0 2 0
 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
 7
```

### See also

`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

```

**See also**

ones, zeros, diag

**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
    4
[s1,s2] = find([1.2,0;0,3.6])
s1 =
    1
    2
s2 =
    1
    2
[s1,s2,x] = find([1.2,0;0,3.6])
s1 =
    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
    5

```

**See also**

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
```

**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

**interp**

Interpolation.

**Syntax**

```
Vi = interp(x1, ..., xn, V, xi1, ..., xin)
Vi = interp(x1, ..., xn, V, xi1, ..., xin, method)
```

**Description**

`interp(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):

<b>Argument</b>	<b>Meaning</b>
'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.

**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;

```

## 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 numerical, 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**

```

a = {'a', 'bc', 'bbb', 'de'};
b = {'z', 'bc', 'aa', 'bbb'};
(c, ia, ib) = intersect(a, b)
c =
    {'bbb', 'bc'}
ia =
    3 2
ib =
    4 2
a(ia)
    {'bbb', 'bc'}
b(ib)
    {'bbb', 'bc'}

```

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

**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 `perm(i)` in A becomes dimension `i` in the result.

**Examples**

```

size(permute(rand(3,4,5), [2,3,1]))
    5 3 4

```

**See also**

permute, ndims, squeeze

**isempty**

Test for empty matrices or empty lists.

**Syntax**

```
b = isempty(M)
b = isempty(list)
```

**Description**

`isempty(obj)` gives true if `obj` is the empty array `[]`, the empty string `''`, or the empty list `{}`, and false otherwise.

**Examples**

```
isempty([])
    true
isempty(0)
    false
isempty('')
    true
isempty({})
    true
isempty({{}})
    false
```

**See also**

size, length

**iscell**

Test for cell arrays.

**Syntax**

```
b = iscell(X)
```

**Description**

`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)
```

**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 numerical array or a cell array, matching type of set `s`.

**Example**

```

s = {'a', 'bc', 'bbb', 'de'};
m = {'d', 'a', 'x'; 'de', 'a', 'z'};
b = ismember(m, s)
b =
  F T F
  T T F

```

**See also**

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`

**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
 1  2  3  4  5
 1  2  3  4  5
Y =
 2  2  2  2  2
 3  3  3  3  3
 4  4  4  4  4
Z = atan2(X, Y)
Z =
 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, ..., Xn) = ndgrid(x1, ..., xn)
(X1, ..., 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
```

**See also**

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.

## Syntax

```
n = nnz(A)
```

## Description

`nnz(A)` returns the number of nonzero elements of array A.

## 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 dimensions `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 numerical 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]}
```

**See also**

`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'})
2
```

## See also

size, length

## ones

Array of ones.

## Syntax

```
A = ones(n)
A = ones(n1, n2, ...)
A = ones([n1, n2, ...])
A = ones(..., type)
```

## Description

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).

## Example

```
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 `i` in `A` becomes dimension `perm(i)` in the result.

**Examples**

```
size(permute(rand(3,4,5), [2,3,1]))
4 5 3
```

**See also**

`ndims`, `squeeze`, `ipermute`, `num2cell`

**rand**

Uniformly-distributed random number.

**Syntax**

```
x = rand
M = rand(n)
M = rand(n1, n2, ...)
M = rand([n1, n2, ...])
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 current implementation is based on a scalar 64-bit seed, which theoretically allows  $2^{64}$  different numbers. This seed can be set with the arguments `rand('seed', s)`, where `s` is a scalar or a vector of two components. `rand('seed', s)` returns the empty array `[]` as output argument. To discard it, the statement should be followed by a semicolon.

`rand(n)`, `rand(n1, n2, ...)` and `rand([n1, n2, ...])` return an `n`-by-`n` square matrix or an array of arbitrary size whose elements are pseudo-random numbers uniformly distributed between 0 and 1.

**Examples**

```

rand
  0.2361
rand(1, 3)
  0.6679 0.8195 0.2786
rand('seed', 0);
rand
  0.2361

```

**See also**

randn

**randn**

Normally-distributed random number

**Syntax**

```

x = randn
M = randn(n)
M = randn(n1, n2, ...)
M = randn([n1, n2, ...])
randn('seed', s);

```

**Description**

randn builds a scalar pseudo-random number chosen from a normal distribution with zero mean and unit variance. The current implementation is based on a scalar 64-bit seed, which theoretically allows  $2^{64}$  different numbers. This seed can be set with the arguments `randn('seed', s)`, where `s` is a scalar or a vector of two components. The seed is not the same as the seed of `rand`. `randn('seed', s)` returns the empty array `[]` as output argument. To discard it, the statement should be followed by a semicolon.

`randn(n)`, `randn(n1, n2, ...)` and `randn([n1, n2, ...])` return an `n`-by-`n` square matrix or an array of arbitrary size whose elements are pseudo-random numbers chosen from a normal distribution.

**Examples**

```

randn
  1.5927
randn(1, 3)
  0.7856 0.6489 -0.8141
randn('seed', 0);

```

```
randn
    1.5927
```

**See also**

rand

**repmat**

Replicate an array.

**Syntax**

```
A2 = repmat(A1, n)
A2 = repmat(A1, m, n)
A2 = repmat(A1, [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. The number of copies is `m` in the vertical direction, and `n` in the horizontal direction. The type of the first argument (number, character or logical value) is preserved. With a vector as second argument, the array can be replicated along more than two dimensions.

**Examples**

```
repmat([1,2;3,4],1,2)
    1 2 1 2
    3 4 3 4
repmat('abc',3)
    abcabcabc
    abcabcabc
    abcabcabc
```

**See also**

`zeros`, `ones`, operator `:`, `kron`, `replist`

**reshape**

Rearrange the elements of an array to change its shape.

**Syntax**

```
A2 = reshape(A1)
A2 = reshape(A1, n1, n2, ...)
A2 = reshape(A1, [n1, n2, ...])
```

**Description**

`reshape(A1)` gives a column vector with all the elements of array `A1`, which is read row-wise. 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 line-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.

**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 ()

**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
```

**See also**

`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 numerical, 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  $(c, ia) = \text{setdiff}(a, b)$ , 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

$\text{setxor}(a, b)$  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 numerical, 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  $(c, ia, ib) = \text{setxor}(a, b)$ , then  $c$  is the union of  $a(ia)$  and  $b(ib)$ .

**Example**

```

a = {'a','bc','bbb','de'};
b = {'z','bc','aa','bbb'};
(c, ia, ib) = setxor(a, b)
c =
    {'a','aa','de','z'}
ia =
    1 4
ib =
    3 1
union(a(ia),b(ib))
    {'a','aa','de','z'}

```

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.

**Syntax**

```

(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)

```

**Description**

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  1
  6  4  2
  nan 5  3
sort([2,5,3;nan,4,2;6,1,1], 'd')
  6  5  3
  2  4  2
  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.

**Syntax**

```
ind = sub2ind(size, i, j)
```

**Description**

`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.

**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

`triu(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
```

### See also

`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 numerical, 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  $(c, ia, ib) = \text{union}(a, b)$ , then elements of `c` are the elements of `a(ia)` or `b(ib)`; the intersection of `a(ia)` and `b(ib)` is empty.

**Example**

```
a = {'a','bc','bbb','de'};
b = {'z','bc','aa','bbb'};
(c, ia, ib) = union(a, b)
c =
    {'a','aa','bbb','bc','de','z'}
ia =
    1 3 2 4
ib =
    3 1
a(ia)
    {'a','bbb','bc','de'}
b(ib)
    {'aa','z'}
```

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`

## 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 0
```

### See also

ones, cell, eye, rand, randn, repmat

## 3.18 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**

delaulnay, 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:

<b>Argument</b>	<b>Meaning</b>
'0' or 'nearest'	nearest value
'1' or 'linear'	linear

**See also**

delaulnay, 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

## See also

`deLaunay`, `tsearchn`, `griddata`, `interp`

## 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

**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);
```

**See also**

deleunayn, voronoin

**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
```

**See also**

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  $n = \text{size}(x, 2)$ . 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

## 3.19 Integer Functions

<b>uint8 uint16 uint32 uint64 int8 int16 int32 int64</b>
--

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.

`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
int8([50:50:400])
 1x8 int8 array
 50 100 -106 -56 -6 44 94 -112

```

**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`, `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
```

**See also**

uint8 and related functions.

## 3.20 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  $\mathbb{R}^n$ .

### 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  $\text{fun}(x)$  is minimized, starting from point  $x_0$ .

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  $\text{fun}(x)$  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  $x_0$  (the "simplex"). Then at each iteration, either vertex  $x_i$  with the maximum value  $\text{fun}(x_i)$  is moved to decrease it with a reflexion-expansion, a reflexion, or a contraction; or the simplex is shrunk 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  $x e^{-x^2 y^2} x y - 0.1 x^2$  The function if 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  $R^2$ :

```
mfun = @(X) -(X(1)*exp(-(X(1)*X(2))^2)*X(1)*X(2)-0.1*X(1)^2);
fminsearch(mfun, [1, 2])
2.1444 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):

```
mfunc = @(X) ...
    X(1) < 1 ...
    ? -(X(1)*exp(-(X(1)*X(2))^2)*X(1)*X(2) - 0.1*X(1)^2) ...
    : inf;
fminsearch(mfunc, [1, 2])
1 0.7071
```

## See also

optimset, fminbnd, 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
```

**See also**

`optimset`, `fminsearch`, `inline`, `operator @`, `roots`

**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) = ode45(fun,[t0,tend],y0)
(t,y) = ode45(fun,[t0,tend],y0,options)
(t,y) = ode45(fun,[t0,tend],y0,options,...)
```

## 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.)

Additional input arguments of `ode45` are given as additional input arguments to the function specified by `fun`. They permit to parameterize the ODE.

## Example

Let us integrate the following ordinary differential equation (Van Der Pol equation), parameterized by  $\mu$ :

$$x'' = \mu(1 - x^2)x' - x$$

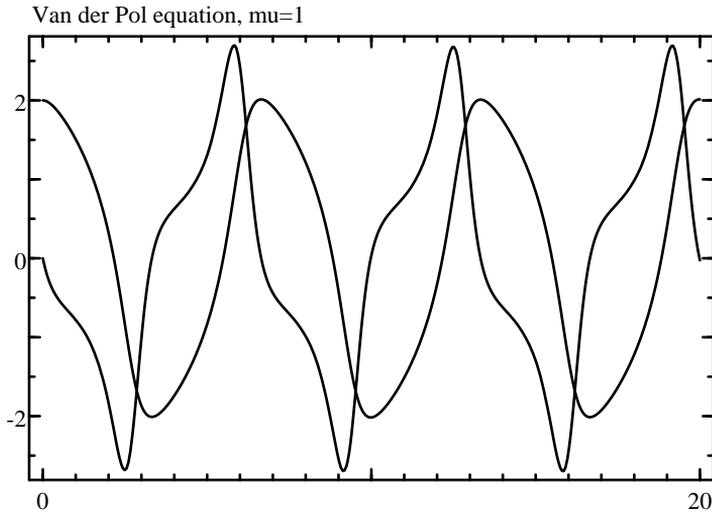
Let  $y_1 = x$  and  $y_2 = x'$ ; their derivatives are

$$\begin{aligned} y_1' &= y_2 \\ y_2' &= \mu(1 - y_1^2)y_2 - y_1 \end{aligned}$$

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. 3.1):



**Figure 3.1** Van der Pol equation with  $\mu = 1$  integrated with ode45

```
(t,y)=ode45(@f,[0,10],[2;0],[],1);
```

The `plot` command expects traces along the second dimension; consequently, the result of `ode45` should be transposed.

```
plot(t', y');
```

**See also**

`odeset`, `quad`, `inline`, `operator @`, `expm`

**odeset**

Options for ordinary differential equation integration.

**Syntax**

```
options = odeset  
options = odeset(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. 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.

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 "automatic"):

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
<code>AbsTol</code>	<code>1e-6</code>	maximum absolute error
<code>Events</code>	<code>[]</code> (none)	state-based event function
<code>EventTime</code>	<code>[]</code> (none)	time-based event function
<code>InitialStep</code>	<code>[]</code> ( <code>10*MinStep</code> )	initial time step
<code>MaxStep</code>	<code>[]</code> (time range/10)	maximum time step
<code>MinStep</code>	<code>[]</code> (time range/1e6)	minimum time step
<code>NormControl</code>	<code>false</code>	error control on state norm
<code>OnEvent</code>	<code>[]</code> (none)	event function
<code>OutputFcn</code>	<code>[]</code> (none)	output function
<code>PreArg</code>	<code>{}</code>	list of prepended input arguments
<code>Refine</code>	<code>[]</code> (1, 4 for <code>ode45</code> )	refinement factor
<code>RelTol</code>	<code>1e-3</code>	maximum relative error
<code>Stats</code>	<code>false</code>	statistics display

### ***Time steps and output***

Several options control how the time step is tuned during the numerical 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  $t_n$  is the time of the new samples,  $y_n$  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,  $t_n$  is a row vector and  $y_n$  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  $t_s$ , and the second column gives the offset  $t_o$ . Non-repeating events are specified with an infinite time step  $t_s$ . Events occur at time  $t = t_o + k * t_s$ , 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

```
function (value, isterminal, direction) ...
    = eventsfun(t, y, ...)
```

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 OnEvent 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.

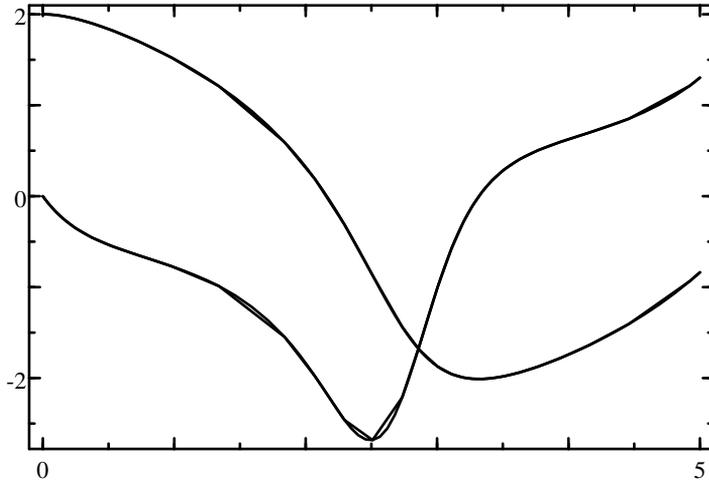
### **Additional arguments**

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
```



**Figure 3.2** Van der Pol equation with Refine set to 1 and 4

```
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. 3.2). 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, y) = 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(y)
    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');
```

### **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:

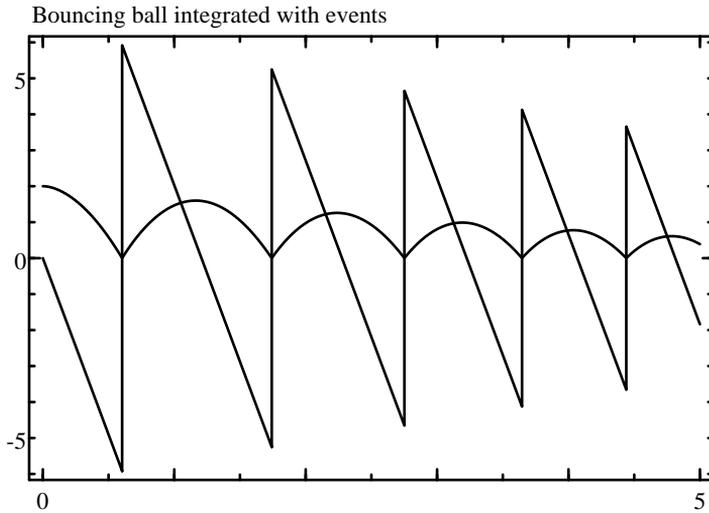
$$y_n = [0; -\text{damping} \cdot y(2)];$$

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)];
```

Ball state is integrated during 5 s (see Fig. 3.3) with

```
opt = odeset('Events', @ballevents, ...
    'OnEvent', @ballonevent);
(t, y) = ode45(@ballfun, [0, 5], [2; 0], opt, 1);
plot(t', y');
```



**Figure 3.3** Bouncing ball integrated with events

**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);
```

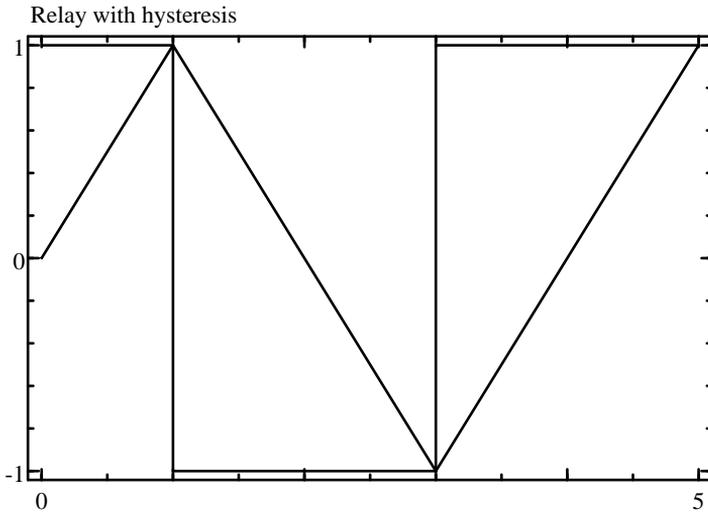
A single time event is generated at  $t = 1$ :

```
opt = odeset('EventTime', [inf, 1]);
(t, y) = ode45(@filterfun, [0, 5], 0, opt);
plot(t', y');
```

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:



**Figure 3.4** Relay with hysteresis integrated with events

```
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. 3.4) with

```
(t, y) = ode45(@relayfun, [0, 5], [0; 1], ...
    odeset('Events', @relayevents, 'OnEvent', @relayonevent));
plot(t', y');
```

**See also**

ode23, ode45, optimset

**optimset**

Options for minimization and zero finding.

**Syntax**

```
options = optimset
options = optimset(name1, value1, ...)
options = optimset(options0, name1, value1, ...)
```

**Description**

optimset(name1,value1,...) creates the option argument used by fminbnd, fminsearch, and fzero. 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, 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.

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"):

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
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
ToIX: []

```

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

```

## See also

fzero, fminbnd, fminsearch, odeset

## 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 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**

$$\int_0^2 te^{-t} dt$$

```
quad(@(t) t*exp(-t), 0, 2)
0.5940
```

**See also**

sum, ode45, inline, operator @

## 3.21 String Functions

**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

**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 '='; 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 1521.

## 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`

## 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\r\n\0'))
  32  9 65 66 32 32 67 68
```

**See also**

strtrim

**findstr**

Find a substring in a string.

**Syntax**

```
pos = findstr(str, sub)
```

**Description**

`findstr(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**

```
findstr('ababcdbaaab','ab')
    1 3 10
findstr('ababcdbaaab','ac')
    []
findstr('aaaaaa','aaa')
    1 2 3
```

**See also**

`find`, `strcmp`, `strmatch`, `strtok`

**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.

**Examples**

```
isdigit('a123bAB12* ')  
F T T T F F F T T F F
```

**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 a letter and false otherwise. Letters with diacritical signs are not considered as letters.

**Examples**

```
isletter('abAB12* ')  
T T T T F F F F
```

**See also**

isdigit, isspace, lower, upper, ischar

**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.

**Example**

```
isspace('a\tb c\n d')  
0 1 0 1 0 1 0
```

**See also**

isletter, isdigit, ischar

**lower**

Convert all uppercase letters to lowercase.

**Syntax**

```
s2 = lower(s1)
```

**Description**

`lower(s1)` converts all the uppercase letters of string *s1* to lowercase. Currently, only ASCII letters (without diacritic) are converted.

**Example**

```
lower('abcABC123')
abcabc123
```

**See also**

upper, isletter

**md5**

Calculate MD5 digest.

**Syntax**

```
digest = md5(strb)
digest = md5(fd)
```

**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.

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**

sha1

**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**

Calculate SHA1 digest.

**Syntax**

```
digest = sha1(strb)
digest = sha1(fd)
```

**Description**

sha1(strb) calculates the SHA1 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.

`sha1(fd)` calculates the SHA1 digest of the bytes read from file descriptor `fd` until the end of the file. The file is left open.

SHA1 digest is an Internet standard described in RFC 3174.

