

# Package ‘cmna’

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**Description** Provides the source and examples for James P. Howard, II,  
``Computational Methods for Numerical Analysis with R,"  
<<https://jameshoward.us/cmna>>, a forthcoming book on  
numerical methods in R.

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**Author** James P. Howard, II [aut, cre]

**Maintainer** ``James P. Howard, II" <jh@jameshoward.us>

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**Description**

Provides the source and examples for *Computational Methods for Numerical Analysis with R*.

**Details**

This package provides a suite of simple implementations of standard methods from numerical analysis. The collection is designed to accompany *Computational Methods for Numerical Analysis with R* by James P. Howard, II. Together, these functions provide methods to support linear algebra, interpolation, integration, root finding, optimization, and differential equations.

**Author(s)**

James P. Howard, II <jh@jameshoward.us>

**See Also**

Useful links:

- <https://jameshoward.us/cmna/>
- <https://github.com/howardjp/cmna>
- Report bugs at <https://github.com/howardjp/cmna/issues>

**Description**

Adaptive integration

**Usage**

```
adaptint(f, a, b, n = 10, tol = 1e-06)
```

**Arguments**

f	function to integrate
a	the a-bound of integration
b	the b-bound of integration
n	the maximum recursive depth
tol	the maximum error tolerance

**Details**

The `adaptint` function uses Romberg's rule to calculate the integral of the function `f` over the interval from `a` to `b`. The parameter `n` sets the number of intervals to use when evaluating. Additional options are passed to the function `f` when evaluating.

**Value**

the value of the integral

**See Also**

Other integration: [gaussint](#), [giniquintile](#), [mcint](#), [midpt](#), [revolution-solid](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

Other newton-cotes: [giniquintile](#), [midpt](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

**Examples**

```
f <- function(x) { sin(x)^2 + log(x) }
adaptint(f, 1, 10, n = 4)
adaptint(f, 1, 10, n = 5)
adaptint(f, 1, 10, n = 10)
```

---

bezier

*Bezier curves*


---

**Description**

Find the quadratic and cubic Bezier curve for the given points

**Usage**

```
qbezier(x, y, t)
```

```
cbezier(x, y, t)
```

**Arguments**

<code>x</code>	a vector of x values
<code>y</code>	a vector of y values
<code>t</code>	a vector of t values for which the curve will be computed

**Details**

`qbezier` finds the quadratic Bezier curve for the given three points and `cbezier` finds the cubic Bezier curve for the given four points. The curve will be computed at all values in the vector `t` and a list of x and y values returned.

**Value**

a list composed of an x-vector and a y-vector

**See Also**

Other interp: [bilinear](#), [cubicspline](#), [linterp](#), [nn](#), [polyinterp](#), [pwiselinterp](#)

**Examples**

```
x <- c(1, 2, 3)
y <- c(2, 3, 5)
f <- qbezier(x, y, seq(0, 1, 1/100))
```

```
x <- c(-1, 1, 0, -2)
y <- c(-2, 2, -1, -1)
f <- cbezier(x, y, seq(0, 1, 1/100))
```

---

bilinear

*Bilinear interpolation*

---

**Description**

Finds a bilinear interpolation bounded by four points

**Usage**

```
bilinear(x, y, z, newx, newy)
```

**Arguments**

x	vector of two x values representing x_1 and x_2
y	vector of two y values representing y_1 and y_2
z	2x2 matrix if z values
newx	vector of new x values to interpolate
newy	vector of new y values to interpolate

**Details**

bilinear finds a bilinear interpolation bounded by four corners

**Value**

a vector of interpolated z values at (x, y)

**See Also**

Other interp: [bezier](#), [cubicspline](#), [linterp](#), [nn](#), [polyinterp](#), [pwiselinterp](#)

Other algebra: [cubicspline](#), [division](#), [fibonacci](#), [horner](#), [isPrime](#), [linterp](#), [nthroot](#), [polyinterp](#), [pwiselinterp](#), [quadratic](#)

**Examples**

```
x <- c(2, 4)
y <- c(4, 7)
z <- matrix(c(81, 84, 85, 89), nrow = 2)
newx <- c(2.5, 3, 3.5)
newy <- c(5, 5.5, 6)
bilinear(x, y, z, newx, newy)
```

---

bisection

*The Bisection Method*

---

**Description**

Use the bisection method to find real roots

**Usage**

```
bisection(f, a, b, tol = 0.001, m = 100)
```

**Arguments**

f	function to locate a root for
a	the a bound of the search region
b	the b bound of the search region
tol	the error tolerance
m	the maximum number of iterations

**Details**

The bisection method functions by repeatedly halving the interval between a and b and will return when the interval between them is less than tol, the error tolerance. However, this implementation also stops if after m iterations.