### Example

SHA1 digest of the three characters 'a', 'b', and 'c':

```
sha1('abc')
a9993e364706816aba3e25717850c26c9cd0d89d
```

### See also

`md5`

### `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', 'abc', 5)
false
```

### See also

`strcmpi`, `operator ===`, `operator ~=`, `operator ==`, `findstr`, `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 ~==, operator ==, findstr, strmatch

## 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 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', ['xyz'; 'uabc'; 'abcd'; 'efgh'])
    3
strmatch('abc', ['xyz'; 'uabc'; 'abcd'; 'efgh'], 'exact')
    0
strmatch('abc', {'ABC', 'xyz', 'abcdefg', 'ab', 'abcd'})
    3

```

**See also**

strcmp, findstr

**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.

**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, findstr, 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\r\n\0'))
65 66 32 32 67 68
```

### See also

deblank, strtok

## upper

Convert all lowercase letters to uppercase.

### Syntax

```
s2 = upper(s1)
```

**Description**

`upper(s1)` converts all the lowercase letters of string `s1` to uppercase. Currently, only ASCII letters (without diacritic) are converted.

**Example**

```
upper('abcABC123')
ABCABC123
```

**See also**

`lower`, `isletter`

**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 one, two, or three bytes of UTF-8 code. The result is a standard character array with a single row. Invalid codes (for example when the beginning of the decoded data does not correspond to a character boundary) are ignored.

**See also**

`utf8encode`

**utf8encode**

Encode a string of Unicode characters using UTF-8.

**Syntax**

```
b = utf8encode(str)
```

## Description

`utf8encode(b)` encodes the contents of character array `str` using UTF-8. Each Unicode character in `str` corresponds to one, two, or three 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);
+str
    1x6 uint16 array
    97 98 99 200 2000 20000
```

## See also

`utf8decode`

## 3.22 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:

Function	Operator	Purpose
ctranspose	'	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.

<b>Function</b>	<b>Purpose</b>
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.

<b>Function</b>	<b>Purpose</b>
cumsum	cumulative sum
diff	differences
double	conversion to array of double
mean	arithmetic mean
sum	sum

Functions below are related to array size.

<b>Function</b>	<b>Purpose</b>
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.

<b>Function</b>	<b>Purpose</b>
<code>disp</code>	display
<code>dumpvar</code>	conversion to string
<code>subsasgn</code>	assignment to subarrays or to quaternion parts
<code>subsref</code>	reference 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**

$R = \text{q2mat}(q)$

**Description**

$R = \text{q2mat}(q)$  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  $aq = \text{quaternion}(x, y, z)$ , 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
```

$$\frac{\text{abs}(p)}{2}$$
**See also**

quaternion, qnorm, conj

**qnorm**

Quaternion scalar norm.

**Syntax**
$$n = \text{qnorm}(q)$$
**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**
$$q = \text{qslerp}(q1, q2, t)$$
**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

Quaternion 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)
q =
    1+2i+3j+4k
q + 5
    6+2i+3j+4k
q * q
    -28+4i+6j+8k
Q = [q, 2; 2*q, 5]
    2x2 quaternion array
Q.y
    3  0
    6  0
q = quaternion(1, [5; 3; 7])
q =
    1+5i+3j+7k
q.v
    5
    3
    7

```

**See also**

real, qimag, 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(θ/2) and imaginary part sin(θ/2)(v\_xi + v\_yj + v\_zk), for a rotation of θ around unit vector

$[v_x v_y v_z]^T$ . The rotation is applied to a point  $[x y z]^T$  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

## 3.23 List Functions

### apply

Function evaluation with arguments in lists.

## Syntax

```
listout = apply(fun, listin)
listout = apply(fun, listin, nargout)
```

## 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 either by its name as a string or by an inline function.

The number of expected output arguments can be specified with an optional third input argument `nargout`.

## Examples

```
apply('min', {5, 7})
    {5}
apply('size',{magic(3)},2)
    {3, 3}
apply(inline('2*x+3*y','x','y'), {5, 10})
    {40}
```

## See also

`map`, `feval`, `inline`, `operator @`

## join

List concatenation.

## 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'}
```

**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. fun is the name of a function as a string, a function reference, or an inline function.

**Examples**

```
map('max', {[2,6,4], [7,-1], 1:100})
  {6, 7, 100}
map(inline('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
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

## 3.24 Structure Functions

**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(struct('a', 1, 'b', 1:5))
{'a', 'b'}
```

**See also**

struct, isfield, orderfields, rmfield

**getfield**

Value of a field in a structure.

**Syntax**

```
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(struct('a', 1:3, 'x', 'abc'), 'x')
true
isfield(struct('a', 1:3, 'x', 'abc'), 'X')
false
```

## See also

`isstruct`, `struct`

## isstruct

Test for a structure object.

## Syntax

```
b = isstruct(obj)
```

## Description

`isstruct(obj)` is true if the object `obj` is a structure, false otherwise. Structures are lists whose at least one field has a name.

## Examples

```
isstruct(struct('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 = struct('a',123,'c',1:3,'b','123')
s =
  a: 123
  c: real 1x3
  b: 'abcde'
(t, p) = orderfields(s)
t =
  a: 123
  b: 'abcde'
  c: real 1x3
p =
  1
  3
  2
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(struct('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`. `setfield` 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`, `getfield`, `rmfield`

**struct**

Creation of a structure

**Syntax**

```
strct = struct(fieldname1, value1, fieldname2, value2, ...)
```

**Description**

`struct` builds a new structure. 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.

**Example**

```
x = struct('a', 1, 'b', 2:5);  
x.a  
    1  
x.b  
    2 3 4 5
```

**See also**

isstruct, isfield, rmfield, fieldnames, operator {}

## 3.25 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.

<b>Class</b>	<b>Native type</b>
double	real, complex, or logical scalar or array
char	character or character array
list	list or structure
inline	inline function
funref	function reference

#### Examples

```
o1 = class(struct('fld1', 1, 'fld2', rand(4)), 'c1');
o2 = class(struct('fld3', 'abc'), 'c2', o1);
class(o2)
c2
```

**See also**

map, isa, isobject, methods

**isa**

Test for an object of a given class.

**Syntax**

```
b = isa(object, 'classname')
```

**Description**

`isa(object, 'classname')` returns true if object is an object of class `class`, directly or by inheritance.

**Example**

```
isa(pi, 'double')  
true
```

**See also**

class, isobject, methods

**isobject**

Test for an object.

**Syntax**

```
b = isobject(a)
```

**Description**

`isobject(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`

# 3.26 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.

**Syntax**

```
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).

**Syntax**

```
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([0uint8, 1uint8, 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.

**Syntax**

```
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  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*.

**Syntax**

```
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 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.

**Syntax**

```
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.

### Syntax

```

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 `|`

## 3.27 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

$$\begin{aligned}\frac{dx}{dt}(t) &= A_c x(t) + B_c u(t) \\ y(t) &= C_c x(t) + D_c u(t)\end{aligned}$$

and

$$\begin{aligned}x(k+1) &= A_d x(k) + B_d u(k) \\ y(k) &= C_d x(k) + D_d u(k)\end{aligned}$$

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 `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`.

`d2cm(Ad, Bd, Cd, Dd, Ts, method)` performs the conversion from discrete-time state-space model  $(A_d, B_d, C_d, D_d)$  to continuous-time state-space model  $(A_c, B_c, C_c, D_c)$ , defined by

$$\begin{aligned}x(k+1) &= A_d x(k) + B_d u(k) \\ y(k) &= C_d x(k) + D_d u(k)\end{aligned}$$

and

$$\begin{aligned}\frac{dx}{dt}(t) &= A_c x(t) + B_c u(t) \\ y(t) &= C_c x(t) + D_c u(t)\end{aligned}$$

### 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 `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) = 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`

## 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

$$\begin{aligned}\frac{dx}{dt}(t) &= Ax(t) + Bu(t) \\ y(t) &= Cx(t) + Du(t)\end{aligned}$$

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 null, and one less if  $D$  is null.

## Example

```
(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`

## 3.28 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

## 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)
```

## Description

`disp(obj)` displays the object `obj`. Command format may be used to control how numbers are formatted.

## 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`

**format**

Default output format.

**Syntax**

```
format
format short
format short e
format short eng
format short g
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 '+'
format i
format j
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:

<b>Arguments</b>	<b>Meaning</b>
(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)
+	'+', '-' 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 '+' 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

## fprintf

Formatted output.

## Syntax

```
n = fprintf(fd,format,a,b,...)
n = fprintf(format,a,b,...)
```

## 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. 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
```

## Caveat

Same limitations as `sprintf`

## 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 bytes 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.

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.

<b>precision</b>	<b>meaning</b>
<code>int8</code>	signed 8-bit integer ( $-128 \leq x \leq 127$ )
<code>char</code>	signed 8-bit integer ( $-128 \leq x \leq 127$ )
<code>int16</code>	signed 16-bit integer ( $-32768 \leq x \leq 32767$ )
<code>int32</code>	signed 32-bit integer ( $-2147483648 \leq x \leq 2147483647$ )
<code>int64</code>	signed 64-bit integer ( $-9.223372e18 \leq x \leq 9.223372e18$ )
<code>uint8</code>	unsigned 8-bit integer ( $0 \leq x \leq 255$ )
<code>uchar</code>	unsigned 8-bit integer ( $0 \leq x \leq 255$ )
<code>uint16</code>	unsigned 16-bit integer ( $0 \leq x \leq 65535$ )
<code>uint32</code>	unsigned 32-bit integer ( $0 \leq x \leq 4294967295$ )
<code>uint64</code>	unsigned 64-bit integer ( $0 \leq x \leq 18.446744e18$ )
<code>single</code>	32-bit IEEE floating-point
<code>double</code>	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).

## See also

`fwrite`, `sread`

## 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**

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

```

## sprintf

Formatted conversion of objects into a string.

### Syntax

```
s = sprintf(format,a,b, ...)
```

### 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, 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; by default, it is 4 for a number or the length of the string for a string), and `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**

<code>%</code>	single <code>%</code>
<code>d</code>	decimal number as an integer
<code>i</code>	same as <code>d</code>
<code>x</code>	hexadecimal number (for integers between 0 and $2^{32}-1$ )
<code>X</code>	same as <code>x</code> , with uppercase digits
<code>o</code>	octal number (for integers between 0 and $2^{32}-1$ )
<code>f</code>	fixed number of decimals (exp. notation if $\text{abs}(x) > 1e18$ )
<code>F</code>	same as <code>f</code> , with an uppercase <code>E</code>
<code>e</code>	scientific notation such as <code>1e5</code>
<code>E</code>	scientific notation such as <code>1E5</code>
<code>n</code>	engineering notation such as <code>100e3</code>
<code>N</code>	engineering notation such as <code>100E3</code>
<code>g</code>	decimal or scientific notation
<code>G</code>	same as <code>g</code> , with an uppercase <code>E</code>
<code>k</code>	same as <code>g</code> , with as few characters as possible
<code>K</code>	same as <code>k</code> , with an uppercase <code>E</code>
<code>c</code>	character
<code>s</code>	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.

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**

```

printf('%d %.2f %.2e %.2E %.2g', pi*ones(1,5))
3 3.14 3.14e0 3.14E0 3.14
printf('%%.1k ', 0.001, 0.11, 111, 1000)
1e-3 0.11 111 1e3
printf('%*8.3f%*8.6s%*-8.6s*', pi, 'abcdefg', 'abcdefg')
* 3.142* abcdef*abcdef *
printf('%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_
printf('%*.*f', 15, 7, pi)
3.1415927
printf('%%.3d,%.3d', 12, 12345)
012,12345

```

**Caveat**

Exotic formats unsupported.

**See also**

fprintf, sscanf, write

**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

## sscanf

Decoding of formatted numbers.

### Syntax

```

r = sscanf(str, format)
(r, count) = sscanf(str, format)
(r, count, nchar) = sscanf(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:

<b>Substring in format</b>	<b>Meaning</b>
%C	single character
%S	string
%d	integer number in decimal
%X	unsigned integer number in hexadecimal
%O	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)')
2.3
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')
34
sscanf('0275a0ff', '%2x')
2
117
160
255

```

## See also

`sprintf`

## swrite

Store data in a string.

## Syntax

```

s = swrite(data)
s = swrite(data, precision)

```

## Description

`swrite(fd, data)` stores the contents of the matrix data in a string. The third parameter is the precision, whose meaning is the same as for `fread`. Its default value is `'uint8'`.

**Examples**

```

write(65:68)
  ABCD
double(write([1,2], 'int16'))
  1 0 2 0
double(write([1,2], 'int16;b'))
  0 1 0 2

```

**See also**

bwrite, fwrite, sprintf

## 3.29 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>	<b>Description</b>
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
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

## 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`

## 3.30 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 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**

`tic`, `toc`

**tic**

Start stopwatch.

**Syntax**

```
tic
```

**Description**

`tic` resets the stopwatch. Typically, `tic` is used once at the beginning of the block to be timed.

**See also**

`toc`, `clock`

**toc**

Elapsed time of stopwatch.

**Syntax**

```
elapsed_time = toc
```

**Description**

`toc` gets the time elapsed since the last execution of `tic`. Typically, `toc` is used at the end of the block of statements to be timed.

On multi-tasking operating systems like Windows, Mac OS X and Unix, `toc` measures only the time spent in the LME application. Other processes do not have a large impact. For instance, typing `tic` at the command-line prompt, waiting 5 seconds, and typing `toc` will show a value much smaller than 5.

**Example**

```
tic; x = eig(rand(200)); toc  
0.3046
```

**See also**

`tic`, `clock`

## 3.31 Date Conversion 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 as 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:

```
julian2cal(cal2julian(2005, 4, 18) + 1000)
2006 11 14 0 0 0
```

**See also**

`cal2julian`

## 3.32 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	

*Scaling and labels*

label	plotoption	scalefactor
legend	scale	title

*3D*

contour3	plot3	surf
line3	plotpoly	
mesh	sensor3	

*3D scaling and lighting*

camdolly	camroll	daspect
camorbit	camtarget	lightangle
campan	camup	material
campos	camva	
camproj	camzoom	

### 3.33 Remarks on graphics

Most 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. The possible values 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.

<b>Color</b>	<b>String</b>
black	k
blue	b
green	g
cyan	c
red	r
magenta	m
yellow	y
white	w
RGB	h(rrggbb)
RGB	h(rgb)

<b>Dash Pattern</b>	<b>String</b>
solid	_ (underscore)
dashed	-
dotted	:
dash-dot	!

<b>Shape</b>	<b>String</b>
none (invisible)	(space)
point	.
circle	o
cross	x
plus	+
star	*
triangle up	^
triangle down	v
square	[
diamond	<

Miscellaneous	String
stairs	s
fill	f
arrow at end	a
arrows at beginning and end	A

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 `'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(cccccc)')
```

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:

<b>Command</b>	<b>Intended use</b>
<code>hgrid</code>	<code>nyquist</code> , <code>dnyquist</code>
<code>ngrid</code>	<code>nichols</code> , <code>dnichols</code>
<code>sgrid</code>	<code>plotroots</code> , <code>rlocus</code> (continuous-time)
<code>zgrid</code>	<code>plotroots</code> , <code>rlocus</code> (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]);
```

## 3.34 Base Graphical Functions

### activerregion

Region associated with an ID.

#### Syntax

```
activerregion(xmin, xmax, ymin, ymax, id)
activerregion(X, Y, id)
```

## Description

The command `activerregion` 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.

`activerregion(xmin, xmax, ymin, ymax, id)` defines a rectangular shape.

`activerregion(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 `activerregion`, a hit would be detected when the mouse is close to any of the four corners; with `activerregion`, a hit is detected when the mouse is inside the rectangle.

```
plot([50, 70, 70, 50, 50], [10, 10, 30, 30, 10]);
activerregion(50, 70, 10, 30, 1);
```

## See also

`plot`, `image`

## 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  $y=y_0$  and values defined by `y`.

The optional arguments `style` and `id` have their usual meaning. `area` uses default colors when argument `style` is missing.

**Example**

```
area(rand(20,10))
```

**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)
```

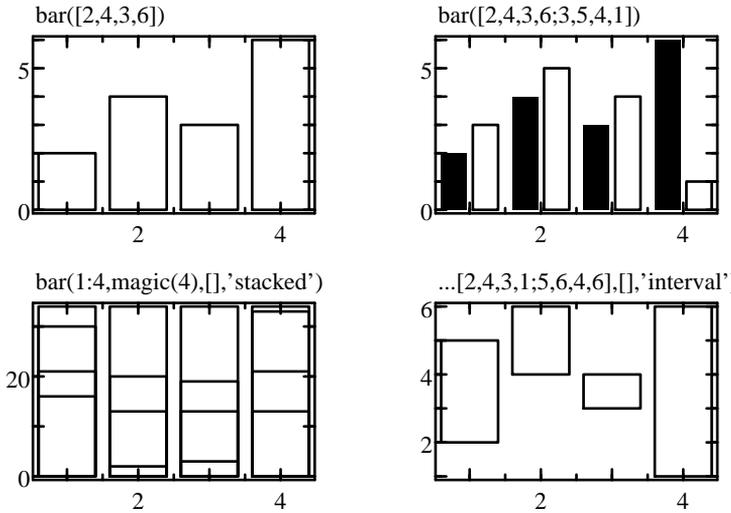
**Description**

`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.



**Figure 3.5** Example of bar with different options

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. 3.5):

```
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*

### **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` axis 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.

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.

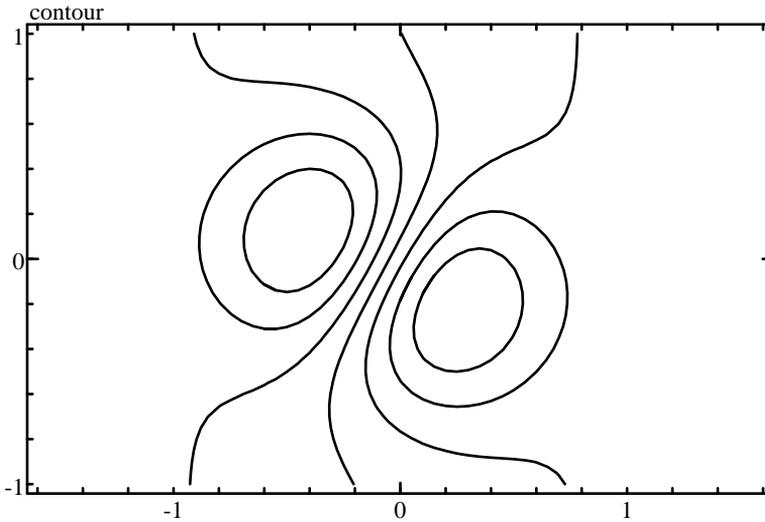
The optional fourth argument is the style of each line, from the minimum to the maximum level (styles are recycled if necessary). The default style is `'kbrmgcy'`.

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. 3.6):

```
(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]);
```



**Figure 3.6** Example of contour

Filled contours:

```

u = -2 + (0:80) / 20;
x = repmat(u, 81, 1);
y = x';
z = exp(-((x-0.2).^2+(y+0.3).^2)) ...
    - exp(-((x+0.5).^2+(y-0.1).^2)) ...
    + 0.1 * x ...
    + 0.5 * sin(y);
levels = -1:0.2:1;
scale equal;
contour(z, [-1,1,-1,1], levels, 'f');

```

### See also

image, quiver

### fontset

Options for fonts.

### Syntax

```

options = fontset
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. 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.

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"):

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
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
```

**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 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**

Plot a sine:

```
fplot(@sin);
```

Plot  $(x + 0.3)^2 + a \exp -3x^2$  in red for  $x \in [-2, 3]$  with  $a = 7.2$  and an identifier of 1:

```
fun = inline(...
'function y=f(x,a); y=(x+0.3)^2+a*exp(-3*x^2);');
fplot(fun, [-2,3], 'r', 1, 7.2);
```

**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(..., style)
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), or by a uint8 matrix in the range 0 (black) to 255 (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, or between 0 (black) to 255 (maximum intensity) with uint8 values.

The position is defined by the the minimum and maximum coordinates along the horizontal and vertical axis. The raster image is scaled to fit. The first line of the matrix or matrices is displayed at the top. If `style` 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 `style` is '1', and the bicubic method if `style` is '3'.

## Examples

Two ways to display a table of 10-by-10 random color cells (see Fig. 3.7):

```
image(rand(10), rand(10), rand(10));
image(rand(10, 10, 3));
```

A ramp of gray shades:

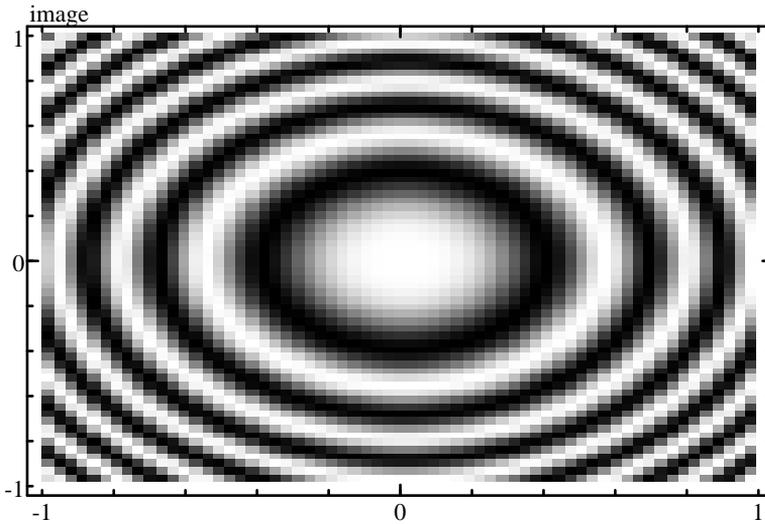
```
image(uint8(0:255));
```

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`



**Figure 3.7** Example of image

## label

Plot labels.

### Syntax

```
label(label_x)
label(label_x, label_y)
```

### Description

`label(label_x, label_y)` displays labels for the X and Y axis. Its arguments are strings. The label for the Y axis may be omitted.

### Examples

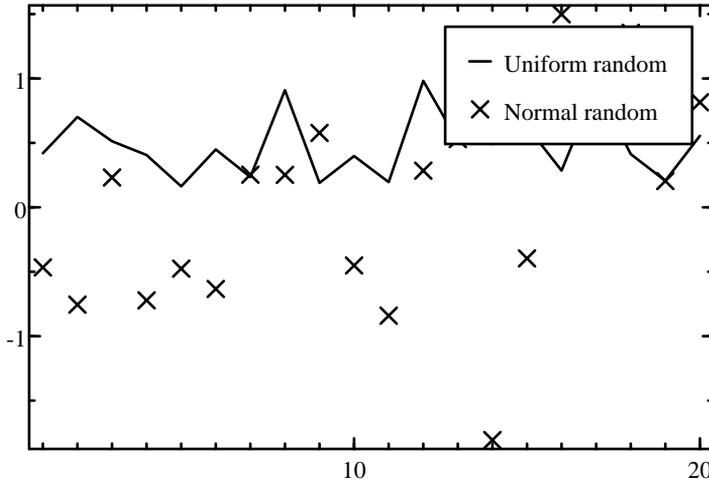
```
step(1,[1,2,3,4]);
label('t [s]', 'y [m]');
```

With literal strings, the command syntax may be more convenient:

```
label Re Im;
```

### See also

`text`, `legend`, `title`



**Figure 3.8** Example of legend

## legend

Plot legend.

### Syntax

```
legend(str)
legend(str, style)
```

### 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.

With a single input argument, legend(str) uses the default style 'k'.

### Example

Legend for two traces (see Fig. 3.8).

```
plot(1:20, [rand(1,20); randn(1,20)], '-x');
legend('Uniform random\nNormal random', '-x');
```

**See also**

label, title

**line**

Plot lines.

**Syntax**

```
line(A, b)
line(A, b, style)
line(A, b, style, id)
```

**Description**

line displays one or several straight line(s). Each line is defined by an equation 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; each row corresponds to a different line. The second argument is a column vector which contains the coefficients  $b$ . If one of these 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 lines at  $y=2$  and  $y=3$ :

```
line([0,1],[2;3], 'b', 1)
```

**See also**

plot, circle

## **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`

## 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 `_x0`, `_y0`, and `_z0` are set to the position of the point before it is moved; the variable `_id` is set to the ID as defined by the command `plot`; the variable `_nb` is set to the number of the row, and the variable `_ix` is set to the index of the column of the matrix `x` and `y`.

### Examples

Sine between 0 and  $2\pi$ :

```
x = 2 * pi * (0:100) * 0.01;
y = sin(x);
plot(x, y);
```

Ten random crosses:

```
plot(rand(1,10), rand(1,10), 'x');
```

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, 'rf', 1); % filled red triangle with 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.

## Syntax

```
plotoption(str)
```

## Description

plotoption sets the initial value of one of the plot options the user can change. Its single argument, a character string, can take one of the following values.

'frame' Rectangular frame with tick marks and a white background around the plot.

'noframe' No frame, no tickmarks, no white background.

- 'label' Subplot name above the frame.
- 'nolabel' No subplot name.
- 'legend' Legend (if it has been set with legend).
- 'nolegend' Hidden legend.
- 'trlegend' Legend in top right corner (default).
- 'tllegend' Legend in top left corner.
- 'brlegend' Legend in bottom right corner.
- 'bllegend' Legend in bottom left corner.
- 'margin' Margin (space for title and labels).
- 'nomargin' No margin.
- 'xgrid' Grid of vertical lines for the x axis.
- 'noxgrid' No grid for the x axis.
- 'ygrid' Grid of horizontal lines for the y axis.
- 'noygrid' No grid for the y axis.
- 'xygrid' Grid of vertical and horizontal lines for the x and y axis.
- 'noxygrid' No grid for the x and y axis.
- 'grid' Normal details for grids displayed by sgrid, zgrid, etc.
- 'nogrid' Removal of grids displayed by sgrid, zgrid, etc.
- 'fullgrid' More details for grids displayed by sgrid, zgrid, etc.
- 'fill3d' In 3D graphics, zoom in so that the bounding box fills the figure.

### Example

Display of a photographic image without frame:

```
plotoption noframe;  
image(photo);
```

### See also

scale, legend

## 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 axis, 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  $[1,2,\dots,m]$  and  $[1,2,\dots,n]$  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])
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

```

## 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 axis or only for the x axis in 2D graphics, or for x, y and z axis 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.

- 'equal' Same linear scale for x and y axis. Typically used for representation of the complex plane, such as the roots of a polynomial or a Nyquist diagram. For 3D graphics, same effect as `daspect([1,1,1])`.
- 'lock' See below.
- 'linlin' Linear scale for both axis.
- '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 axis.
- '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.

This last 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.

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.

<b>Bit</b>	<b>Meaning</b>
0	<b>log x</b>
2	tick on x axis
3	grid for x axis
4	labels on x axis
6	<b>log y</b>
7	<b>dB y</b>
8	tick on y axis
9	grid for y axis
10	labels on y axis
12	same scale on both axis
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 numerical 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 an output argument, 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 x and y 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.

**Examples**

Here are some suggestions for the most usual graphics:

Time response	(default linlin is fine)
Bode mag	scale logdb
Bode phase	scale loglin
D bode mag	scale('lindb/logdb',[0,pi/Ts])
D bode phase	scale('linlin/loglin',[0,pi/Ts])
Poles	scale equal
D poles	scale('equal',[-1,1,-1,1])
Nyquist	scale('equal',[-1.5,1.5,-1.5,1.5])
Nichols	scale lindb

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 real positive numbers to set separately the x axis and the y axis, or a real positive scalar to set the same factor for both axis. 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`

### text

Display text in a figure.

### Syntax

```
text(x, y, string)
text(x, y, string, justification)
text(..., font)
```

### 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:

<b>Char.</b>	<b>Alignment</b>
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`.

## 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);
```

## See also

label, fontset, sprintf

## title

Subplot title.

## Syntax

```
title(string)
```

## Description

title(string) sets or changes the title of the current subplot.

## See also

label, legend, text, sprintf

## 3.35 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).

## 3.36 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`

## 3.37 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.

## 3.38 Functions

### `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)
```

**Description**

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
```

## Description

`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.

**Syntax**

```
camroll(dalpha)
```

**Description**

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.

## Syntax

```
camup(p)
camup auto
camup manual
p = camup
```

## Description

`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.

**Syntax**

```
camzoom(f)
```

**Description**

`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 azimuth `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(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 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.

The optional third and fourth arguments are the same as for all graphical commands.

**Example**

Vertical line at  $x=5$ ,  $y=6$ :

```
line3([1,0;0,1],[5;6])
```

**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 ambient 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.

## 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.

**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 sensitive to 3D interactive displacement.

## Syntax

```
sensor3(type, param, id)
sensor3(type, param, typeAlt, paramAlt, id)
```

**Description**

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\_(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.

## See also

`plot3`, `mesh`, `plotpoly`

## 3.39 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

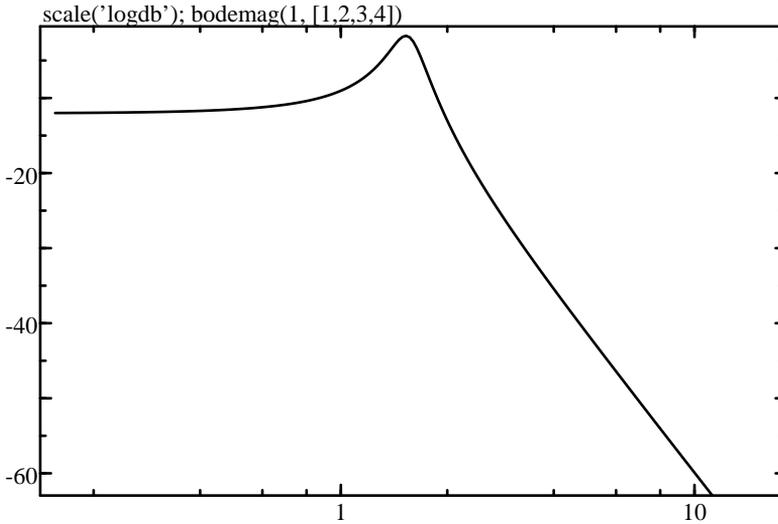
`bodemag(numc, denc)` plots the magnitude of the frequency response of the continuous-time transfer function  $\text{numc}/\text{denc}$ . The range of frequencies is selected automatically or can be specified in an optional argument `w`, a vector of frequencies.

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.

`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

$$\begin{aligned}j\omega X(j\omega) &= A_c X(j\omega) + B_c U(j\omega) \\ Y(j\omega) &= C_c X(j\omega) + D_c U(j\omega)\end{aligned}$$

With output arguments, `bodemag` gives the magnitude and the frequency as column vectors. No display is produced.



**Figure 3.9** `scale('logdb'); bodemag(1, [1,2,3,4])`

### Examples

Green plot for  $|1/(s^3 + 2s^2 + 3s + 4)|$  with  $s = j\omega$  (see Fig. 3.9):

```
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`

## 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  $\text{numc}/\text{denc}$ . 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.

`bodemag(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

$$\begin{aligned}
 j\omega X(j\omega) &= A_c X(j\omega) + B_c U(j\omega) \\
 Y(j\omega) &= C_c X(j\omega) + D_c U(j\omega)
 \end{aligned}$$

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. 3.10):

```
bodephase(1, [1, 2, 3, 4], 'g');
```

## See also

`bodemag`, `dbodephase`, `responseset`

## dbodemag

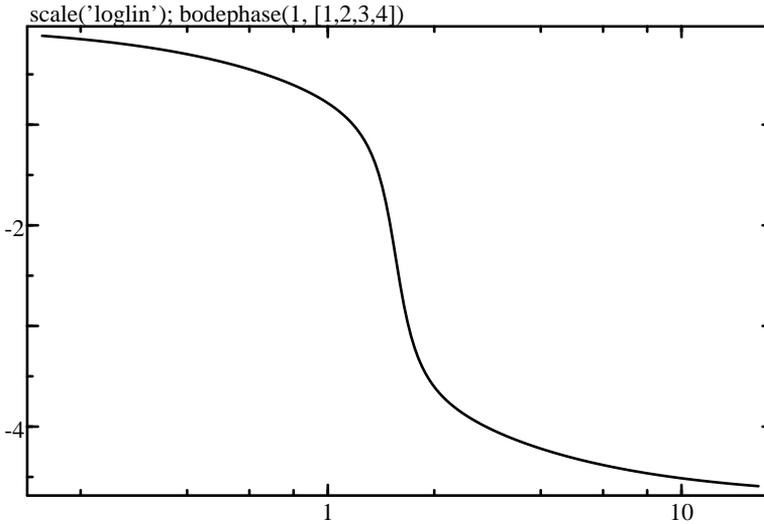
Magnitude Bode diagram for a discrete-time system.

## Syntax

```

dbodemag(numd, dend, Ts)
dbodemag(numd, dend, Ts, w)
dbodemag(numd, dend, Ts, opt)

```



**Figure 3.10** `scale('loglin'); bodephase(1, [1,2,3,4])`

```

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, 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  $(A_d, B_d, C_d, D_d)$  defined as

$$\begin{aligned}
 zX(z) &= A_d X(z) + B_d U(z) \\
 Y(z) &= C_d X(z) + D_d U(z)
 \end{aligned}$$

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`

## 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, 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

$$\begin{aligned} zX(z) &= A_d X(z) + B_d U(z) \\ Y(z) &= C_d X(z) + D_d U(z) \end{aligned}$$

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`

## 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 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.

`dimpulse(Ad,Bd,Cd,Dd,Ts)` plots the impulse response of the discrete-time state-space model `(Ad,Bd,Cd,Dd)` defined as

$$\begin{aligned}x(k+1) &= A_d x(k) + B_d u(k) \\ y(k) &= C_d x(k) + D_d u(k)\end{aligned}$$

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

**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, Bd, Cd, Dd, Ts, x0, opt)
dinitial(..., style)
dinitial(..., style, id)
(y, t) = dinitial(...)
```

**Description**

`dinitial(Ad,Bd,Cd,Dd,Ts,x0)` plots the output(s) of the discrete-time state-space model (Ad,Bd,Cd,Dd) with null input and initial state `x0`. The model is defined as

$$\begin{aligned}x(k+1) &= A_d x(k) + B_d u(t) \\ y(k) &= C_d x(k) + D_d u(k)\end{aligned}$$

where `u(k)` is null. Sampling period is `Ts`. The state-space model may have a scalar or vector output.

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

## 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 discrete-time state-space model  $(Ad, Bd, Cd, Dd)$  defined as

$$\begin{aligned}x(k+1) &= A_d x(k) + B_d u(t) \\ y(k) &= C_d x(k) + D_d u(k)\end{aligned}$$

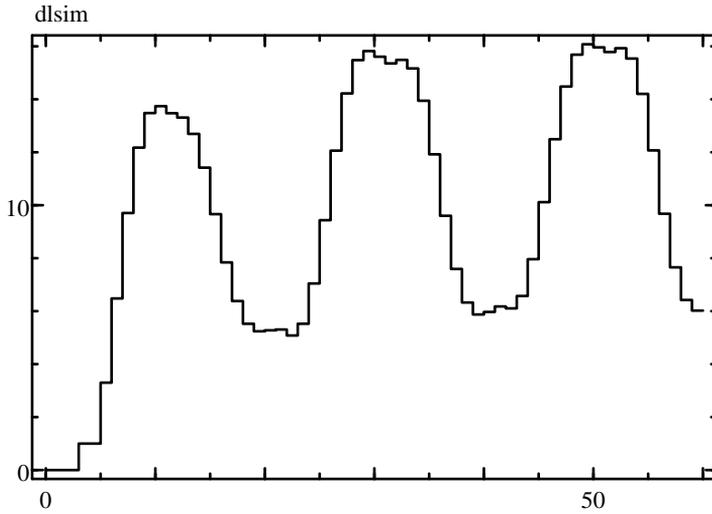
where the system input at time sample `k` is `u(k, :)'`. For single-input systems, `u` can also be a row vector.

`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.



**Figure 3.11** `dlsim(1, poly([0.9,0.7+0.6j,0.7-0.6j]), u)`

**Example**

Simulation of a third-order system with a rectangular input (see Fig. 3.11):

```
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`

**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`

## 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) (see Fig. 3.12)

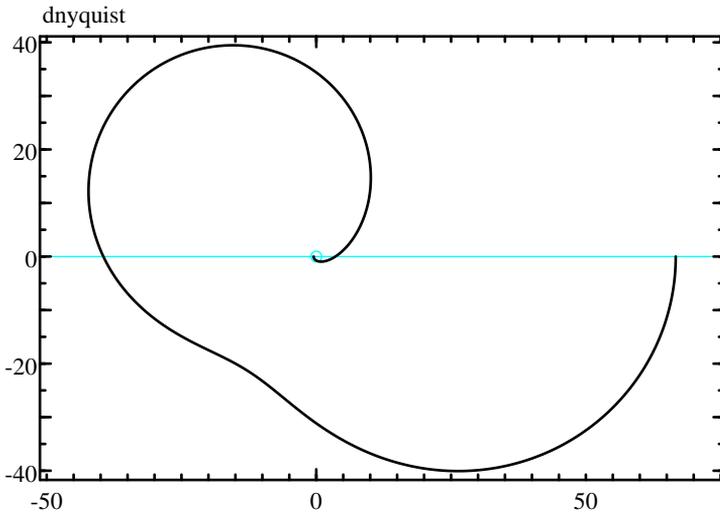
```
scale equal;
hgrid;
dnyquist(3, poly([0.9,0.7+0.6j,0.7-0.6j]))
```

## See also

`nyquist`, `hgrid`, `dnichols`, `responseset`

## **dsigma**

Singular value plot for a discrete-time state-space model.



**Figure 3.12** `dnyquist(3, poly([0.9,0.7+0.6j,0.7-0.6j]))`

### 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  $(A_d, B_d, C_d, D_d)$  defined as

$$\begin{aligned}
 zX(z) &= A_d X(z) + B_d U(z) \\
 Y(z) &= C_d X(z) + D_d U(z)
 \end{aligned}$$

where  $z = e^{j\omega T_s}$  and  $T_s$  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*

## 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

$$\begin{aligned}x(k+1) &= A_d x(k) + B_d u(k) \\ y(k) &= C_d x(k) + D_d u(k)\end{aligned}$$

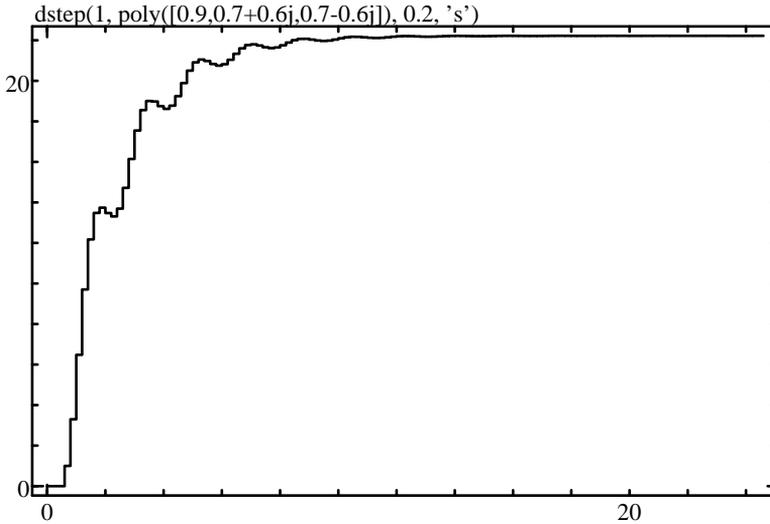
where *u(k)* is a unit step. The state-space model must have a scalar input, and may have a scalar or vector output.

With output arguments, *dstep* gives the output and the time as column vectors. No display is produced.

### Example

Step response of a discrete-time third-order system (see Fig. 3.13):

```
dstep(1, poly([0.9,0.7+0.6j,0.7-0.6j]), 1, 'g');
```



**Figure 3.13** `dstep(1, poly([.9, .7+.6j, .7-.6j]), 0.2, 's')`

### See also

`dimpulse`, `dlsim`, `step`, `hstep`, `responseset`

### erlocus

Root locus of a polynomial with coefficients bounded by an ellipsoid.

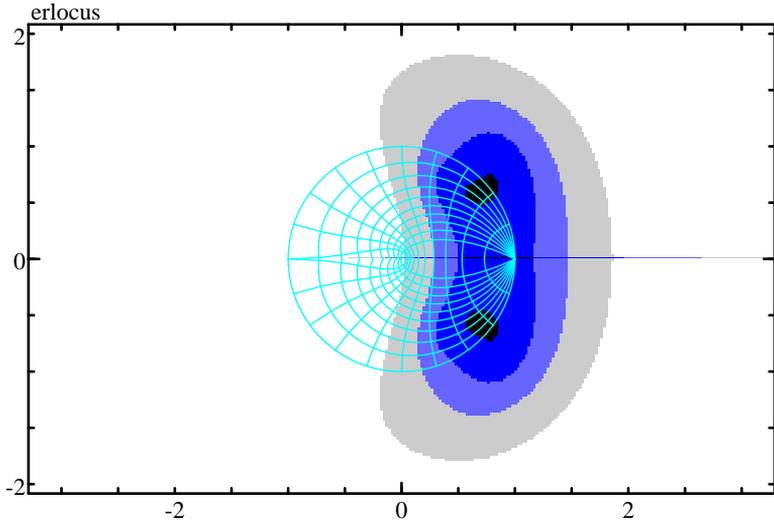
### Syntax

```
erlocus(C0, P)
erlocus(C0, 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  $dC \cdot \text{inv}(P) \cdot dC' < 1$ .

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  $dC \cdot \text{inv}(P) \cdot dC' < \text{sizes}(i)^2$  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 =`



**Figure 3.14** `erlocus(poly([.8,.7+.6j,.7-.6j]), eye(3))`

`[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. 3.14).

```

scale('equal', [-2,2,-2,2]);
erlocus(poly([0.8, 0.7+0.6j, 0.7-0.6j]), eye(3));
zgrid;

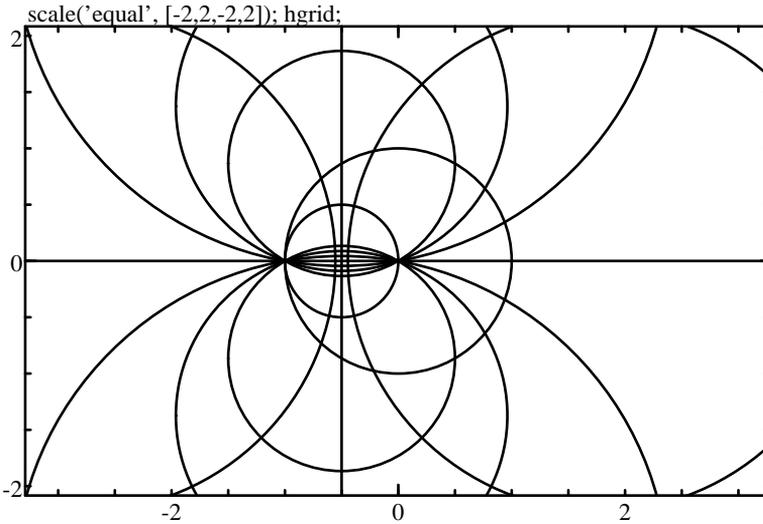
```

**See also**

`plotroots`, `rlocus`

**hgrid**

Hall chart grid.