**Value**

the real root found

**See Also**

Other optimz: [goldsect](#), [gradient](#), [hillclimbing](#), [newton](#), [sa](#), [secant](#)

**Examples**

```
f <- function(x) { x^3 - 2 * x^2 - 159 * x - 540 }  
bisection(f, 0, 10)
```

---

**bvp***Boundary value problems*

---

**Description**

solve boundary value problems for ordinary differential equations

**Usage**

```
bvpexample(x)  
  
bvpexample10(x)
```

**Arguments**

x                    proposed initial x-value

**Details**

The euler method implements the Euler method for solving differential equations. The codemidp-  
tvp method solves initial value problems using the second-order Runge-Kutta method. The rungekutta4  
method is the fourth-order Runge-Kutta method.

**Value**

a data frame of x and y values

**Examples**

```
bvpexample(-2)  
bvpexample(-1)  
bvpexample(0)  
bvpexample(1)  
bvpexample(2)  
## (bvp.b <- bisection(bvpexample, 0, 1))  
## (bvp.s <- secant(bvpexample, 0))
```

choleskymatrix      *Cholesky Decomposition*

---

**Description**

Decompose a matrix into the Cholesky

**Usage**

```
choleskymatrix(m)
```

**Arguments**

m                    a matrix

**Details**

choleskymatrix decomposes the matrix m into the LU decomposition, such that  $m == L$

**Value**

the matrix L

**See Also**

Other linear: [detmatrix](#), [gdls](#), [invmatrix](#), [iterativematrix](#), [lumatrix](#), [refmatrix](#), [rowops](#), [tridiagmatrix](#), [vecnorm](#)

**Examples**

```
(A <- matrix(c(5, 1, 2, 1, 9, 3, 2, 3, 7), 3))
(L <- choleskymatrix(A))
t(L) %**% L
```

---

cubicspline      *Natural cubic spline interpolation*

---

**Description**

Finds a piecewise linear function that interpolates the data points

**Usage**

```
cubicspline(x, y)
```



**Arguments**

x                    a vector of x values  
y                    a vector of y values

**Details**

cubicspline finds a piecewise cubic spline function that interpolates the data points. For each x-y ordered pair. The function will return a list of four vectors representing the coefficients.

**Value**

a list of coefficient vectors

**See Also**

Other interp: [bezier](#), [bilinear](#), [linterp](#), [nn](#), [polyinterp](#), [pwiselinterp](#)

Other algebra: [bilinear](#), [division](#), [fibonacci](#), [horner](#), [isPrime](#), [linterp](#), [nthroot](#), [polyinterp](#), [pwiselinterp](#), [quadratic](#)

**Examples**

```
x <- c(1, 2, 3)
y <- c(2, 3, 5)
f <- cubicspline(x, y)
```

```
x <- c(-1, 1, 0, -2)
y <- c(-2, 2, -1, -1)
f <- cubicspline(x, y)
```

---

detmatrix

*Calculate the determinant of the matrix*

---

**Description**

Calculate the determinant of the matrix

**Usage**

```
detmatrix(m)
```

**Arguments**

m                    a matrix

**Details**

detmatrix calculates the determinant of the matrix given.

**Value**

the determinant

**See Also**

Other linear: [choleskymatrix](#), [gdls](#), [invmatrix](#), [iterativematrix](#), [lumatrix](#), [refmatrix](#), [rowops](#), [tridiagmatrix](#), [vecnorm](#)

**Examples**

```
A <- matrix(c(1, 2, -7, -1, -1, 1, 2, 1, 5), 3)
detmatrix(A)
```

---

division

*Algorithms for divisions*

---

**Description**

Algorithms for division that provide a quotient and remainder.

**Usage**

```
naivediv(m, n)
```

```
longdiv(m, n)
```

**Arguments**

m            the dividend

n            the divisor

**Details**

The `naivediv` divides `m` by `n` by using repeated division. The `longdiv` function uses the long division algorithm in binary.