**Figure 3.15** `scale('equal', [-2,2,-2,2]); hgrid`

## Syntax

```
hgrid
hgrid(style)
```

## 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 0.2, 0.5, 0.8, 1, 1/0.8, 2, and 5; 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. 3.15):

```
scale('equal', [-1.5, 1.5, -1.5, 1.5]);
hgrid;
nyquist(20, poly([-1,-2+1j,-2-1j]))
```

**See also**

ngrid, nyquist, plotoption

**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  $\text{numd}/\text{dend}$ , a discrete-time transfer function with sampling period  $T_s$ ; the resulting signal is converted to continuous-time with a zero-order hold, and filtered by the continuous-time transfer function  $\text{numc}/\text{denc}$ .

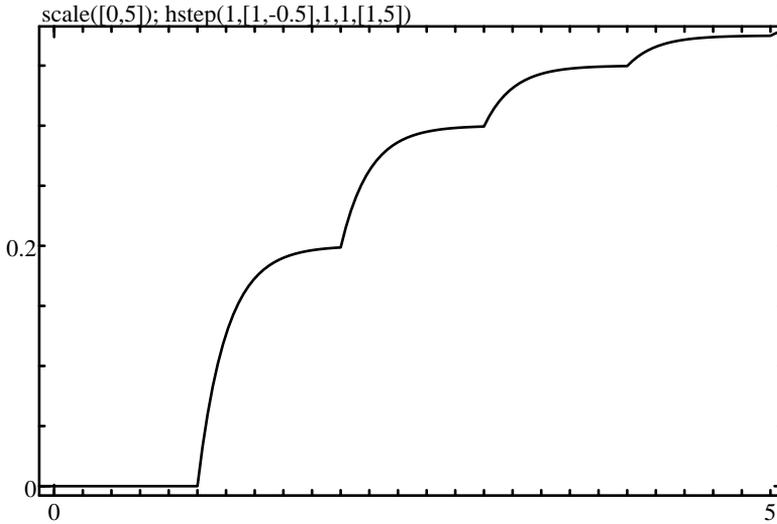
Most discrete-time controllers are used with a zero-order hold and a continuous-time system. `hstep` can be used to display the simulated output of the system when a step is applied somewhere in the loop, e.g. as a reference signal or a disturbance. The transfer function  $\text{numd}/\text{dend}$  should correspond to the transfer function between the step and the system input; the transfer function  $\text{numc}/\text{denc}$  should be the model of the system.

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. 3.16):

```
% 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
```



**Figure 3.16** `scale([0,5]); hstep(1,[1,-0.5],1,1,[1,5])`

```

kp = 1.5;
% transfer function between ref. signal and input
b = conv(kp, denc);
a = addpol(conv(kp, numd), denc);
% 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, denc), Ts, 'o');

```

### See also

step, dstep

### 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 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.

`impulse(Ac,Bc,Cc,Dc)` plots the impulse response of the continuous-time state-space model  $(A_c, B_c, C_c, D_c)$  defined as

$$\begin{aligned}\frac{dx}{dt}(t) &= A_c x(t) + B_c u(t) \\ y(t) &= C_c x(t) + D_c u(t)\end{aligned}$$

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');
```

## See also

`dimpulse`, `step`, `lsim`, `initial`, `responseset`

## initial

Time response plot for a continuous-time state-space model with initial conditions.

## Syntax

```
initial(Ac, Bc, Cc, Dc, 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

$$\begin{aligned}\frac{dx}{dt}(t) &= A_c x(t) + B_c u(t) \\ y(t) &= C_c x(t) + D_c u(t)\end{aligned}$$

where `u(t)` is null. The state-space model may have a scalar or vector output.

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. 3.17):

```
initial([-0.3,0.1;-0.8,-0.4],[2;3],[1,3;2,1],[2;1],[5;3])
```

## See also

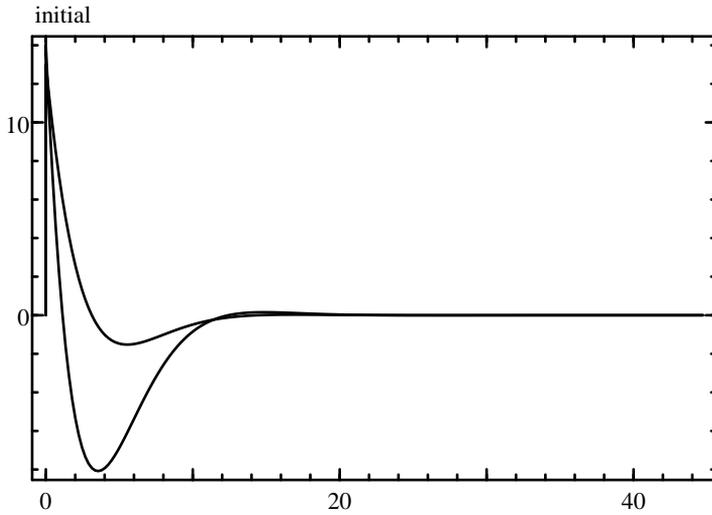
`dinitial`, `impulse`, `responseset`

## lsim

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)
```



**Figure 3.17** Example of initial

```
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 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. The input used for the simulation is interpolated to have a smooth response.

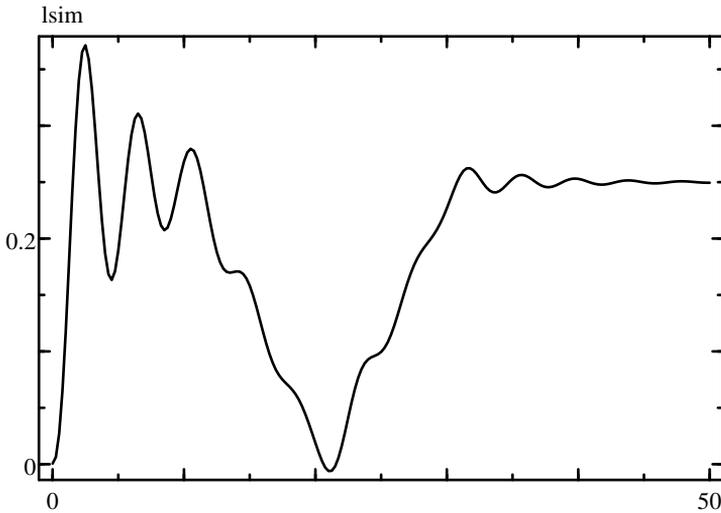
lsim(Ac,Bc,Cc,Dc,u,t) plots the time response of the continuous-time state-space model (Ac,Bc,Cc,Dc) defined as

$$\begin{aligned} \frac{dx}{dt}(t) &= A_c x(t) + B_c u(t) \\ y(t) &= C_c x(t) + D_c u(t) \end{aligned}$$

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:



**Figure 3.18** `lsim(1, [1,2,3,4], u, t)`

'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 state-space model, where each row represents a sample). No display is produced.

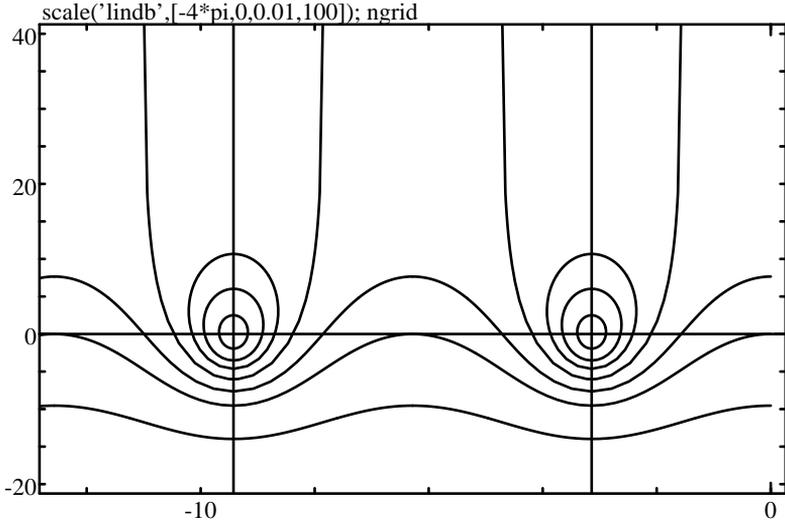
### Example

Response of continuous-time system given by its transfer function with an input defined by linear segments (see Fig. 3.18):

```
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`



**Figure 3.19** `scale('lindb',[-4*pi,0,0.01,100]); ngrid`

### ngrid

Nichols chart grid.

### Syntax

```
ngrid  
ngrid(style)
```

### Description

ngrid plots a Nichols chart in the complex plane of the Nichols diagram (see Fig. 3.19). 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.

**Example**

```
ngrid;
nichols(7, 1:3);
```

**See also**

hgrid, nichols, 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(...)
```

**Description**

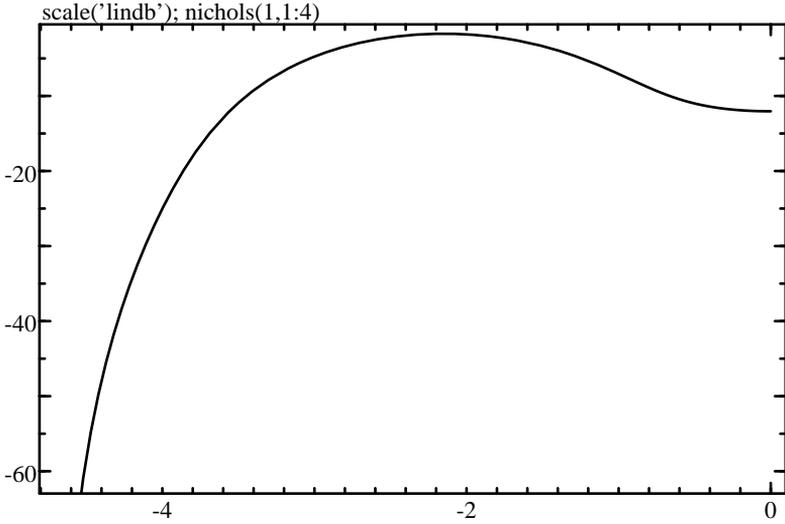
`nichols(numc, denc)` displays the Nichols diagram of the continuous-time transfer function given by polynomials `numc` and `denc`. In continuous time, the Nichols diagram is the locus of the complex values of the transfer function evaluated at  $j\omega$ , where  $\omega$  is real positive, displayed 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`.



**Figure 3.20** `scale('lindb'); nichols(1,1:4)`

**Example**

Nichols diagram of a third-order system (see Fig. 3.20):

```
scale('lindb');  
ngrid;  
nichols(20,poly([-1,-2+1j,-2-1j]))
```

**See also**

`dnichols`, `ngrid`, `nyquist`, `responseset`

**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.

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. 3.21):

```
scale equal;
hgrid;
nyquist(20, poly([-1, -2+1j, -2-1j]))
```

## See also

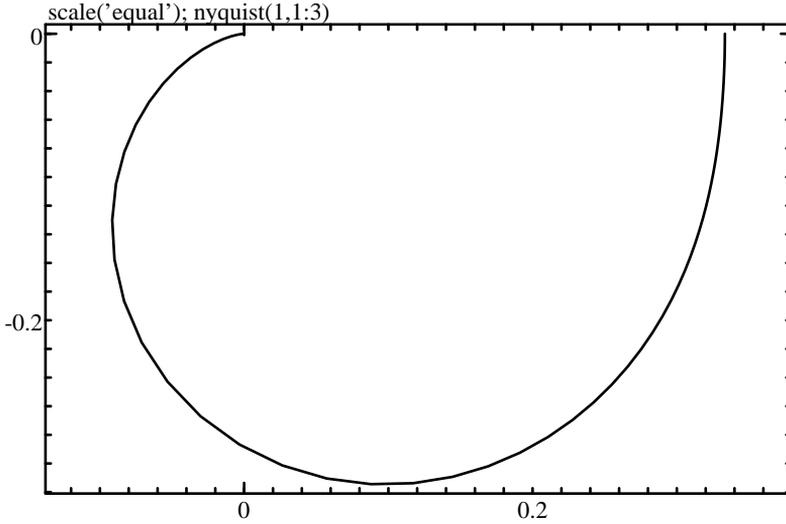
`dnyquist`, `hgrid`, `nichols`, `responseset`

## plotroots

Roots plot.

## Syntax

```
plotroots(pol)
plotroots(pol, style)
plotroots(pol, style, id)
```



**Figure 3.21** `scale equal; nyquist(1,[1,2,3])`

**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.

**Example**

```
scale equal;  
plotroots(den,'x');  
plotroots(num,'o');
```

**See also**

`rlocus`, `erlocus`, `sgrid`, `zgrid`, `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 numerical 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
Range	[]	time or frequency range
tOnly	false	samples for specified time only ( <code>lsim</code> )

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 `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

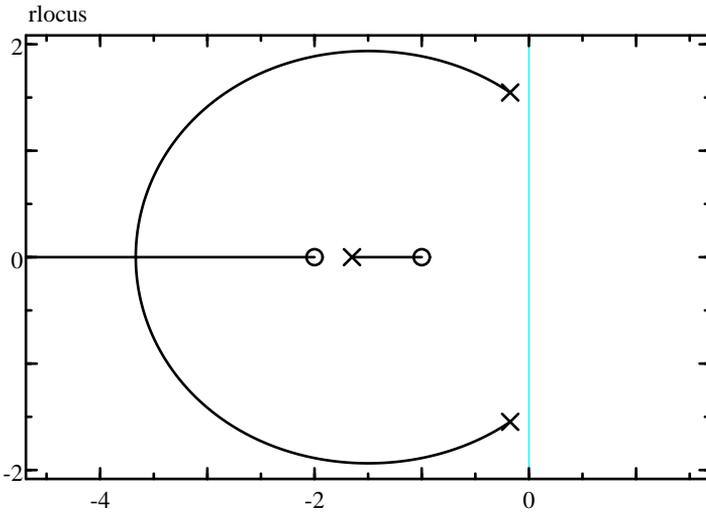
```
rlocus(num, den)  
rlocus(num, den, style)  
rlocus(num, den, style, id)
```

### 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  $\text{num}/\text{den}$  when the gain is between 0 and  $+\infty$  inclusive. The characteristic polynomial is  $\text{num} + k \cdot \text{den}$ , with  $k \geq 0$ . `rlocus` requires a causal system with real coefficients, i.e.  $\text{length}(\text{den}) \geq \text{length}(\text{num})$ . 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  $\text{length}(\text{den}) - 1$  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  $\text{num} + \text{den}$ ) 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.

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`



**Figure 3.22** Example of `rlocus`

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. 3.22):

```
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
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`, `erlocus`, `sgrid`, `zgrid`

### sgrid

Relative damping and natural frequency grid for the poles of a continuous-time system.

### Syntax

```

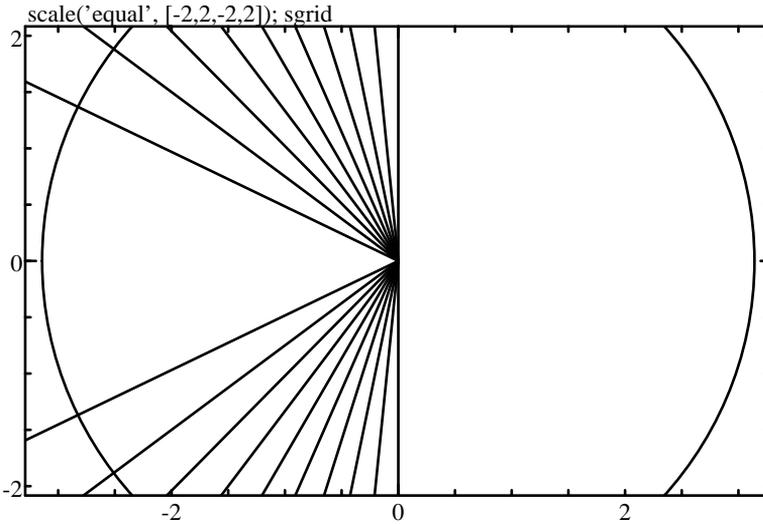
sgrid
sgrid(style)
sgrid(damping, freq)
sgrid(damping, freq, style)

```

### Description

With no or one argument, `sgrid` plots a grid of lines with constant relative damping and natural frequencies in the complex plane of  $s$  (see Fig. 3.23). The `style` argument has its usual meaning.

With two or three arguments, `sgrid` plots only the lines for the specified values of damping and natural frequency. Let  $p$  and  $\bar{p}$  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



**Figure 3.23** `scale('equal', [-2,2,-2,2]); sgrid`

with a constant damping  $\zeta$  is generated by  $|\text{Im } p| = \sqrt{1 - \zeta^2} \text{Re } p$  with  $\text{Re } p < 0$ . The locus of roots with a constant natural frequency  $\omega$  is a circle of radius  $\omega$ .

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) 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`, `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  $(A_c, B_c, C_c, D_c)$  defined as

$$\begin{aligned}
 j\omega X(j\omega) &= A_c X(j\omega) + B_c U(j\omega) \\
 Y(j\omega) &= C_c X(j\omega) + D_c U(j\omega)
 \end{aligned}$$

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`

**step**

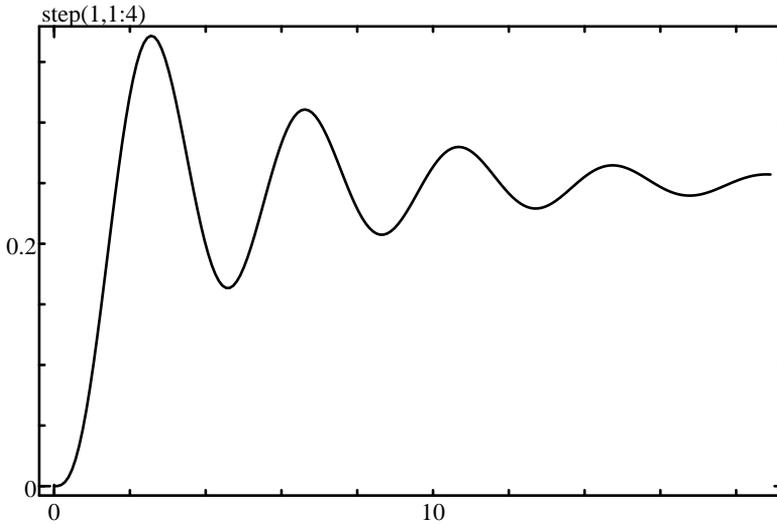
Step response plot of a continuous-time linear system.

**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(...)

```



**Figure 3.24** `step(1, [1,2,3,4])`

### Description

`step(numc,denc)` plots the step response of the continuous-time transfer function  $\text{numc}/\text{denc}$ .

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  $(A_c, B_c, C_c, D_c)$  defined as

$$\begin{aligned}\frac{dx}{dt}(t) &= A_c x(t) + B_c u(t) \\ y(t) &= C_c x(t) + D_c u(t)\end{aligned}$$

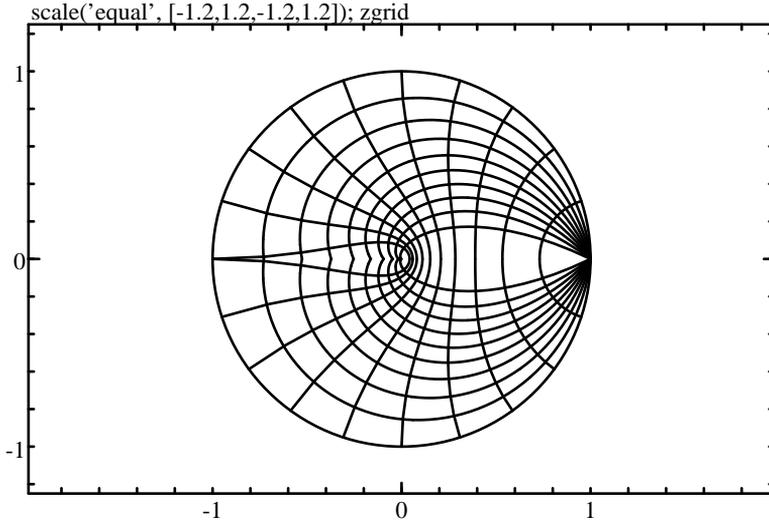
where  $u$  is a unit step. The state-space model must have a scalar input, and may have a scalar or vector output.

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. 3.24):

```
step(1, 1:4, 'b');
```



**Figure 3.25** `scale('equal', [-1.2,1.2,-1.2,1.2]); zgrid`

**See also**

`impulse`, `lsim`, `dstep`, `hstep`, `responseset`

**zgrid**

Relative damping and natural frequency grid for the poles of a discrete-time system.

**Syntax**

```
zgrid
zgrid(style)
zgrid(damping, freq)
zgrid(damping, freq, style)
```

**Description**

With no or one argument, `zgrid` plots a grid of lines with constant relative damping and natural frequencies in the complex plane of  $z$  (see Fig. 3.25). The `style` argument has its usual meaning.

With two or three 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  $\bar{z}$  are generated by  $z = e^{(-1+j\sqrt{1-\zeta^2}/\zeta)u}$ , with  $0 \leq u \leq u_{\max}$  and  $u_{\max}$  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  $e^{\omega e^{j\nu}}$ , where  $\nu$  is such that the curve is inside the unit circle.

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) is displayed.

### 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`, `plotoption`

## Chapter 4

# Extensions

Extensions are additional functions, usually developed in C or Fortran, which extend the core functionality of LME, the programming language of Sysquake. Extensions are grouped in so-called *shared libraries* or *dynamically-linked libraries* (DLL) files. At startup, Sysquake loads all extensions it finds in the folder LMExt in the same location as the Sysquake program file. Each extension initializes itself and usually displays a line of information in the Command window. No further action is needed in order to use the new functions.

You can also develop and add your own extensions, as explained in the next chapter.

Here is the list of the extensions currently provided with Sysquake.

### Mathematics

**Lapack (Windows, Mac, Unix)** LAPACK-based linear algebra functions.

**Long integers (Windows, Mac, Unix)** Arithmetic on arbitrary-length integer numbers.

### File input/output and data compression

**Memory mapping (Mac OS X, Unix)** Mapping of files in memory, which can be read and written like regular arrays.

**Data compression (Windows, Mac, Unix)** Support for compressing and uncompressing data using ZLib.

**Image Input/Output (Windows, Mac, Unix)** Support for reading and writing arrays as PNG or JPEG image files.

**MAT-file (Windows, Mac, Unix)** Support for reading and writing MAT-files (native MATLAB binary files).

## Databases

**ODBC (Windows)** Connection to a relational database using ODBC.

**MySQL (Windows, Mac OS X, Unix)** Connection to a MySQL relational database, locally or on a remote server.

**SQLite (Windows, Mac OS X, Unix)** SQLite, an embedded relational database in single files also using SQL as its query language.

## Operating system

**Socket (Windows, Mac, Unix)** TCP/IP communication with servers or clients on the same computer, on a local network or on the Internet.

**Launch URL (Windows, Mac, Unix)** Opening of documents in a World Wide Web browser.

**Download URL (Windows, Mac, Linux)** Download of documents from the World Wide Web.

**Open Script Architecture (Mac)** Communication with other applications.

**Power Management (Windows, Mac)** Functions related to power management.

**System Log (Mac OS X, Unix)** Output to system log.

**Shell (Windows, Mac OS X, Unix)** Shell related functions.

**Signal (Mac OS X, Unix)** Support for signals (POSIX functions kill and signal).

**Web Services (Windows, Mac OS X, Unix)** Web Services (standard remote procedure calls using XML-RPC and SOAP).

## Hardware support

**Serial port (Windows, Mac, Unix)** Communication with the serial port.

**I2C bus (Linux)** Communication with devices on an I2C bus.

**Joystick (Windows, Mac OS X, Linux)** Support for reading the state of a joystick or other similar device.

**Audio playback (Windows, Mac OS X, Linux)** Audio output.

**Audio recording (Windows, Mac OS X, Linux)** Audio input.

**Speech (Windows, Mac)** Speech output.

**Image Capture (Mac OS X)** Support for getting images from digital cameras.

## 4.1 Serial Port Functions

Serial port functions enable communication with devices connected to the computer via an RS-232 interface. Such devices include modems, printers, and many scientific instruments. The operating system can also emulate RS-232 connections with other devices, such as built-in modems or USB (Universal Serial Bus) devices.

Functions described in this section include only those required for opening and configuring the connection. They correspond to `fopen` for files. Input, output, and control are done with the following generic functions:

Function	Description
<code>fclose</code>	close the file
<code>fflush</code>	flush I/O buffers
<code>fgetl</code>	read a line
<code>fgets</code>	read a line
<code>fprintf</code>	write formatted data
<code>fread</code>	read data
<code>fscanf</code>	read formatted data
<code>fwrite</code>	write data
<code>redirect</code>	redirect output

Functions `opendev`, `devicename`, `closedev`, and `flushdev` are obsolete and may be removed in the future. They are replaced with `serialdevopen` and `serialdevset` to specify configuration settings, `serialdevname`, `fclose`, and `fflush`.

### **serialdevname**

Serial device name.

### **Syntax**

```
name = serialdevname(n)
list = serialdevname
```

**Description**

`serialdevname(n)` returns the name of the `n`:th serial device which can be opened by `serialdevopen`. Argument `n` must be 1 or higher; with other values, such as those larger than the number of serial devices available on your computer, `serialdevname` returns the empty string.

Without input argument, `serialdevname` gives the list of serial device names.

**Examples**

On a Macintosh with internal modem:

```
serialdevname(1)
  Internal Modem
```

Under Windows:

```
serialdevname(1)
  COM1
```

**See also**

`serialdevopen`

**serialdevopen**

Open a serial port.

**Syntax**

```
fd = serialdevopen(portname, options)
fd = serialdevopen(portname)
```

**Description**

`serialdevopen(portname)` opens a connection to the serial port whose name is `portname` and returns a file descriptor `fd`. Names depend on the operating system and can be obtained with `serialdevname`.

Some platforms do not provide a complete list of all ports; `serialdevopen` may accept additional device names and pass them directly to the corresponding function of the operating system.

The second argument of `serialdevopen(portname,options)` is a structure which contains configuration settings. It is set with `serialdevset`.

Once a connection has been opened, the file descriptor `fd` can be used with functions such as `fread`, `fwrite`, `fscanf`, and `fprintf`. The connection is closed with `fclose`.

### Example

```
fd = serialdevopen(serialdevname(1), ...
    serialdevset('BPS',19200,'TextMode',true,'Timeout',2));
fprintf(fd, 'L,%d,2\n', 1);
reply = fgetl(fd)
fclose(fd);
```

### See also

`fclose`, `serialdevname`, `serialdevset`, `fflush`, `fread`, `fwrite`, `fscanf`, `fgetl`, `fgets`, `fprintf`

## serialdevset

Configuration settings for serial port.

### Syntax

```
options = serialdevset
options = serialdevset(name1, value1, ...)
options = serialdevset(options0, name1, value1, ...)
```

### Description

`serialdevset(name1,value1,...)` creates the option argument used by `serialdevopen`. 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, `serialdevset` creates a structure with all the default settings. Note that `serialdevopen` also interprets the lack of an option argument, or the empty array `[]`, as a request to use the default values.

When its first input argument is a structure, `serialdevset` adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
BPS	19200	bit per seconds
Delay	0	delay after character output in seconds
Handshake	false	hardware handshake
StopBits	2	number of stop bits (1, 1.5, or 2)
TextMode	false	text mode
Timeout	1	timeout in seconds

Output operations wait for the specified delay after each character; this can be useful with slow devices without handshake.

When text mode is set, input CR and CR/LF sequences are converted to LF. Output CR and LF are not converted.

Depending on the platform, operations which use the timeout value (such as input) can be interrupted with the platform-dependent abort key(s) (typically Escape or Control-C) or are limited to 10 seconds.

### Example

```
serialdevset
  BPS: 19200
  Handshake: false
  StopBits: 2
  TextMode: false
  Timeout: 1
```

### See also

serialdevopen, serialdevname

## Chapter 5

# Libraries

Libraries are collections of functions which complement the set of built-in functions and operators of LME. To use them, type (or add at the beginning of the libraries 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.

`classes` `classes` implements constructors and methods for polynomial and rational functions. With them, you can use standard operator notations such as `+` or `*`.

`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.

**ratio** `ratio` implements constructors and methods for rational numbers based on long integers. Standard arithmetic and boolean operators can be used.

**sigenc** `sigenc` implements functions related to signal encoding to and decoding from a digital representation.

**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.

**wsserver** `wsserver` implements a server for web services using the XML-RPC or SOAP protocol. It requires the TCP/IP and WebServices functions.

## 5.1 **stdlib**

`stdlib` is a library which extends the native LME functions in the following areas:

- creation of matrices: `blkdiag`, `compan`, `hankel`, `linspace`, `logspace`, `toeplitz`
- geometry: `cart2sph`, `cart2pol`, `pol2cart`, `sph2cart`, `subspace`
- functions on integers: `factor`, `isprime`, `primes`
- statistics: `corrcoef`, `median`, `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
```

### **cart2pol**

Cartesian to polar coordinates transform.

## Syntax

```
use stdlib
(phi, r) = cart2pol(x, y)
(phi, r, z) = cart2pol(x, y, z)
```

## Description

$(\text{phi}, r) = \text{cart2pol}(x, y)$  transforms Cartesian coordinates  $x$  and  $y$  to polar coordinates  $\text{phi}$  and  $r$  such that  $x = r \cos(\varphi)$  and  $y = r \sin(\varphi)$ .

$(\text{phi}, r, z) = \text{cart2pol}(x, y, z)$  transform Cartesian coordinates to cylindrical coordinates, leaving  $z$  unchanged.

## Example

```
use stdlib
(phi, r) = cart2pol(1, 2)
phi =
  1.1071
r =
  2.2361
```

## See also

[cart2sph](#), [pol2cart](#), [sph2cart](#)

## cart2sph

Cartesian to spherical coordinates transform.

## Syntax

```
use stdlib
(phi, theta, r) = cart2sph(x, y, z)
```

## Description

$(\text{phi}, \text{theta}, r) = \text{cart2sph}(x, y, z)$  transforms Cartesian coordinates  $x$ ,  $y$ , and  $z$  to polar coordinates  $\text{phi}$ ,  $\text{theta}$ , and  $r$  such that  $x = r \cos(\varphi) \cos(\vartheta)$ ,  $y = r \sin(\varphi) \cos(\vartheta)$ , and  $z = r \sin(\vartheta)$ .

**Example**

```
use stdlib
(phi, theta, r) = cart2sph(1, 2, 3)
phi =
  1.1071
theta =
  0.9303
r =
  3.7417
```

**See also**

cart2pol, pol2cart, sph2cart

**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`

**companion**

Companion matrix.

**Syntax**

```
use stdlib
X = companion(pol)
```

**Description**

`companion(pol)` gives the companion matrix of polynomial `pol`, a square matrix whose eigenvalues are the roots of `pol`.

**Example**

```
use stdlib
companion([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.

**Syntax**

```
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 numerical 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`

## factor

Prime factors.

## Syntax

```
use stdlib
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.

**Example**

```
use stdlib
factor(350)
  2 5 5 7
```

**See also**

isprime

**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.

**Syntax**

```
use stdlib
Y = filter2(F, X)
Y = filter2(F, X, shape)
```

**Description**

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 equally-spaced 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.

**Syntax**

```

use stdlib
Y = ifftshift(X)

```

**Description**

`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

**isprime**

Prime number test.

**Syntax**

```
use stdlib
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**

```
use stdlib
isprime(7)
true
isprime([0, 2, 10])
F T F
```

**See also**

primes, factor

**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

**linspace**

Sequence of linearly-spaced elements.

**Syntax**

```

use stdlib
v = linspace(x1, x2)
v = linspace(x1, x2, n)

```

**Description**

`linspace(x1,x2)` produces a row vector of 100 values spaced linearly from `x1` and `x2` inclusive. With a third argument, `linspace(x1,x2,n)` produces a row vector of `n` values.

**Examples**

```

use stdlib
linspace(1,10)
  1.0000 1.0909 1.1818 ... 9.9091 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**

```
use stdlib
v = logspace(x1, x2)
v = logspace(x1, x2, n)
```

**Description**

`logspace(x1,x2)` produces a row vector of 100 values spaced logarithmically from  $10^{x1}$  and  $10^{x2}$  inclusive. With a third argument, `logspace(x1,x2,n)` produces a row vector of  $n$  values.

**Example**

```
logspace(0,1)
1.0000 1.0235 1.0476 ... 9.5455 9.7701 10.0000
```

**See also**

`linspace`, operator :

**median**

Median.

**Syntax**

```
use stdlib
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.

`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**

```
use stdlib
median([1, 2, 5, 6, inf])
5
```

**See also**

mean, sort

**perms**

Array of permutations.

**Syntax**

```
use stdlib
M = perms(v)
```

**Description**

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

**pol2cart**

Polar to Cartesian coordinates transform.

**Syntax**

```
use stdlib
(x, y) = pol2cart(phi, r)
(x, y, z) = pol2cart(phi, r, z)
```

## Description

$(x,y)=\text{pol2cart}(\text{phi},r)$  transforms polar coordinates  $\text{phi}$  and  $r$  to Cartesian coordinates  $x$  and  $y$  such that  $x = r \cos(\varphi)$  and  $y = r \sin(\varphi)$ .

$(x,y,z)=\text{pol2cart}(\text{phi},r,z)$  transforms cylindrical coordinates to Cartesian coordinates, leaving  $z$  unchanged.

## Example

```
use stdlib
(x, y) = pol2cart(1, 2)
x =
  1.0806
y =
  1.6829
```

## See also

`cart2pol`, `cart2sph`, `sph2cart`

## 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.

### Syntax

```
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

`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
```

**See also**

isprime

**sortrows**

Sort matrix rows.

**Syntax**

```
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

**sph2cart**

Spherical to Cartesian coordinates transform.

**Syntax**

```
use stdlib
(x, y, z) = sph2cart(phi, theta, r)
```

**Description**

$(x,y,z)=\text{sph2cart}(\phi,\theta,r)$  transforms polar coordinates  $\phi$ ,  $\theta$ , and  $r$  to Cartesian coordinates  $x$ ,  $y$ , and  $z$  such that  $x = r \cos(\phi) \cos(\theta)$ ,  $y = r \sin(\phi) \cos(\theta)$ , and  $z = r \sin(\theta)$ .

**Example**

```
use stdlib
(x, y, z) = sph2cart(1, 2, 3)
x =
  -0.6745
y =
  -1.0505
z =
  2.7279
```

**See also**

`cart2pol`, `cart2sph`, `pol2cart`

**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  $\mathbb{R}^2$ :

```
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  $\mathbb{R}^3$ :

```
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

Numerical integration with trapezoidal approximation.

### Syntax

```
use stdlib  
s = trapz(Y)  
s = trapz(X, Y)  
s = trapz(X, Y, dim)
```

**Description**

trapez(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.

trapez(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
trapez([2, 3, 5])
    6.5
trapez([1, 2, 5], [2, 3, 5])
    14.5
```

**See also**

sum, cumtrapz

**5.2 stat**

stat is a library which adds to LME advanced statistical functions.

The following statement makes available functions defined in stat:

```
use stat
```

**bootstrp**

Bootstrap estimate.

**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.

### 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, iqr

**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)`.

`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.

### Syntax

```
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
```

## See also

`nanmedian`, `nanstd`, `mean`

## nansum

Sum after discarding NaNs.

### Syntax

```
use stat
y = nansum(A)
y = 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

```
use stat
pdist((1:3)')
  1 2 1
squareform(pdist((1:3)'))
  0 1 2
  1 0 1
  2 1 0
squareform(pdist([1,2,6; 3,1,7;6,1,2]))
  0      2.4495    6.4807
  2.4495    0      5.831
  6.4807    5.831    0
```

## 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.

**Example**

```
prctile(rand(1,1000),90)
0.8966
```

**See also**

trimmean, iqr

**range**

Mean absolute deviation.

**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.

**Syntax**

```
use stat
D = squareform(d)
```

**Description**

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).

**Syntax**

```
use stat
Y = zscore(X)
Y = zscore(X, dim)
```

**Description**

`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.

**5.3 classes**

Library `classes` 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 classes:

```
use classes
```

**`polynom::polynom`**

Polynom object constructor.

**Syntax**

```
use classes
a = polynom
a = polynom(coef)
```

**Description**

`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.

```
- minus      + plus
^ mpower     rem
\ mldivide   roots
/ mrdivide   - uminus
mtimes       + uplus
```

**Examples**

```

use classes
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 classes
disp(a)

```

**Description**

disp(a) displays polynomial a. It is also executed implicitly when LME displays the polynom result of an expression which does not end with a semicolon.

**Example**

```

use classes
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 classes
coef = double(a)
```

## Description

`double(a)` converts polynomial `a` to a row vector of descending-power coefficients.

## Example

```
use classes
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.

## Syntax

```
use classes
subst(a, b)
```

## Description

`subst(a, b)` substitutes the variable of `polynom a` with `polynom b`.

## Example

```
use classes
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 classes
diff(a)
```

**Description**

diff(a) differentiates polynomial a.

**Example**

```
use classes
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 classes
int(a)
```

**Description**

int(a) integrates polynomial a.

**Example**

```
use classes
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 classes
fun = inline(a)
```

**Description**

`inline(a)` converts polynomial `a` to an inline function which can then be used with functions such as `feval` and `ode45`.

**Example**

```
use classes
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 classes
y = feval(a, x)
```

**Description**

`feval(a,x)` evaluates polynomial `a` 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 classes
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`

**ratfun::ratfun**

Ratfun object constructor.

**Syntax**

```
use classes
a = ratfun
a = ratfun(coefnum)
a = 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 classes
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 classes
disp(a)

```

### Description

disp(a) displays rational function a. 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 classes
coef = num(a)

```

**Description**

`num(a)` gets the numerator of `a` 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 classes
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 classes
diff(a)
```

**Description**

`diff(a)` differentiates `ratfun a`.

**Example**

```
use classes
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 classes
fun = inline(a)
```

**Description**

inline(a) converts ratfun a 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 classes
y = feval(a, x)
```

**Description**

feval(a, x) evaluates ratfun a 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 classes
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

**5.4 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
```

**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  $\text{num}/\text{den}$ . 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.

```

==  eq      \  mldivide
>=  ge      ^  mpower
>   gt      /  mrdivide
    inv     *  mtimes
<=  le      ~= ne
<   lt      +  plus
    max     -  uminus
    min     +  uplus
-   minus

```

## Examples

```

use ratio
r = ratio(2, 3)
r =
  2/3
q = 5 * r - 1
q =
  7/3

```

## 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`

## 5.5 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
```

### **bitfield::beginning**

First bit position in a bitfield.

#### **Syntax**

```
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 if `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.

<code>&amp;</code>	and	<code>~</code>	not
<code>==</code>	eq	<code> </code>	or
<code>-</code>	minus	<code>+</code>	plus
<code>\</code>	mldivide	<code>-</code>	uminus
<code>/</code>	mrdivide	<code>+</code>	uplus
	mtimes		xor
<code>~=</code>	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

```
use bitfield
a = bitfield(123, 16)
a =
    0b0000000001111011
b = ~a
b =
    0b111111110000100
b = a * 5
b =
    0b0000001001100111
```

## 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.

**Syntax**

```
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

```
use bitfield
a = bitfield(9, 4);
a =
  0x1001
i = int8(a)
i =
  210
b = bitfield(i)
b =
  0b11111001
```

## 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`

## 5.6 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`, `cheb1ap`, `cheb2ap`, 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 band-stop 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`.

## besselap

Bessel analog filter prototype.

### Syntax

```
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  $n$ th-order digital low-pass filter 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 2 $n$ th-order digital bandpass filter with pass-band between `wl` and `wh` relatively to half the sampling frequency.

`besself(n,w0,'high')` gives a  $n$ th-order digital highpass filter 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 2 $n$ th-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', `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 transformation  $z_d = (1 + z_c/2f_s)/(1 - z_c/2f_s)$ . 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, besslap, cheblap, 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 low-pass 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 pass-band between wl and wh relatively to half the sampling frequency.

`butter(n, w0, 'high')` gives a  $n$ th-order digital highpass filter with a cutoff frequency of  $w_0$  relatively to half the sampling frequency.

`butter(n, [wl, wh], 'stop')`, where the second input argument is a vector of two numbers, gives a 2 $n$ th-order digital bandstop filter with stopband between  $w_l$  and  $w_h$  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.

### Syntax

```
use filter
(z, p, k) = cheb1ap(n, rp)
```

### Description

`cheb1ap(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.

### 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.

## See also

`cheby1`, `cheblap`, `ellipap`, `besselap`, `buttap`

## cheby1

Chebyshev type 1 filter.

## Syntax

```
use filter
(z, p, k) = cheby1(n, w0)
(num, den) = cheby1(n, w0)
(...) = cheby1(n, [wl, wh])
(...) = cheby1(n, w0, 'high')
(...) = cheby1(n, [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,w0)`, where  $w_0$  is a scalar, gives a  $n$ th-order digital low-pass filter with a cutoff frequency of  $w_0$  relatively to half the sampling frequency.

`cheby1(n,[wl,wh])`, where the second input argument is a vector of two numbers, gives a  $2n$ th-order digital bandpass filter with passband between  $w_l$  and  $w_h$  relatively to half the sampling frequency.