**Value**

the quotient and remainder as a list

**See Also**

Other algebra: [bilinear](#), [cubicspline](#), [fibonacci](#), [horner](#), [isPrime](#), [linterp](#), [nthroot](#), [polyinterp](#), [pwiselinterp](#), [quadratic](#)

**Examples**

```
a <- floor(runif(1, 1, 1000))
b <- floor(runif(1, 1, 100))
naivediv(a, b)
longdiv(a, b)
```

---

fibonacci

*Fibonacci numbers*

---

**Description**

Return the n-th Fibonacci number

**Usage**

```
fibonacci(n)
```

**Arguments**

n                    n

**Details**

This function recursively implements the famous Fibonacci sequence. The function returns the nth member of the sequence.

**Value**

the sequence element

**See Also**

Other algebra: [bilinear](#), [cubicspline](#), [division](#), [horner](#), [isPrime](#), [linterp](#), [nthroot](#), [polyinterp](#), [pwiselinterp](#), [quadratic](#)

**Examples**

```
fibonacci(10)
```

---

`findiff`*Finite Differences*

---

**Description**

Finite differences formulas

**Usage**

```
findiff(f, x, h = x * sqrt(.Machine$double.eps))
```

```
syndiff(f, x, h = x * .Machine$double.eps^(1/3))
```

```
findiff2(f, x, h)
```

```
rdiff(f, x, n = 10, h = 1e-04)
```

**Arguments**

<code>f</code>	function to differentiate
<code>x</code>	the x-value to differentiate at
<code>h</code>	the step-size for evaluation
<code>n</code>	the maximum number of convergence steps in <code>rdiff</code>

**Details**

The `findiff` formula uses the finite differences formula to find the derivative of `f` at `x`. The value of `h` is the step size of the evaluation. The function `findiff2` provides the second derivative.

**Value**

the value of the derivative

**Examples**

```
findiff(sin, pi, 1e-3)
syndiff(sin, pi, 1e-3)
```

---

`gaussint`*Gaussian integration method driver*

---

**Description**

Use the Gaussian method to evaluate integrals

**Usage**

```
gaussint(f, x, w)
```

```
gauss.legendre(f, m = 5)
```

```
gauss.laguerre(f, m = 5)
```

```
gauss.hermite(f, m = 5)
```

**Arguments**

<code>f</code>	function to integrate
<code>x</code>	list of evaluation points
<code>w</code>	list of weights
<code>m</code>	number of evaluation points

**Details**

The `gaussint` function uses the Gaussian integration to evaluate an integral. The function itself is a driver and expects the integration points and associated weights as options.

**Value**

the value of the integral

**See Also**

Other integration: [adaptint](#), [giniquintile](#), [mcint](#), [midpt](#), [revolution-solid](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

**Examples**

```
w = c(1, 1)
x = c(-1 / sqrt(3), 1 / sqrt(3))
f <- function(x) { x^3 + x + 1 }
gaussint(f, x, w)
```

---

`gdls`*Least squares with gradient descent*

---

**Description**

Solve least squares with gradient descent

**Usage**

```
gdls(A, b, alpha = 0.05, tol = 1e-06, m = 1e+05)
```

**Arguments**

<code>A</code>	a square matrix representing the coefficients of a linear system
<code>b</code>	a vector representing the right-hand side of the linear system
<code>alpha</code>	the learning rate
<code>tol</code>	the expected error tolerance
<code>m</code>	the maximum number of iterations

**Details**

`gdls` solves a linear system using gradient descent.

**Value**

the modified matrix

**See Also**

Other linear: [choleskymatrix](#), [detmatrix](#), [invmatrix](#), [iterativematrix](#), [lumatrix](#), [refmatrix](#), [rowops](#), [tridiagmatrix](#), [vecnorm](#)

**Examples**

```
head(b <- iris$Sepal.Length)
head(A <- matrix(cbind(1, iris$Sepal.Width, iris$Petal.Length, iris$Petal.Width), ncol = 4))
gdls(A, b, alpha = 0.05, m = 10000)
```

---

giniquintile	<i>Gini coefficients</i>
--------------	--------------------------

---

**Description**

Calculate the Gini coefficient from quintile data

**Usage**

```
giniquintile(L)
```

**Arguments**

L                    vector of percentages at 20th, 40th, 60th, and 80th percentiles

**Details**

Calculate the Gini coefficient given the quintile data.