`cheby1(n,w0,'high')` gives a  $n$ th-order digital highpass filter with a cutoff frequency of  $w_0$  relatively to half the sampling frequency.

`cheby1(n,[wl,wh],'stop')`, where the second input argument is a vector of two numbers, gives a  $2n$ th-order digital bandstop filter with stopband between  $w_l$  and  $w_h$  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, w0)
(num, den) = cheby2(n, w0)
(...) = cheby2(n, [wl, wh])
(...) = cheby2(n, w0, 'high')
(...) = cheby2(n, [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 numerator and denominator coefficient vectors if there are two output arguments.

cheby2(n, w0), where w0 is a scalar, gives a nth-order digital low-pass filter with a cutoff frequency of w0 relatively to half the sampling frequency.

cheby2(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.

cheby2(n, w0, 'high') gives a nth-order digital highpass filter with a cutoff frequency of w0 relatively to half the sampling frequency.

cheby2(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', 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, w0)
(num, den) = ellip(n, w0)
(...) = ellip(n, [wl, wh])
(...) = ellip(n, w0, 'high')
(...) = ellip(n, [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,w0)`, where `w0` is a scalar, gives a `n`th-order digital low-pass filter with a cutoff frequency of `w0` relatively to half the sampling frequency.

`ellip(n, [wl,wh])`, where the second input argument is a vector of two numbers, gives a 2`n`th-order digital bandpass filter with passband between `wl` and `wh` relatively to half the sampling frequency.

`ellip(n,w0,'high')` gives a `n`th-order digital highpass filter with a cutoff frequency of `w0` relatively to half the sampling frequency.

`ellip(n,[wl,wh],'stop')`, where the second input argument is a vector of two numbers, gives a 2`n`th-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.

**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  $w_0$  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  $w_0$ . `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

$$F(s) = F_0 \left( \frac{1}{\omega_0 s} \right)$$

where  $F_0(s)$  is the filter prototype.

**See also**

lp2lp, lp2bp, lp2bs

**lp2lp**

Lowpass prototype to lowpass filter conversion.

**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  $w_0$ .  $lp2lp(z_0, p_0, k_0, w_0)$  converts a filter given by its zeros, poles, and gain;  $lp2lp(num_0, den_0, w_0)$  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

**5.7 lti**

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 state-space 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
```

## **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 continuous-time state-space model

$$\begin{aligned}x'(t) &= A x(t) + B u(t) \\ y(t) &= C x(t) + D u(t)\end{aligned}$$

`ss(A,B,C,D,Ts)` creates an LTI object which represents the discrete-time state-space model with sampling period  $T_s$

$$\begin{aligned}x(k+1) &= A x(k) + B u(k) \\ y(k) &= C x(k) + D u(k)\end{aligned}$$

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)
sc =
  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
A =
  -2   -3   -4
   1    0    0
   0    1    0
B =
   1
   0
   0
C =
   0    0    1
D =
   0

```

**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 continuous-time 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` may 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:

```
sd = tf({0.1; 0.15}, {[1, -0.8]; [1; -0.78]}, 1e-3)
sd =
  discrete-time transfer function, Ts=1e-3
  y1/u1: 0.1/(s-0.8)
  y2/u1: 0.15/(s-0.78)
```

## See also

`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  $[u_1;u_2]$  and outputs  $[y_1;y_2]$ , where  $u_1$  and  $u_2$  are the inputs of  $a_1$  and  $y_1$  and  $y_2$  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`.

## 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`

## lti::d2c

Conversion from discrete time to continuous time.

## Syntax

```
use lti
b = d2c(a)
b = d2c(a, method)
```

## Description

`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`.

## See also

`lti::c2d`, `d2cm`

## 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 =
    0
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                0.5
    (:,:,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

**ss::ctrb**

Controllability matrix.

**Syntax**

```

use lti
C = ctrb(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::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::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`

**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 continuous-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

**lti::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.

**Syntax**

```
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`

## **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.

## **lti::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  $\text{inv}(a)*b$ .

**See also**

`lti::mrdivide`, `lti::times`, `lti::inv`

**lti::mrdivide**

System right division.

**Syntax**

```
use lti
c = a / b
c = mrdivide(a, b)
```

**Description**

$a/b$  is equivalent to  $a*\text{inv}(b)$ .

**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
O = 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::plus

System sum.

### Syntax

```
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::series**

Series connection.

**Syntax**

```
use lti
c = series(a, b)
c = series(a, b, outa, inb)
```

**Description**

`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::repmat**

Replicate a system.

**Syntax**

```
use lti
b = repmat(a, n)
b = repmat(a, [m,n])
b = repmat(a, m, n)
```

**Description**

`repmat(a,n)`, when `a` is a transfer function or a matrix of transfer functions, creates a new system described by a matrix of transfer functions where `a` is repeated `n` times horizontally and vertically. If `a` is a state-space system, matrices `B`, `C`, and `D` are replicated to obtain the same effect.

`repmat(a,[m,n])` or `repmat(a,m,n)` repeats matrix `a` `m` times vertically and `n` times horizontally.

**See also**`lti::append`**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.

**Syntax**

```
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, gives the four matrices of the state-space model, and optionally the sampling period or the empty array `[]` for continuous-time systems.

**See also**`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 right-hand 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 both, `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. 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.

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 both, `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) = ssdata(a)

```

**Description**

`tfdata(a)`, where `a` is any kind of LTI object, 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**

`lti:ssdata`

**lti::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`

**lti::uplus**

Negative.

**Syntax**

```
use lti
b = +a
b = uplus(a)
```

**Description**

+a gives a.

**See also**

`lti::uminus`, `lti::plus`

## zpk

LTI transfer function constructor using zeros and poles.

### Syntax

```

use lti
a = zpk(z, p, k)
a = zpk(zeroslist, poleslist, gainlist)
a = zpk(..., Ts)
a = zpk(..., Ts, var)
a = zpk(..., b)
a = zpk(b)

```

### Description

zpk creates transfer-function LTI systems like `tf::tf`. Instead of using transfer function coefficients as input, it accepts a vector of zeros, a vector of poles, and a gain for a simple-input simple-output (SISO) system; or lists of sublists of zeros, poles and gains for multiple-input multiple-output (MIMO) systems.

### Examples

```

use lti
sd = zpk(0.3, [0.8+0.5j; 0.8-0.5j], 10, 0.1)
sd =
  discrete-time transfer function, Ts=0.1
  (10z-3)/(z^2-1.6z+0.89)

```

### See also

`tf::tf`

### Iti::bodemag

Magnitude of the Bode plot.

### Syntax

```

use lti
bodemag(a, ...)
... = bodemag(a, ...)

```

### Description

`bodemag(a)` plots the magnitude of the Bode diagram of system `a`.

**See also**

`lti::bodephase`, `lti::nichols`, `lti::nyquist`

**lti::bodephase**

Phase of the Bode plot.

**Syntax**

```
use lti
bodephase(a, ...)
... = bodephase(a, ...)
```

**Description**

`bodephase(a)` plots the magnitude of the Bode diagram of system `a`.

**See also**

`lti::bodemag`, `lti::nichols`, `lti::nyquist`

**lti::impulse**

Impulse response.

**Syntax**

```
use lti
impulse(a, ...)
... = impulse(a, ...)
```

**Description**

`impulse(a)` plots the impulse response of system `a`.

**See also**

`lti::step`, `lti::lsim`, `lti::initial`

**lti::initial**

Time response with initial conditions.

**Syntax**

```
use lti
initial(a, x0, ...)
... = initial(a, x0, ...)
```

**Description**

`initial(a,x0)` plots the time response of state-space system `a` with initial state `x0` and null input.

**See also**

`lti::impulse`, `lti::step`, `lti::lsim`

**lti::lsim**

Time response.

**Syntax**

```
use lti
lsim(a, u, t, ...)
... = lsim(a, u, t)
```

**Description**

`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.

**See also**

`lti::impulse`, `lti::step`, `lti::initial`

**lti::nichols**

Nichols plot.

**Syntax**

```
use lti
nichols(a, ...)
... = nichols(a, ...)
```

**Description**

`nichols(a)` plots the Nichols diagram of system `a`.

**See also**

`lti::nyquist`, `lti::bodemag`, `lti::bodephase`

**lti::nyquist**

Nyquist plot.

**Syntax**

```
use lti
nyquist(a, ...)
... = nyquist(a, ...)
```

**Description**

`nyquist(a)` plots the Nyquist diagram of system `a`.

**See also**

`lti::nichols`, `lti::bodemag`, `lti::bodephase`

**lti::step**

Step response.

**Syntax**

```
use lti
step(a, ...)
... = step(a, ...)
```

**Description**

`step(a)` plots the step response of system `a`.

**See also**

`lti::impulse`, `lti::lsim`, `lti::initial`

## 5.8 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
```

### 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.

#### Syntax

```
use sigenc
output = alawexpand(input)
output = alawexpand(input, a)
```

**Description**

`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  $k$  is

$$y^*(k) = \sum_{i=1}^{\text{degpredictor}} \text{predictor}_i \cdot y_q(k-i)$$

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 <sup>1</sup>, 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>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.

```

(partition_opt, codebook_opt) = lloyds(in, codebook)
    partition_opt =
        -2.9  -0.5   1.3
    codebook_opt =
        -4.1  -1.7   0.4   2.2
(i, out, dist) = quantiz(in, partition_opt, codebook_opt);
dist
    1.0543

```

## 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 look-up 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)
    i =
        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,a)` 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.

**Syntax**

```

use sigenc
output = ulawexpand(input)
output = ulawexpand(input, mu)

```

**Description**

`ulawexpand(input, a)` 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`

**5.9 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
```

**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;
t = (0:sr)' / sr;
s = sin(2 * pi * 740 * t);
wavwrite(map2int(s, -1, 1, 'int16'), sr, 16, 'beep.wav');
```

**See also**

wavread

## 5.10 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
```

### 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:

<b>Sequence</b>	<b>Replaced with</b>
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)
yy	year (2 digits)
yyyy	year (4 digits)

If the sequence PM is found, the hour is between 1 and 12; otherwise, between 0 and 23.

## 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
```

## 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:

<b>Example</b>	<b>Value</b>
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

```
use date
datevec('Date and time: 20050418T162603')
  2005  4  18  16  26  3
datevec('03:57 PM')
  0  0  0  15  57  0
datevec('01-Aug-1291')
  1291  8  1  0  0  0
datevec('At 16:30 on 11/04/07')
  2007  11  4  16  30  0
```

## 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

## 5.11 constants

constants is a library which defines physical constants in SI units (meter, kilogram, second, ampere).

The following statement makes available constants defined in constants:

```
use constants;
```

The following constants are defined:

<b>Name</b>	<b>Value</b>	<b>Unit</b>
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	C
electron_mass	9.1093897e-31	kg
faraday_constant	9.6485309e4	C/mole
gravitational_constant	6.672659e-11	N m <sup>2</sup> /kg <sup>2</sup>
gravity_acceleration	9.80655	m/s <sup>2</sup>
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 <sup>3</sup> /mole
muon_mass	1.8835327e-28	kg
neutron_mass	1.6749286e-27	kg
plank_constant	6.6260755e-34	J s
plank_constant_reduced	1.0545727e-34	J s
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	5.67051e-8	W/m <sup>2</sup> K <sup>-4</sup>
vacuum_permittivity	8.854187817e-12	A s/V m

## Chapter 6

# Extensions

Extensions are additional functions, usually developed in C or Fortran, which extend the core functionality of LME. Extensions are grouped in so-called *dynamically-linked libraries* (DLL) files. At startup, LME loads all extensions it finds in the folder LMExt in the same location as the LME program file. Each extension initializes itself and usually displays a line of information in the Command window. No further action is needed in order to use the new functions.

Note that the current release includes some extensions inside the EXE application itself.

You can also develop and add your own extensions, as explained in the next chapter.

Here is the list of the extensions currently provided with LME for Pocket PC.

### Mathematics

**Long integers (Mac, Windows, Unix)** Arithmetic on arbitrary-length integer numbers.

### File input/output and data compression

**Memory mapping (Mac OS X, Unix)** Mapping of files in memory, which can be read and written like regular arrays.

**Data compression (Mac, Windows, Unix)** Support for compressing and uncompressing data using ZLib.

**Image Input/Output (Mac, Windows, Unix)** Support for reading and writing arrays as PNG or JPEG image files.

**MAT-file (Mac, Windows, Unix)** Support for reading and writing MAT-files (native MATLAB binary files).

## Operating system

**Download URL (Mac, Windows, Linux)** Download of documents from the World Wide Web.

**Launch URL (Mac, Windows)** Opening of documents in a World Wide Web browser.

**Web Services (Windows, Mac OS X, Unix)** Web Services (standard remote procedure calls using XML-RPC and SOAP).

## Hardware support

**Audio playback (Mac OS X, Windows, Linux)** Audio output.

**Audio recording (Mac OS X, Windows)** Audio input.

## 6.1 Long Integers

This section describes functions which support long integers (*longint*), i.e. integer numbers with an arbitrary number of digits limited only by the memory available. Some LME functions have been overloaded: new definitions have been added and are used when at least one of their arguments is of type *longint*. These functions are listed in the table below.

LME	Operator	Purpose
abs		absolute value
char		conversion to string
disp		display
double		conversion to floating-point
gcd		greatest common divisor
lcm		least common multiple
minus	-	subtraction
mldivide	\	left division
mpower	^	power
mrdivide	/	right division
mtimes	*	multiplication
plus	+	addition
rem		remainder
uminus	-	negation
uplus	+	no operation

### longint

Creation of a long integer.

**Syntax**

```
li = longint(i)
li = longint(str)
```

**Description**

`longint(i)` creates a long integer from a native LME floating-point number. `longint(str)` creates a long integer from a string of decimal digits.

**Examples**

```
longint('1234567890')
1234567890
longint(2)^100
1267650600228229401496703205376
```

13th Mersenne prime:

```
longint(2)^521-1
6864797660130609714981900799081393217269
4353001433054093944634591855431833976560
5212255964066145455497729631139148085803
7121987999716643812574028291115057151
```

Number of decimal digits in the 27th Mersenne prime:

```
length(char(longint(2)^44497-1))
13395
```

## 6.2 Memory Mapping

This section describes functions which offer support for file memory mapping. Once a file is mapped in memory, its elements (bytes, 16-bit words or 32-bit words) can be accessed like a normal array, thanks to virtual memory.

These functions are available only on Unix (or Unix-like) systems, such as Mac OS X.

**mmap**

Map a file in memory.

## Syntax

```
m = mmap(filename, n)
m = mmap(filename, n, type)
m = mmap(filename, n, type, perm)
m = mmap(filename, n, type, perm, offset)
```

## Description

`mmap(filename,n)` maps in memory the `n` first bytes of file whose name is given by string `filename`. It returns an object which can be used to read bytes with regular array indexing, with the first byte at offset 0. The file is created if necessary.

`mmap(filename,n,type)` specifies the type of the elements. `type` is one of the strings in the table below.

Type	Range	Description
'uint8'	0 - 255	unsigned byte
'char'	char(0) - char(255)	character
'uint16'	0 - 65535	unsigned 16-bit word
'uint32'	0 - 4294967295	unsigned 32-bit word
'int8'	-128 - 127	signed byte
'int16'	-32768 - 32767	signed 16-bit word
'int32'	-2147483648 - 2147483647	signed 32-bit word

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).

`mmap(filename,n,type,perm)` specifies permission with string `perm`, which takes one of the values in the table below.

Perm	Description
'r'	read-only
'w'	read/write

`mmap(filename,n,type,perm,offset)` specified the offset of the part being memory-mapped in the file.

The following functions are overloaded to accept the type of objects returned by `mmap`: `beginning`, `disp`, `end`, `length`, `subsasgn`, and `subsref`.

## Example

Bytes 0-3999 of file `'test'` are mapped in memory as 32-bit signed integers. They are multiplied by two.

```
m = mmap('test', 1000, 'int32', 'w');
m(0:999) = 2 * m(0:999);
```

```
unmap(m);
```

**See also**

`munmap`, `beginning`, `disp`, `end`, `length`, `subsasgn`, `subsref`

**munmap**

Unmap a memory-mapped file.

**Syntax**

```
munmap(m)
```

**Description**

`munmap(m)` unmaps a file which has been mapped with `mmap`. Its argument is the object given by `mmap`.

**See also**

`mmap`

## 6.3 Data Compression

This section describes functions which compress and uncompress sequences of bytes, such as text. Often, these sequences present redundancy which can be removed to produce a shorter sequence, while still being able to revert to the initial one.

The ZLib extension is based on `zlib` by J.L. Gailly and M. Adler, whose work is gratefully acknowledged. To preserve their terminology, compression is performed with function `deflate`, and uncompression with `inflate`.

**deflate**

Compress a sequence of bytes.

**Syntax**

```
strc = deflate(str)
```

## Description

`deflate(str)` produces a string `strc` which is usually shorter than its argument `str`. String `str` can be reconstructed with `inflate` using only `strc`. `deflate` and `inflate` process any sequence of bytes (8-bit words); their input argument can be any array. However, their shape and their type are lost (the result of `inflate` is always a row vector of characters) and their elements are restored modulo 256.

Depending on the data, compression rates of 2 or more are typical. Sequences without redundancy (such as random numbers or the result of `deflate`) can produce a result slightly larger than the initial sequence.

## Example

```
str = repmat('abcd ef ', 1, 1000);
length(str)
    8000
strc = deflate(str);
length(strc)
    43
str = repmat('abcd ef ', 1, 1000);
strc = deflate(str);
str2 = inflate(strc);
str == str2
    true
```

To compress objects which are not sequence of bytes, you can use `dumpvar` and `str2obj` to convert them to and from a textual representation:

```
A = repmat(600, 2, 2)
A =
    600    600
    600    600
inflate(deflate(A))
    1x4 uint8 array
    88    88    88    88
str = dumpvar(A);
str2obj(deflate(inflate(str)))
    600    600
    600    600
```

## See also

`inflate`

## **inflate**

Uncompress the result of deflate.

### **Syntax**

```
str = inflate(strc)
```

### **Description**

`inflate(strc)` uncompresses `strc` to undo the effect of `compress`. The output is always a character string with one row, whose characters are coded on one byte.

### **See also**

`deflate`

## **zread**

Read deflated data and uncompress them.

### **Syntax**

```
(data, nin) = zread(fd, n)  
(data, nin) = zread(fd)
```

### **Description**

`zread(fd, n)` reads up to `n` bytes from file descriptor `fd`, uncompresses them using the inflate algorithm, and returns the result as a row vector of type `uint8`. An optional second output argument is set to the number of bytes which have actually been read; it is less than `n` if the end-of-file is reached.

With a single input argument, `zread(fd)` reads data until the end of the file.

Note that you must read a whole segment of deflated data with one call. Inflation is restarted every time `zread` is called.

### **See also**

`zwrite`, `inflate`

## **zwrite**

Compress a sequence of bytes and write the result.

### **Syntax**

```
nout = zwrite(fd, data)
```

### **Description**

`zwrite(fd, data)` compresses the array data, of type `int8` or `uint8`, and writes the result to the file descriptor `fd`.

Note that you must write a whole segment of data with one call. Deflation is restarted every time `zwrite` is called.

### **See also**

`zread`, `deflate`

## **6.4 Image Files**

This section describes functions which offer support for reading and writing image files. Formats supported include PNG and JPEG.

Calerga gratefully acknowledges the following contributions: PNG encoding and decoding are based on `libpng`; and JPEG encoding and decoding are based on the work of the Independent JPEG Group.

## **imageread**

Read an image file.

### **Syntax**

```
A = imageread(fd)
```

### **Description**

`imageread(fd)` reads a PNG or JPEG file from file descriptor `fd` and returns it as an array whose first dimension is the image height and second dimension the image width. Grayscale images give a third dimension equal to 1 (i.e. plain matrices). Color images give a third dimension equal to 3; first plane is the red component, second plane the green component, and third plane the blue component. In both cases, value range is 0 for black to 1 for maximum intensity.

The file descriptor is usually obtained by opening a file with `fopen` in binary mode (text mode, with end-of-line translation, would produce garbage or cause a decoding error).

**Example**

```
fd = fopen('image.png', 'r');
im = imageread(fd);
fclose(fd);
```

**See also**

`imagemwrite`

**imageset**

Options for image output.

**Syntax**

```
options = imageset
options = imageset(name1, value1, ...)
options = imageset(options0, name1, value1, ...)
```

**Description**

`imageset(name1,value1,...)` creates the option argument used by `imagemwrite`. 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, `imageset` creates a structure with all the default options. Note that `imagemwrite` also interpret the lack of an option argument, or the empty array `[]`, as a request to use the default values.

When its first input argument is a structure, `imageset` adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
Type	'PNG'	'PNG' or 'JPG'/'JPEG'
Quality	80	JPEG quality (0=worst,100=best)
Progressive	false	true to permit progressive decoding

## Examples

Default options:

```
imageset
  Type: 'png'
  Quality: 80
  Progressive: false
```

Writing the contents of array A into a small, low-quality JPEG file:

```
fd = fopen('A.jpg', 'w');
imagemwrite(fd, A, imageset('Type','JPG','Quality',20));
fclose(fd);
```

## See also

`imagemwrite`

## imagemwrite

Write an image file.