**Value**

the estimated Gini coefficient

**References**

Leon Gerber, "A Quintile Rule for the Gini Coefficient", *Mathematics Magazine*, 80:2, April 2007.

**See Also**

Other integration: [adaptint](#), [gaussint](#), [mcint](#), [midpt](#), [revolution-solid](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

Other newton-cotes: [adaptint](#), [midpt](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

**Examples**

```
L <- c(4.3, 9.8, 15.4, 22.7)
giniquintile(L)
```

---

`goldsect`*Golden Section Search*

---

**Description**

Use golden section search to find local extrema

**Usage**

```
goldsectmin(f, a, b, tol = 0.001, m = 100)
```

```
goldsectmax(f, a, b, tol = 0.001, m = 100)
```

**Arguments**

<code>f</code>	function to integrate
<code>a</code>	the a bound of the search region
<code>b</code>	the b bound of the search region
<code>tol</code>	the error tolerance
<code>m</code>	the maximum number of iterations

**Details**

The golden section search method functions by repeatedly dividing the interval between a and b and will return when the interval between them is less than `tol`, the error tolerance. However, this implementation also stop if after `m` iterations.

**Value**

the x value of the minimum found

**See Also**

Other optimz: [bisection](#), [gradient](#), [hillclimbing](#), [newton](#), [sa](#), [secant](#)

**Examples**

```
f <- function(x) { x^2 - 3 * x + 3 }  
goldsectmin(f, 0, 5)
```



---

gradient	<i>Gradient descent</i>
----------	-------------------------

---

### Description

Use gradient descent to find local minima

### Usage

```
graddsc(fp, x, h = 0.001, tol = 1e-04, m = 1000)
```

```
gradasc(fp, x, h = 0.001, tol = 1e-04, m = 1000)
```

```
gd(fp, x, h = 100, tol = 1e-04, m = 1000)
```

### Arguments

fp	function representing the derivative of f
x	an initial estimate of the minima
h	the step size
tol	the error tolerance
m	the maximum number of iterations

### Details

Gradient descent can be used to find local minima of functions. It will return an approximation based on the step size  $h$  and  $fp$ . The  $tol$  is the error tolerance,  $x$  is the initial guess at the minimum. This implementation also stops after  $m$  iterations.

### Value

the  $x$  value of the minimum found

### See Also

Other optimz: [bisection](#), [goldsect](#), [hillclimbing](#), [newton](#), [sa](#), [secant](#)

### Examples

```
fp <- function(x) { x^3 + 3 * x^2 - 1 }  
graddsc(fp, 0)
```

```
f <- function(x) { (x[1] - 1)^2 + (x[2] - 1)^2 }  
fp <-function(x) {  
  x1 <- 2 * x[1] - 2  
  x2 <- 8 * x[2] - 8
```

```
    return(c(x1, x2))
  }
gd(fp, c(0, 0), 0.05)
```

---

heat

*Heat Equation via Forward-Time Central-Space*

---

### Description

solve heat equation via forward-time central-space method

### Usage

```
heat(u, alpha, xdelta, tdelta, n)
```

### Arguments

u	the initial values of u
alpha	the thermal diffusivity coefficient
xdelta	the change in x at each step in u
tdelta	the time step
n	the number of steps to take

### Details

The heat solves the heat equation using the forward-time central-space method in one-dimension.

### Value

a matrix of u values at each time step

### Examples

```
alpha <- 1
x0 <- 0
xdelta <- .05
x <- seq(x0, 1, xdelta)
u <- sin(x^4 * pi)
tdelta <- .001
n <- 25
z <- heat(u, alpha, xdelta, tdelta, n)
```

---

hillclimbing	<i>Hill climbing</i>
--------------	----------------------

---

**Description**

Use hill climbing to find the global minimum

**Usage**

```
hillclimbing(f, x, h = 1, m = 1000)
```

**Arguments**

f	function representing the derivative of f
x	an initial estimate of the minimum
h	the step size
m	the maximum number of iterations

**Details**

Hill climbing

**Value**

the x value of the minimum found

**See Also**

Other optimz: [bisection](#), [goldsect](#), [gradient](#), [newton](#), [sa](#), [secant](#)

**Examples**

```
f <- function(x) {  
  (x[1]^2 + x[2] - 11)^2 + (x[1] + x[2]^2 - 7)^2  
}  
hillclimbing(f, c(0,0))  
hillclimbing(f, c(-1,-1))  
hillclimbing(f, c(10,10))
```

---

`himmelblau`*Himmelblau Function*

---

**Description**

Generate the Himmelblau function

**Usage**`himmelblau(x)`**Arguments**

`x` a vector of x-values

**Details**

Generate the Himmelblau function

**Value**

the value of the function at `x`.