## Syntax

```
imagemwrite(fd, A)
imagemwrite(fd, A, options)
```

## Description

`imagemwrite(fd,A)` writes array A to a PNG file specified by file descriptor `fd`. Image A is an array whose first dimension is the image height and second dimension the image width. Grayscale images have their third dimension equal to 1 (i.e. they are plain matrices). Color images have a third dimension equal to 3; first plane is the red component, second plane the green component, and third plane the blue component. In both cases, value range is 0 for black to 1 for maximum intensity. Values outside this range are clipped.

`imagemwrite(fd,A,options)` uses structure options to specify image file options. Options are usually created with function `imageset`; they include the file type.

The file descriptor is usually obtained by opening a file with `fopen` in binary mode (text mode, with end-of-line translation, would produce a corrupted image file).

### Example

Write the image contained in the matrix `im` to a file "image.png", using the default options.

```
fd = fopen('image.png', 'w');
imagewrite(fd, im);
fclose(fd);
```

Write the same image as a JPEG file.

```
fd = fopen('image.jpg', 'w');
imagewrite(fd, im, imageset('Type', 'JPEG'));
fclose(fd);
```

### See also

`imageset`, `imagerread`

## 6.5 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, numerical, 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, numerical, 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

## 6.6 SQLite

This section describes functions which SQLite relational databases. SQLite is a public-domain relational database stored locally in a single file, which uses SQL as its query language. There are two main advantages of SQLite with respect to larger relational database systems: there is no need to install any additional software or to have access to a remote database, and the database file can be archived and restored extremely easily. On the other hand, it lacks concurrent access, stored procedures, etc. Its SQL compatibility permits the developer to port easily applications to other databases, should it be necessary.

This documentation assumes you have a basic knowledge of SQL. Even if you do not, the examples should help you to get started. For more informations about SQLite, please visit the Web site <http://www.sqlite.org>.

The creator of SQLite, D. Richard Hipp, is gratefully acknowledged.

The following functions are defined.

<b>Function</b>	<b>Purpose</b>
sqlite_changes	Number of affected rows in the last command
sqlite_close	Close an SQLite database
sqlite_exec	Execute an SQL query
sqlite_last_insert_rowid	Index of the last row inserted
sqlite_open	Open an SQLite database
sqlite_set	Options for sqlite_open
sqlite_version	Get the version of SQLite

## 6.7 Compiling the extension

The extension is installed with Sysquake or LME and ready to use; but it is also provided as source code. If you want, you can check on the Web if there is a more recent version of SQLite and compile the extension again with it. The steps below show the simplest way to do it.

**Check your development tools** Make sure you have the development tools required for compiling the extension. Typically, you need a C compiler chain like gcc. You can get it as free software from GNU.

**Get SQLite distribution** Download the latest distribution from the site <http://www.sqlite.org>.

**Locate the required files** To compile the extension, you will need the following files:

- LMESQLite.c, the main source code of the extension which defines new functions for LME.
- LME\_Ext.h, the header file for LME extensions, which is provided with all LME applications which support extensions; it is typically stored in a directory named ExtDevel. Let *extdevel* be its path.
- The source code of SQLite, typically in the directory *src* of the SQLite distribution.

**Compile the extension** Create a new directory, cd to it, and run the Make file of the SQLite extension. For example:

```
$ cd
$ mkdir mysql-build
$ cd mysql_buid
$ ext= extpath sqlite= sqlitepath
  make -f extpath/Makefile.lme-sqlite
```

**Install the extension** For most LME applications, just move or copy the extension (*sqlite.so* if you have used the command above) to the directory where LME looks for extensions (usually *LMEExt*). For Sysquake Remote, you also have to add the following line to the configuration file of Apache (please read Sysquake Remote documentation for more information):

```
SQLLoadExtension extpath/sqlite.so
```

where *extpath/sqlite.so* is the absolute path of the extension.

**Start or restart the LME application** To check that LME has loaded the extension successfully, check the information line starting with *SQLite*. You can also try to evaluate *sqlite\_version*, which should display the version of SQLite.

## 6.8 Functions

### *sqlite\_changes*

Number of affected rows in the last command.

**Syntax**

```
n = sqlite_changes(c)
```

**Description**

`sqlite_changes(c)` gives the number of affected rows in the last UPDATE, DELETE, or INSERT command.

**SQLite call**

```
sqlite3_changes
```

**See also**

```
sqlite_exec, sqlite_last_insert_rowid
```

**sqlite\_close**

Close an SQLite database.

**Syntax**

```
sqlite_close(c)
```

**Description**

`sqlite_close(c)` closes the MySQLite database identified by `c`.

**SQLite call**

```
sqlite3_close
```

**See also**

```
sqlite_open
```

**sqlite\_exec**

Execute an SQL query against an SQLite database.

**Syntax**

```
sqlite_exec(c, query)  
table = sqlite_exec(c, query)
```

**Description**

`sqlite_exec(c,query)` executes a query given in SQL in a string, against the SQLite database identified by `c`. The number of modified rows can be obtained with `sqlite_changes`.

With an output argument, `sqlite_exec` returns the resulting table as a list of rows. Each row is given as a list of column values or as a structure, as specified in the option argument of `sqlite_open` created with `sqlite_set`.

**SQLite call**

`sqlite3_exec`

**See also**

`sqlite_open`, `sqlite_set`, `sqlite_changes`

**sqlite\_last\_insert\_rowid**

Row ID of the last row inserted in a SQLite database.

**Syntax**

```
n = sqlite_last_insert_rowid(c)
```

**Description**

`sqlite_last_insert_rowid(c)` gives the last row inserted by the INSERT command with `sqlite_exec`.

**SQLite call**

`sqlite3_last_insert_rowid`

**See also**

`sqlite_exec`, `sqlite_changes`

**sqlite\_open**

Open an SQLite database.

**Syntax**

```
c = sqlite_open(filename)
c = sqlite_open(filename, options)
```

**Description**

sqlite\_open(filename) opens the database in the specified file. If the file does not exist, a new database is created. The result is an identifier which should be used in all other SQLite calls. The database is closed with sqlite\_close.

sqlite\_open(filename,options) specifies options in the second input argument, which is usually the result of sqlite\_set.

**Example**

```
c = sqlite_open('test.sqlite')
c =
  0
rows = sqlite_exec(c, 'select * from person');
sqlite_close(c);
```

**SQLite calls**

sqlite\_open

**See also**

sqlite\_close, sqlite\_set

**sqlite\_set**

Options for SQLite.

**Syntax**

```
options = sqlite_set
options = sqlite_set(name1, value1, ...)
options = sqlite_set(options0, name1, value1, ...)
```

**Description**

sqlite\_set(name1,value1,...) creates the option argument used by sqlite\_open. 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, `sqlite_set` creates a structure with all the default options. Note that `sqlite_open` also interprets the lack of an option argument, or the empty array [], as a request to use the default values.

When its first input argument is a structure, `sqlite_set` adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options (empty arrays mean "automatic"):

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
<code>ExecResultClass</code>	<code>'list'</code>	row type ('list' or 'struct')
<code>ExecResultNumeric</code>	<code>true</code>	conversion of numeric columns to double

SQLite is usually typeless. If `ExecResultNumeric` is true, columns are converted to numbers of class `double` unless they contain a non-numeric value, or the type name used during declaration contains `BLOB`, `CHAR`, `CLOB`, or `TEXT`. This is the same convention as what SQLite uses itself, for example when sorting rows. `NULL` values are always represented as the (double) empty array [].

## Examples

Default options:

```
sqlite_set
  ExecResultClass: 'list'
  ExecResultNumeric: true
```

## See also

`sqlite_open`

## sqlite\_version

Get the version of SQLite.

## Syntax

```
str = sqlite_version
```

## Description

`sqlite_version` gives the version of SQLite compiled in the extension, as a string. No database is required.

**SQLite call**

sqlite3\_version

**6.9 Sockets**

Socket functions enable communication with a server over TCP/IP. Services which can be accessed via TCP/IP include HTTP (most common protocol for WWW documents and Web services), SMTP (for sending e-mail), POP (for receiving mail), and telnet. Both TCP (where the client and the server are connected and communicate with streams of bytes in both directions) and UDP (connectionless exchange of packets without guarantee of transfer and order) are supported.

Functions described in this section include only those required for opening and configuring the connection. They correspond to fopen for files. Input and output are done with the following generic functions:

<b>Function</b>	<b>Description</b>
fclose	close the file
fgetl	read a line
fgets	read a line
fprintf	write formatted data
fread	read data
fscanf	read formatted data
fwrite	write data
redirect	redirect output

fread does not block if there is not enough data; it returns immediately whatever is available in the input buffer.

**gethostbyname**

Resolve host name.

**Syntax**

ip = gethostbyname(host)

**Description**

gethostbyname(host) gives the IP address of host in dot notation as a string.

**Example**

```
gethostbyname('localhost')
127.0.0.1
```

**See also**

gethostname

**gethostname**

Get name of current host.

**Syntax**

```
str = gethostname
```

**Description**

gethostname gives the name of the current host as a string.

**See also**

gethostbyname

**socketaccept**

Accept a connection request.

**Syntax**

```
fd = socketaccept(fds)
```

**Description**

socketaccept(fds) accepts a new connection requested by a client to the server queue created with socketservernew. Its argument fds is the file descriptor returned by socketservernew.

Once a connection has been opened, the file descriptor fd can be used with functions such as fread, fwrite, fscanf, and fprintf. The connection is closed with fclose.

**See also**

`fclose`, `socketconnect`, `socketservernew`, `fread`, `fwrite`, `fscanf`, `fgetl`, `fgets`, `fprintf`

**socketconnect**

Change UDP connection.

**Syntax**

```
socketconnect(fd, hostname, port)
```

**Description**

`socketconnect(fd,hostname,port)` changes the remote host and port of the UDP connection specified by `fd`. An attempt to use `socketconnect` on a TCP connection throws an error.

**See also**

`socketnew`

**socketnew**

Create a new connection to a server.

**Syntax**

```
fd = socketnew(hostname, port, options)
fd = socketnew(hostname, port)
```

**Description**

`socketnew(hostname,port)` creates a new TCP connection to the specified hostname and port and returns a file descriptor `fd`.

The third argument of `socketnew(hostname,port,options)` is a structure which contains configuration settings. It is set with `socketset`.

Once a connection has been opened, the file descriptor `fd` can be used with functions such as `fread`, `fwrite`, `fscanf`, and `fprintf`. The connection is closed with `fclose`.

## Example

```
fd = socketnew('www.somewebsserver.com', 80, ...
              socketset('TextMode',true));
fprintf(fd, 'GET %s HTTP/1.0\n\n', '/');
reply = fgets(fd)
      reply =
      HTTP/1.1 200 OK
fclose(fd);
```

## See also

`fclose`, `socketset`, `socketconnect`, `socketservernew`, `fread`, `fwrite`, `fscanf`, `fgetl`, `fgets`, `fprintf`

## socketservernew

Create a new server queue for accepting connections from clients.

## Syntax

```
fds = socketservernew(port, options)
fds = socketservernew(port)
```

## Description

`socketservernew(hostname, port)` creates a new TCP or UDP socket for accepting incoming connections. Connections from clients are accepted with `socketaccept`, which must provide as input argument the file descriptor returned by `socketservernew`. Using multiple threads, multiple connections can be accepted on the same port, using multiple `socketaccept` for one `socketservernew`.

The second argument of `socketservernew(port,options)` is a structure which contains configuration settings. It is set with `socketset`. Options are inherited by the connections established with `socketaccept`. On platforms where administrator authorizations are enforced, only an administrator account (root account) can listen to a port below 1024. Only one server can listen to the same port.

To stop listening to new connections, the socket is closed with `fclose`. The file descriptor returned by `socketservernew` can be used only with `socketaccept` and `fclose`.

## Example

```
fds = socketservernew(8080);
fd = socketaccept(fds);
```

```

request = fscanf(fd, 'GET %s');
fprintf(fd, 'Your request is "%s"\n', request);
fclose(fd);
fclose(fds);

```

**See also**

fclose, socketset, socketaccept, socketnew

**socketset**

Configuration settings for sockets.

**Syntax**

```

options = socketset
options = socketset(name1, value1, ...)
options = socketset(options0, name1, value1, ...)

```

**Description**

socketset(name1,value1,...) creates the option argument used by socketnew and socketservernew. 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, socketset creates a structure with all the default settings. Note that socketnew also interprets the lack of an option argument, or the empty array [], as a request to use the default values.

When its first input argument is a structure, socketset adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
ListenQueue	5	queue size for incoming connections
Proto	'tcp'	protocol ('tcp' or 'udp')
TextMode	true	text mode
Timeout	30	timeout in seconds

When TextMode is true, input CR and CR-LF sequences are converted to LF, and output LF is converted to CR-LF, to follow the requirements of many Internet protocols where lines are separated with CR-LF. Note that TextMode is true by default.

**Example**

```
socketset
  ListenQueue: 5
  Proto: 'tcp'
  TextMode: true
  Timeout: 30
```

**See also**

socketnew, socketservernew, socketsetopt

**socketsetopt**

Settings change for sockets.

**Syntax**

```
socketsetopt(fd, name1, value1, ...)
socketsetopt(fd, options)
```

**Description**

`socketsetopt(fd, name1, value1, ...)` changes the options for the socket identified by `fd`. Options are specified by pairs of name and value. They are the same as those valid with `socketset`. However, only 'TextMode' and 'Timeout' have an effect; other ones are ignored.

`socketsetopt(fd, options)` takes as second argument a structure of options created with `socketset`.

**See also**

socketset, socketnew, socketservernew

**6.10 Launch URL**

This section describes a function which requests the default WWW browser to open a URL.

The intended use of `launchurl` is the display of local or Web-based documentation. You can add menu entries to your SQ files to help your users, point to updates, or send e-mail.

## launchurl

Launch a URL in the default browser.

### Syntax

```
status = launchurl(url)
```

### Description

`launchurl` asks the current browser to launch a URL given as a string. Exactly what "launching a URL" means depends on the URL protocol, i.e. the part before the colon, and on the program which processes it. If the URL cannot be processed, the status is set to false; otherwise, it is true, which does not mean that a connection has been correctly established on the World Wide Web.

The current implementation uses the method `openURL:` of the AppKit framework on the Macintosh and `ShellExecute` on Windows. On Windows, the URL must begin with `http:`, `ftp:`, `gopher:`, `nntp:`, `news:`, `mailto:`, or `file:`. On Linux, the first application in the following list which is found in the current path is executed: `$X11BROWSER`, `$BROWSER` (environment variables), `htmlview`, `firefox`, `mozilla`, `netscape`, `opera`, `konqueror`; `launchurl` always returns true.

### Example

```
if ~launchurl('http://www.calerga.com')
  dialog('Cannot launch http://www.calerga.com');
end
```

## 6.11 Download URL

This section describes a function which downloads data from the WWW.

### urldownload

Launch a URL in the default browser.

### Syntax

```
contents = urldownload(url)
```

## Description

`urldownload(url)` downloads data referenced by a URL. The result is typically an HTML document, or a data file such as an image. Both input and output arguments are strings.

`urldownload(url,query)` submits a query with the GET method and downloads the result. The URL should use the HTTP or HTTPS protocol.

`urldownload(url,query,method)` query with the specified method ('get' or 'post') and downloads the result.

## Example

```
data = urldownload('http://www.w3.org');
```

## 6.12 Web Services

This section describes functions which implement the client side of the XML-RPC and SOAP protocols, as well as low-level functions which can also be used to implement the server side. XML-RPC and SOAP permit web services, i.e. calling a function on a remote server over the World Wide Web. XML-RPC is based on two standards: XML (eXtended Mark-up Language), used to encode the request to the server and its response to the client, and HTTP (HyperText Transfer Protocol), the main communication protocol used by the World Wide Web. In XML-RPC, RPC means Remote Procedure Call; it is a mechanism used for decades to communicate between a client and a server on a network. The advantages of XML-RPC are that it is based on the same technologies as the Web and it is very simple. Its drawbacks are that it is less efficient than a binary encoding, and it is sometimes too simple and requires encoding of binary data, which defeats its main advantage. For instance strings are encoded in ASCII, and supported types are much less rich than LME's.

SOAP is also a standard used for exchanging data encoded with XML. It is more complicated than XML-RPC and supports more types. Function parameters are referenced by name while XML-RPC uses an ordered list. SOAP requests can be sent with different communication protocols; the implementation described here uses only the most common one, HTTP.

### XML-RPC

In LME, XML-RPC makes calls to remote procedure similar to the use of `feval`. The two main functions are `xmlrpcall` and `xmlrpcallset`. Lower-level functions which encode and decode calls and responses,

while not necessary for standard calls, can be used to understand exactly how data are converted, to implement the server, or for special applications.

Procedure calls can contain parameters (arguments) and always return a single response. These data have different types. XML-RPC converts them automatically, as follows.

<b>XML-RPC</b>	<b>LME</b>
i4	int32 scalar
int	int32 scalar
boolean	logical scalar
string	character 1-by-n array
double	real double scalar
dateTime.iso8601	1-by-6 double array
base64	1-by-n uint8 array
struct	structure
array	list

There is no difference between i4 and int. In strings, only the least-significant byte is transmitted (i.e. only ASCII characters between 0 and 127 are transmitted correctly). Double values do not support an exponent (a sufficient number of zeros are used instead). The XML-RPC standard does not support inf and NaN; XML-RPC functions do, which should not do any harm. In LME, date and time are stored in a row vector which contains the year, month, day, hour, minute, and second (like the result of the function clock), without time zone information.

## SOAP

SOAP calls are very similar to XML-RPC. The main difference is that they use a single structure to represent the parameters. The member fields are used as parameter names. The table below shows the mapping between SOAP types and LME types.

<b>SOAP</b>	<b>LME</b>
xsd:int	int32 scalar
xsd:boolean	logical scalar
xsd:string	character 1-by-n array
xsd:double	real double scalar
xsd:dateTime	1-by-6, 1-by-7, or 1-by-8 double array
SOAP-ENC:base64 (structure)	1-by-n uint8 array structure
SOAP-ENC:array	list

In LME, time instants are stored as a row vector of 6, 7, or 8 elements which contains the year, month, day, hour, minute, second, time zone hour, and time zone minute; the time zone is optional. Ar-

rays which are declared with a single type `xsd:int`, `xsd:boolean`, or `xsd:double` are mapped to LME row vectors of the corresponding class.

The two main functions for performing a SOAP call are `soapcall` and `soapcallset`.

## soapcall

Perform a SOAP remote procedure call.

### Syntax

```
response = soapcall(url, method, ns, action, opt)
response = soapcall(url, method, ns, action, opt, param)
```

### Description

`soapcall(url, method, ns, action, opt, param)` calls a remote procedure using the SOAP protocol. `url` (a string) is either the complete URL beginning with `http://`, or only the absolute path; in the second case, the server address and port come from argument `opt`. `method` is the SOAP method name as a string; `ns` is its XML name space; `action` is the SOAP action. `opt` is a structure which contains the options; it is typically created with `soapcallset`, or can be the empty array `[]` for the default options. `param`, if present, is a structure which contains the parameters of the SOAP call.

### Example

The following call requests a translation from english to french (it assumes that the computer is connected to the Internet and that the service is available).

```
url = 'http://services.xmethods.net/perl/soaplite.cgi';
method = 'BabelFish';
ns = 'urn:xmethodsBabelFish';
action = 'urn:xmethodsBabelFish#BabelFish';
param = struct;
param.translationmode = 'en_fr';
param.sourcedata = 'Hello, Sysquake!';
fr = soapcall(url, method, ns, action, [], param)
fr =
    Bonjour, Sysquake!
```

Note that since the server address is given in the URL, the default options are sufficient. The variable `param` is reset to an empty structure to make sure that no other parameter remains from a previous call.

**See also**

soapcallset

**soapcallset**

Options for SOAP call.

**Syntax**

```

options = soapcallset
options = soapcallset(name1, value1, ...)
options = soapcallset(options0, name1, value1, ...)

```

**Description**

soapcallset(name1,value1,...) creates the option argument used by soapcall, including the server and port. 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, soapcallset creates a structure with all the default options. Note that soapcall also interpret the lack of an option argument, or the empty array [], as a request to use the default values.

When its first input argument is a structure, soapcallset adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
Server	''	server name or IP address
Port	80	port number
Timeout	10	maximum time in seconds
Debug	false	true to display data

If the server is an empty string, it is replaced with 'localhost'. The Debug field is not included in the default options; when set, it causes the display of the request and responses.

**Example**

Default options:

```

soapcallset
Server: ''
Port: 80
Timeout: 10

```

**See also**

soapcall

**soapreadcall**

Decode a SOAP call request.

**Syntax**

```
(method, namespace, pstruct, url) = soapreadcall(fd)
(method, namespace, pstruct, url) = soapreadcall(str)
```

**Description**

soapreadcall(fd), where fd is a file descriptor, reads a complete SOAP call, decodes it, and returns the result in four output arguments: the method name and namespace as strings, a structure which contains the parameters, and the URL as a string.

soapreadcall(str) decodes its string argument which must be a whole SOAP call.