---

`horner`*Horner's rule*

---

**Description**

Use Horner's rule to evaluate a polynomial

**Usage**`horner(x, coefs)``rhorer(x, coefs)``naivepoly(x, coefs)``betterpoly(x, coefs)`**Arguments**

`x` a vector of x values to evaluate the polynomial

`coefs` vector of coefficients of x

### Details

This function implements Horner's rule for fast polynomial evaluation. The implementation expects  $x$  to be a vector of  $x$  values at which to evaluate the polynomial. The parameter `coefs` is a vector of coefficients of  $x$ . The vector order is such that the first element is the constant term, the second element is the coefficient of  $x$ , the so forth to the highest degree term. Terms with a 0 coefficient should have a 0 element in the vector.

The function `rhorners` implements the the Horner algorithm recursively.

The function `naivepoly` implements a polynomial evaluator using the straightforward algebraic approach.

The function `betterpoly` implements a polynomial evaluator using the straightforward algebraic approach with cached  $x$  terms.

### Value

the value of the function at  $x$

### See Also

Other algebra: [bilinear](#), [cubicspline](#), [division](#), [fibonacci](#), [isPrime](#), [linterp](#), [nthroot](#), [polyinterp](#), [pwiselinterp](#), [quadratic](#)

### Examples

```
b <- c(2, 10, 11)
x <- 5
horner(x, b)
b <- c(-1, 0, 1)
x <- c(1, 2, 3, 4)
horner(x, b)
rhorners(x, b)
```

---

invmatrix

*Invert a matrix*

---

### Description

Invert the matrix using Gaussian elimination

### Usage

```
invmatrix(m)
```

### Arguments

`m` a matrix

**Details**

`invmatrix` invertse the given matrix using Gaussian elimination and returns the result.

**Value**

the inverted matrix

**See Also**

Other linear: [choleskymatrix](#), [detmatrix](#), [gdls](#), [iterativematrix](#), [lumatrix](#), [refmatrix](#), [rowops](#), [tridiagmatrix](#), [vecnorm](#)

**Examples**

```
A <- matrix(c(1, 2, -7, -1, -1, 1, 2, 1, 5), 3)
refmatrix(A)
```

---

isPrime

*Test for Primality*

---

**Description**

Test the number given for primality.

**Usage**

```
isPrime(n)
```

**Arguments**

n                    n

**Details**

This function tests `n` if it is prime through repeated division attempts. If a match is found, by finding a remainder of 0, FALSE is returned.

**Value**

boolean TRUE if `n` is prime, FALSE if not

**See Also**

Other algebra: [bilinear](#), [cubicspline](#), [division](#), [fibonacci](#), [horner](#), [linterp](#), [nthroot](#), [polyinterp](#), [pwiselinterp](#), [quadratic](#)

**Examples**

```
isPrime(37)
isPrime(89)
isPrime(100)
```

---

iterativematrix      *Solve a matrix using iterative methods*

---

**Description**

Solve a matrix using iterative methods.

**Usage**

```
jacobi(A, b, tol = 1e-06, maxiter = 100)
gaussseidel(A, b, tol = 1e-06, maxiter = 100)
cgmmatrix(A, b, tol = 1e-06, maxiter = 100)
```

**Arguments**

A	a square matrix representing the coefficients of a linear system
b	a vector representing the right-hand side of the linear system
tol	is a number representing the error tolerance
maxiter	is the maximum number of iterations

**Details**

`jacobi` finds the solution using Jacobi iteration. Jacobi iteration depends on the matrix being diagonally-dominant. The tolerance is specified the norm of the solution vector.

`gaussseidel` finds the solution using Gauss-Seidel iteration. Gauss-Seidel iteration depends on the matrix being either diagonally-dominant or symmetric and positive definite.

`cgmmatrix` finds the solution using the conjugate gradient method. The conjugate gradient method depends on the matrix being symmetric and positive definite.

**Value**

the solution vector

**See Also**

Other linear: [choleskymatrix](#), [detmatrix](#), [gdls](#), [invmatrix](#), [lumatrix](#), [refmatrix](#), [rowops](#), [tridiagmatrix](#), [vecnorm](#)

**Examples**

```
A <- matrix(c(5, 2, 1, 2, 7, 3, 3, 4, 8), 3)
b <- c(40, 39, 55)
jacobi(A, b)
```

---

ivp

*Initial value problems*

---

**Description**

solve initial value problems for ordinary differential equations

**Usage**

```
euler(f, x0, y0, h, n)
midptivp(f, x0, y0, h, n)
rungekutta4(f, x0, y0, h, n)
adamsbashforth(f, x0, y0, h, n)
```

**Arguments**

f	function to integrate
x0	the initial value of x
y0	the initial value of y
h	selected step size
n	the number of steps

**Details**

The `euler` method implements the Euler method for solving differential equations. The `codemidptivp` method solves initial value problems using the second-order Runge-Kutta method. The `rungekutta4` method is the fourth-order Runge-Kutta method.

**Value**

a data frame of x and y values

**Examples**

```
f <- function(x, y) { y / (2 * x + 1) }
ivp.euler <- euler(f, 0, 1, 1/100, 100)
ivp.midpt <- midptivp(f, 0, 1, 1/100, 100)
ivp.rk4 <- rungekutta4(f, 0, 1, 1/100, 100)
```



---

`ivpsys`*Initial value problems for systems of ordinary differential equations*

---

**Description**

solve initial value problems for systems ordinary differential equations

**Usage**

```
eulersys(f, x0, y0, h, n)
```

**Arguments**

<code>f</code>	function to integrate
<code>x0</code>	the initial value of x
<code>y0</code>	the vector initial values of y
<code>h</code>	selected step size
<code>n</code>	the number of steps

**Details**

The euler method implements the Euler method for solving differential equations. The `codemidp-tivp` method solves initial value problems using the second-order Runge-Kutta method. The `rungekutta4` method is the fourth-order Runge-Kutta method.

**Value**

a data frame of x and y values

**Examples**

```
f <- function(x, y) { y / (2 * x + 1) }  
ivp.euler <- euler(f, 0, 1, 1/100, 100)
```

---

`linterp`*Linear interpolation*

---

**Description**

Finds a linear function between two points

**Usage**

```
linterp(x1, y1, x2, y2)
```

**Arguments**

x1	x value of the first point
y1	y value of the first point
x2	x value of the second point
y2	y value of the second point

**Details**

linterp finds a linear function between two points.

**Value**

a linear equation's coefficients

**See Also**

Other interp: [bezier](#), [bilinear](#), [cubicspline](#), [nn](#), [polyinterp](#), [pwiselinterp](#)

Other algebra: [bilinear](#), [cubicspline](#), [division](#), [fibonacci](#), [horner](#), [isPrime](#), [nthroot](#), [polyinterp](#), [pwiselinterp](#), [quadratic](#)

**Examples**

```
f <- linterp(3, 2, 7, -2)
```

---

lumatrix

*LU Decomposition*

---

**Description**

Decompose a matrix into lower- and upper-triangular matrices

**Usage**

```
lumatrix(m)
```

**Arguments**

m                    a matrix

**Details**

lumatrix decomposes the matrix m into the LU decomposition, such that  $m == L$

**Value**

list with matrices L and U representing the LU decomposition

**See Also**

Other linear: [choleskymatrix](#), [detmatrix](#), [gdls](#), [invmatrix](#), [iterativematrix](#), [refmatrix](#), [rowops](#), [tridiagmatrix](#), [vecnorm](#)

**Examples**

```
A <- matrix(c(1, 2, -7, -1, -1, 1, 2, 1, 5), 3)
lumatrix(A)
```

---

mcint

*Monte Carlo Integration*

---

**Description**

Simple Monte Carlo Integrator

**Usage**

```
mcint(f, a, b, m = 1000)
```

```
mcint2(f, xdom, ydom, m = 1000)
```

**Arguments**

f	function to integrate
a	the lower-bound of integration
b	the upper-bound of integration
m	the number of subintervals to calculate
xdom	the domain on x of integration in two dimensions
ydom	the domain on y of integration in two dimensions

**Details**

The `mcint` function uses a simple Monte Carlo algorithm to estimate the value of an integral. The parameter `n` sets the total number of evaluation points. The parameter `max.y` is the maximum expected value of the range of function `f`. The `mcint2` provides Monte Carlo integration in two dimensions.