**Example**

```
param = struct('x',pi,'y',true);
str = soapwritecall('/', '/', '', 'fun', 'namespace', param);
(method, ns, pstruct, url) = soapreadcall(str)
method =
    fun
ns =
    namespace
pstruct =
    x: 3.1416
    y: true
url =
    /
```

**See also**

soapreadresponse, soapwritecall

**soapreadresponse**

Decode a SOAP call response.

**Syntax**

```
(fault, value) = soapreadresponse(fd)
(fault, value) = soapreadresponse(str)
```

**Description**

soapreadresponse(fd), where fd is a file descriptor, reads a complete SOAP response and decodes it. In case of success, it returns true in the first output argument and the decoded response value in the second output argument. In case of failure, it returns false and the fault structure, which contains the fields faultcode (error code as a string) and faultstring (error message as a string).

soapreadresponse(str) decodes its string argument which must be a whole SOAP response.

**Examples**

```
str = soapwriteresponse('fun', 'namespace', 123);
(fault, value) = soapreadresponse(str)
  fault =
    false
  value =
    123
strf = soapwritefault(12int32, 'No power');
(fault, value) = soapreadresponse(strf)
  fault =
    true
  value =
    faultcode: '12'
    faultstring: 'No power'
```

**See also**

soapreadcall, soapwriteresponse, soapwritefault

**soapwritecall**

Encode a SOAP call request.

**Syntax**

```
soapwritecall(fd, server, url, action, method, ns, params)
soapwritecall(server, url, action, method, ns, params)
str = soapwritecall(server, url, action, method, ns, params)
```

## Description

`soapwritecall(fd, server, url, action, method, ns, params)` writes to file descriptor `fd` a complete SOAP call, including the HTTP header. If `fd` is missing, the call is written to standard output (file descriptor 1); since the output contains carriage return characters, it may not be displayed correctly on all platforms. The `server` argument is a string which contains the server name, and, optionally, a colon and the server port number. `url` is a string which contains the absolute path (without the protocol, server, and port part). `action` is a string which contains the SOAP action, or is empty if no action is required for the service. `method` contains the method name sent to the server; `ns` is its XML name space. `param`, if present, is a structure which contains the parameters of the SOAP call.

With an output argument, `soapwritecall` returns the call as a string, without any output.

## Example

```
param = struct('x',pi,'y',true);
soapwritecall('server.com','/', 'action', 'fun', 'ns', param)
POST / HTTP/1.1
User-Agent: LME 4.5
Host: server.com
Content-Type: text/xml; charset=utf-8
Content-Length: 495
SOAPAction: action

<?xml version="1.0"?>
<SOAP-ENV:Envelope
SOAP-ENV:encodingStyle="http://schemas.xmlsoap.org/soap/encoding/"
xmlns:SOAP-ENC="http://schemas.xmlsoap.org/soap/encoding"
xmlns:SOAP-ENV="http://schemas.xmlsoap.org/soap/envelope/"
xmlns:xsd="http://www.w3.org/1999/XMLSchema"
xmlns:xsi="http://www.w3.org/1999/XMLSchema-instance">
<SOAP-ENV:Body>
<m:fun xmlns:m="ns">
<x xsi:type="xsd:double">3.1415926535898</x>
<y xsi:type="xsd:boolean">1</y>
</m:fun>
</SOAP-ENV:Body>
</SOAP-ENV:Envelope>
```

## See also

`soapwriteresponse`, `soapreadcall`, `soapreadresponse`

## soapwritefault

Encode a SOAP call response fault.

### Syntax

```
soapwritefault(fd, faultCode, faultString)
soapwritefault(faultCode, faultString)
str = soapwritefault(faultCode, faultString)
```

### Description

soapwritefault(fd, faultCode, faultString) writes to file descriptor fd a complete SOAP response fault, including the HTTP header. If fd is missing, the response is written to standard output (file descriptor 1); since the output contains carriage return characters, it may not be displayed correctly on all platforms. The faultCode argument is the fault code as an integer or a string, and the faultString is the fault message.

With an output argument, soapwritefault returns the response as a string, without any output.

### See also

soapwriteresponse, soapreadresponse

## soapwriteresponse

Encode a SOAP call response.

### Syntax

```
soapwriteresponse(fd, method, ns, value)
soapwriteresponse(method, ns, value)
str = soapwriteresponse(method, ns, value)
```

### Description

soapwriteresponse(fd, method, ns, value) writes to file descriptor fd a complete SOAP response, including the HTTP header. If fd is missing, the response is written to standard output (file descriptor 1); since the output contains carriage return characters, it may not be displayed correctly on all platforms. The method argument is the method name as a string; ns is the XML name space; and value is the result of the call.

With an output argument, `soapwriteresponse` returns the response as a string, without any output.

### Example

```
soapwriteresponse('fun', 'namespace', 123)
HTTP/1.1 200 OK
Connection: close
Server: LME 4.5
Content-Length: 484
Content-Type: text/xml

<?xml version="1.0"?>
<SOAP-ENV:Envelope
SOAP-ENV:encodingStyle="http://schemas.xmlsoap.org/soap/encoding/"
xmlns:SOAP-ENC="http://schemas.xmlsoap.org/soap/encoding"
xmlns:SOAP-ENV="http://schemas.xmlsoap.org/soap/envelope/"
xmlns:xsd="http://www.w3.org/1999/XMLSchema"
xmlns:xsi="http://www.w3.org/1999/XMLSchema-instance">
<SOAP-ENV:Body>
<m:funResponse xmlns:m="namespace">
<Result xsi:type="xsd:double">123.</Result>
</m:funResponse>
</SOAP-ENV:Body>
</SOAP-ENV:Envelope>
```

### See also

`soapwritercall`, `soapreadresponse`, `soapreadcall`

## xmlrpcall

Perform an XML-RPC remote procedure call.

### Syntax

```
response = xmlrpcall(url, method, opt, params...)
```

### Description

`xmlrpcall(url, method, opt, params)` calls a remote procedure using the XML-RPC protocol. `url` (a string) is either the complete URL beginning with `http://`, or only the absolute path; in the second case, the server address and port come from argument `opt`. `method` is the XML-RPC method name as a string; `opt` is a structure which contains the options; it is typically created with `xmlrpcallset`, or can be the

empty array [] for the default options. The remaining input arguments are sent to the server as parameters of the XML-RPC call.

**Examples**

The following call requests the current time and date with a complete URL (it assumes that the computer is connected to the Internet and that the service is available).

```
url = 'http://time.xmlrpc.com/RPC2';
dateTime = xmlrpccall(url, 'currentTime.getCurrentTime')
dateTime =
    2005 1 20 17 32 47
```

The server address (and the server port if it was not the default value of 80) can also be specified in the options; then the URL contains only the absolute path.

```
server = xmlrpccallset('Server', 'time.xmlrpc.com');
dateTime = xmlrpccall('/RPC2', 'currentTime.getCurrentTime', server)
dateTime =
    2005 1 20 17 32 47
```

**See also**

xmlrpccallset

**xmlrpccallset**

Options for XML-RPC call.

**Syntax**

```
options = xmlrpccallset
options = xmlrpccallset(name1, value1, ...)
options = xmlrpccallset(options0, name1, value1, ...)
```

**Description**

xmlrpccallset(name1,value1,...) creates the option argument used by xmlrpccall, including the server and port. 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, xmlrpccallset creates a structure with all the default options. Note that xmlrpccall also interpret the lack of an option

argument, or the empty array [], as a request to use the default values.

When its first input argument is a structure, `xmlrpcallset` adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
Server	''	server name or IP address
Port	80	port number
Timeout	10	maximum time in seconds
Debug	false	true to display data

If the server is an empty string, it is replaced with 'localhost'. The Debug field is not included in the default options; when set, it causes the display of the request and responses.

### Example

Default options:

```
xmlrpcallset
  Server: ''
  Port: 80
  Timeout: 10
```

### See also

`xmlrpcall`

## xmlrpcreadcall

Decode an XML-RPC call request.

### Syntax

```
(method, arglist, url) = xmlrpcreadcall(fd)
(method, arglist, url) = xmlrpcreadcall(str)
```

### Description

`xmlrpcreadcall(fd)`, where `fd` is a file descriptor, reads a complete XML-RPC call, decodes it, and returns the result in three output arguments: the method name as a string, a list of arguments, and the URL as a string.

`xmlrpcreadcall(str)` decodes its string argument which must be a whole XML-RPC call.

**Example**

```

str = xmlrpcwritecall('rpc.remote.com', '/rpc', 'getPressure');
(method, arglist, url) = xmlrpcreadcall(str)
method =
    getPressure
arglist =
    {}
url =
    /rpc

```

**See also**

xmlrpcreadresponse, xmlrpcwritecall

**xmlrpcreadresponse**

Decode an XML-RPC call response.

**Syntax**

```

(fault, value) = xmlrpcreadresponse(fd)
(fault, value) = xmlrpcreadresponse(str)

```

**Description**

xmlrpcreadresponse(fd), where fd is a file descriptor, reads a complete XML-RPC response and decodes it. In case of success, it returns true in the first output argument and the decoded response value in the second output argument. In case of failure, it returns false and the fault structure, which contains the fields faultCode (error code as an int32) and faultString (error message as a string).

xmlrpcreadresponse(str) decodes its string argument which must be a whole XML-RPC response.

**Examples**

```

str = xmlrpcwriteresponse(123);
(fault, value) = xmlrpcreadresponse(str)
fault =
    false
value =
    123
strf = xmlrpcwritefault(12int32, 'No power');
(fault, value) = xmlrpcreadresponse(strf)
fault =
    true

```

```
value =
  faultCode: 12int32
  faultString: 'No power'
```

## See also

`xmlrpcreadcall`, `xmlrpcwriteresponse`, `xmlrpcwritefault`

## xmlrpcwritecall

Encode an XML-RPC call request.

### Syntax

```
xmlrpcwritecall(fd, server, url, method, params...)
xmlrpcwritecall(server, url, method, params...)
str = xmlrpcwritecall(server, url, method, params...)
```

### Description

`xmlrpcwritecall(fd, server, url, method, params...)` writes to file descriptor `fd` a complete XML-RPC call, including the HTTP header. If `fd` is missing, the call is written to standard output (file descriptor 1); since the output contains carriage return characters, it may not be displayed correctly on all platforms. The `server` argument is a string which contains the server name, and, optionally, a colon and the server port number. The `url` argument is a string which contains the absolute path (without the protocol, server, and port part). The `method` argument contains the method name sent to the server. Remaining input arguments, if any, are sent as parameters.

With an output argument, `xmlrpcwritecall` returns the call as a string, without any output.

### Example

```
xmlrpcwritecall('rpc.remote.com', '/rpc', 'getPressure', 1int32)
POST /rpc HTTP/1.0
User-Agent: LME 4.5
Host: rpc.remote.com
Content-Type: text/xml
Content-Length: 111

<?xml version="1.0"?>
<methodCall>
<methodName>getPressure</methodName>
```

```
<params>
<param>
<value>
<int>1</int>
</value>
</param>
</params>
</methodCall>
```

### See also

`xmlrpcwriteresponse`, `xmlrpcreadcall`, `xmlrpcreadresponse`

## xmlrpcwritedata

Encode an XML-RPC value.

### Syntax

```
xmlrpcwritedata(fd, val)
xmlrpcwritedata(val)
str = xmlrpcwritedata(val)
```

### Description

`xmlrpcwritedata(fd, val)` writes to file descriptor `fd` the value `val` encoded for XML-RPC. If `fd` is missing, the value is written to standard output (file descriptor 1); since the output contains carriage return characters, it may not be displayed correctly on all platforms.

With an output argument, `xmlrpcwritedata` returns the encoded value as a string, without any output.

### Example

```
xmlrpcwritedata(pi)
<double>3.141592653589</double>
```

### See also

`xmlrpcwritecall`, `xmlrpcwriteresponse`

## xmlrpcwritefault

Encode an XML-RPC call response fault.

## Syntax

```
xmlrpcwritefault(fd, faultCode, faultString)
xmlrpcwritefault(faultCode, faultString)
str = xmlrpcwritefault(faultCode, faultString)
```

## Description

`xmlrpcwritefault(fd, faultCode, faultString)` writes to file descriptor `fd` a complete XML-RPC response fault, including the HTTP header. If `fd` is missing, the response is written to standard output (file descriptor 1); since the output contains carriage return characters, it may not be displayed correctly on all platforms. The `faultCode` argument is the numeric fault code, and the `faultString` is the fault message.

With an output argument, `xmlrpcwritefault` returns the response fault as a string, without any output.

## See also

`xmlrpcwriteresponse`, `xmlrpcreadresponse`

## xmlrpcwriteresponse

Encode an XML-RPC call response.

## Syntax

```
xmlrpcwriteresponse(fd, value)
xmlrpcwriteresponse(value)
str = xmlrpcwriteresponse(value)
```

## Description

`xmlrpcwriteresponse(fd, value)` writes to file descriptor `fd` a complete XML-RPC response, including the HTTP header. If `fd` is missing, the response is written to standard output (file descriptor 1); since the output contains carriage return characters, it may not be displayed correctly on all platforms. The `value` argument is the result of the call.

With an output argument, `xmlrpcwriteresponse` returns the response as a string, without any output.

**Example**

```
xmlrpcwriteresponse(123)
HTTP/1.1 200 OK
Connection: close
Server: LME 4.5
Content-Length: 123
Content-Type: text/xml

<?xml version="1.0"?>
<methodResponse>
<params>
<param>
<double>123.</double>
</param>
</params>
</methodResponse>
```

**See also**

xmlrpcwritecall, xmlrpcreadresponse, xmlrpcreadcall

## 6.13 Serial port

Serial port functions enable communication with devices connected to the computer via an RS-232 interface. Such devices include modems, printers, and many scientific instruments. The operating system can also emulate RS-232 connections with other devices, such as built-in modems or USB (Universal Serial Bus) devices.

Functions described in this section include only those required for opening and configuring the connection. They correspond to fopen for files. Input, output, and control are done with the following generic functions:

<b>Function</b>	<b>Description</b>
fclose	close the file
fflush	flush I/O buffers
fgetl	read a line
fgets	read a line
fprintf	write formatted data
fread	read data
fscanf	read formatted data
fwrite	write data
redirect	redirect output

Functions opendirdevice, devicename, closedevice, and flushdevice are obsolete and may be removed in the future. They

are replaced with `serialdevopen` and `serialdevset` to specify configuration settings, `serialdevname`, `fclose`, and `fflush`.

## **serialdevname**

Serial device name.

### **Syntax**

```
name = serialdevname(n)
list = serialdevname
```

### **Description**

`serialdevname(n)` returns the name of the `n`:th serial device which can be opened by `serialdevopen`. Argument `n` must be 1 or higher; with other values, such as those larger than the number of serial devices available on your computer, `serialdevname` returns the empty string.

Without input argument, `serialdevname` gives the list of serial device names.

### **Examples**

On a Macintosh with internal modem:

```
serialdevname(1)
Internal Modem
```

Under Windows:

```
serialdevname(1)
COM1
```

### **See also**

`serialdevopen`

## **serialdevopen**

Open a serial port.

### **Syntax**

```
fd = serialdevopen(portname, options)
fd = serialdevopen(portname)
```

**Description**

`serialdevopen(portname)` opens a connection to the serial port whose name is `portname` and returns a file descriptor `fd`. Names depend on the operating system and can be obtained with `serialdevname`.

Some platforms do not provide a complete list of all ports; `serialdevopen` may accept additional device names and pass them directly to the corresponding function of the operating system.

The second argument of `serialdevopen(portname,options)` is a structure which contains configuration settings. It is set with `serialdevset`.

Once a connection has been opened, the file descriptor `fd` can be used with functions such as `fread`, `fwrite`, `fscanf`, and `fprintf`. The connection is closed with `fclose`.

**Example**

```
fd = serialdevopen(serialdevname(1), ...
    serialdevset('BPS',19200,'TextMode',true,'Timeout',2));
fprintf(fd, 'L,%d,2\n', 1);
reply = fgetl(fd)
fclose(fd);
```

**See also**

`fclose`, `serialdevname`, `serialdevset`, `fflush`, `fread`, `fwrite`, `fscanf`, `fgetl`, `fgets`, `fprintf`

**serialdevset**

Configuration settings for serial port.

**Syntax**

```
options = serialdevset
options = serialdevset(name1, value1, ...)
options = serialdevset(options0, name1, value1, ...)
```

**Description**

`serialdevset(name1,value1,...)` creates the option argument used by `serialdevopen`. 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, `serialdevset` creates a structure with all the default settings. Note that `serialdevopen` also interprets the lack of an option argument, or the empty array [], as a request to use the default values.

When its first input argument is a structure, `serialdevset` adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
BPS	19200	bit per seconds
Delay	0	delay after character output in seconds
Handshake	false	hardware handshake
StopBits	2	number of stop bits (1, 1.5, or 2)
TextMode	false	text mode
Timeout	1	timeout in seconds

Output operations wait for the specified delay after each character; this can be useful with slow devices without handshake.

When text mode is set, input CR and CR/LF sequences are converted to LF. Output CR and LF are not converted.

Depending on the platform, operations which use the timeout value (such as input) can be interrupted with the platform-dependent abort key(s) (typically Escape or Control-C) or are limited to 10 seconds.

## Example

```
serialdevset
  BPS: 19200
  Handshake: false
  StopBits: 2
  TextMode: false
  Timeout: 1
```

## See also

`serialdevopen`, `serialdevname`

## 6.14 Audio output

This section describes functions which play sounds.

### **audioplay**

Play audio samples.

## Syntax

```
audioplay(samples)
audioplay(samples, options)
```

## Description

`audioplay(samples)` plays the audio samples in array `samples` at a sample rate of 44.1 kHz. Each column of `samples` is a channel (i.e. `samples` is a column vector for monophonic sound and a two-column array for stereophonic sound), and each row is a sample. Samples are stored as double or single numbers between -1 and 1, `int8` numbers between -128 and 127, or `int16` numbers between -32768 and 32767.

`audioplay(samples,options)` uses the specified options, which are typically built with `audioset`.

## Examples

A monophonic bell-like sound of two seconds with a frequency of 740 Hz and a damping time constant of 0.5 second:

```
t = (0:88200)'/44100;
samples = sin(2*pi*740*t).*exp(-t/0.5);
audioplay(samples);
```

Some white noise which oscillates 5 times between left and right:

```
t = (0:44099)' / 44100;
noise = 0.1 * randn(length(t), 1);
left = cos(2 * pi * t) .* noise;
right = sin(2 * pi * t) .* noise;
opt = audioset('Repeat', 5);
audioplay([left, right], opt);
```

## See also

`audioset`

## audioset

Options for audio.

## Syntax

```
options = audioset
options = audioset(name1, value1, ...)
options = audioset(options0, name1, value1, ...)
```

## Description

`audioset(name1,value1,...)` creates the option argument used by `audioplay`. 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, `audioset` creates a structure with all the default options. Note that `audioplay` also interprets the lack of an option argument, or the empty array `[]`, as a request to use the default values.

When its first input argument is a structure, `audioset` adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
Repeat	1	number of repetitions
SampleRate	44100	sample rate in Hz

Default values may be different on platforms with limited audio capabilities.

## Example

Default options:

```
audioset
  Repeat: 1
  SampleRate: 44100
```

## See also

`audioplay`

## 6.15 Audio Input

This section describes functions which record sounds. Currently, these functions are available on Windows and Mac OS X.

### **audiorecord**

Record audio samples.

## Syntax

```
samples = audiorecord(t)
samples = audiorecord(t, options)
```

## Description

`audiorecord(t)` records audio samples for `t` seconds and returns them in an array of single numbers. On computers which support them, the default sample rate is 44.1 kHz and the default number of channels is 2 (stereo). The result has 2 columns (one per channel) and as many rows as samples.

`audiorecord(t,options)` uses the specified options, which are typically built with `audiorecordset`.

## Example

```
samples = audiorecord(1);
audioplay(samples);
```

## See also

`audiorecordset`, `audioplay`

## audiorecordset

Options for audio.

## Syntax

```
options = audiorecordset
options = audiorecordset(name1, value1, ...)
options = audiorecordset(options0, name1, value1, ...)
```

## Description

`audiorecordset(name1,value1,...)` creates the option argument used by `audiorecord`. 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, `audiorecordset` creates a structure with all the default options. Note that `audiorecord` also interprets the lack of an option argument, or the empty array `[]`, as a request to use the default values.

When its first input argument is a structure, `audiorecordset` adds or changes fields which correspond to the name/value pairs which follow.

Here is the list of permissible options:

<b>Name</b>	<b>Default</b>	<b>Meaning</b>
<code>SampleRate</code>	44100	sample rate in Hz
<code>Stereo</code>	true	true for stereo, false for mono

Default values may be different on platforms with limited audio capabilities.

## Examples

Default options:

```
audiorecordset
  SampleRate: 44100
  Stereo: true
```

Record in mono:

```
samples = audiorecord(1, audiorecordset('Stereo', false));
```

## See also

`audiorecord`

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