**Value**

the value of the integral

**See Also**

Other integration: [adaptint](#), [gaussint](#), [giniquintile](#), [midpt](#), [revolution-solid](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

**Examples**

```
f <- function(x) { sin(x)^2 + log(x) }
mcint(f, 0, 1)
mcint(f, 0, 1, m = 10e6)
```

---

midpt

*rectangle method*

---

**Description**

Use the rectangle method to integrate a function

**Usage**

```
midpt(f, a, b, m = 100)
```

**Arguments**

f	function to integrate
a	the a-bound of integration
b	the b-bound of integration
m	the number of subintervals to calculate

**Details**

The midpt function uses the rectangle method to calculate the integral of the function f over the interval from a to b. The parameter m sets the number of intervals to use when evaluating the rectangles. Additional options are passed to the function f when evaluating.

**Value**

the value of the integral

**See Also**

Other integration: [adaptint](#), [gaussint](#), [giniquintile](#), [mcint](#), [revolution-solid](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

Other newton-cotes: [adaptint](#), [giniquintile](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

**Examples**

```
f <- function(x) { sin(x)^2 + cos(x)^2 }
midpt(f, -pi, pi, m = 10)
midpt(f, -pi, pi, m = 100)
midpt(f, -pi, pi, m = 1000)
```

---

newton	<i>Newton's method</i>
--------	------------------------

---

### Description

Use Newton's method to find real roots

### Usage

```
newton(f, fp, x, tol = 0.001, m = 100)
```

### Arguments

f	function to integrate
fp	function representing the derivative of f
x	an initial estimate of the root
tol	the error tolerance
m	the maximum number of iterations

### Details

Newton's method finds real roots of a function, but requires knowing the function derivative. It will return when the interval between them is less than `tol`, the error tolerance. However, this implementation also stops after `m` iterations.

### Value

the real root found

### See Also

Other optimz: [bisection](#), [goldsect](#), [gradient](#), [hillclimbing](#), [sa](#), [secant](#)

### Examples

```
f <- function(x) { x^3 - 2 * x^2 - 159 * x - 540 }  
fp <- function(x) { 3 * x^2 - 4 * x - 159 }  
newton(f, fp, 1)
```

---

nn	<i>Nearest interpolation</i>
----	------------------------------

---

### Description

Find the nearest neighbor for a set of data points

### Usage

```
nn(p, y, q)
```

### Arguments

p	matrix of variable values, each row is a data point
y	vector of values, each entry corresponds to one row in p
q	vector of variable values, each entry corresponds to one column of p

### Details

nn finds the n-dimensional nearest neighbor for given datapoint

### Value

an interpolated value for  $q$

### See Also

Other interp: [bezier](#), [bilinear](#), [cubicspline](#), [linterp](#), [polyinterp](#), [pwiselinterp](#)

### Examples

```
p <- matrix(floor(runif(100, 0, 9)), 20)
y <- floor(runif(20, 0, 9))
q <- matrix(floor(runif(5, 0, 9)), 1)
nn(p, y, q)
```

---

nthroot	<i>The n-th root formula</i>
---------	------------------------------

---

### Description

Find the n-th root of real numbers

### Usage

```
nthroot(a, n, tol = 1/1000)
```

### Arguments

a	a positive real number
n	n
tol	the permitted error tolerance

### Details

The nthroot function finds the nth root of a via an iterative process.

### Value

the root

### See Also

Other algebra: [bilinear](#), [cubicspline](#), [division](#), [fibonacci](#), [horner](#), [isPrime](#), [linterp](#), [polyinterp](#), [pwiselinterp](#), [quadratic](#)

### Examples

```
nthroot(100, 2)
nthroot(65536, 4)
nthroot(1000, 3)
```

---

polyinterp	<i>Polynomial interpolation</i>
------------	---------------------------------

---

**Description**

Finds a polynomial function interpolating the given points

**Usage**

```
polyinterp(x, y)
```

**Arguments**

x	a vector of x values
y	a vector of y values

**Details**

polyinterp finds a polynomial that interpolates the given points.

**Value**

a polynomial equation's coefficients

**See Also**

Other interp: [bezier](#), [bilinear](#), [cubicspline](#), [linterp](#), [nn](#), [pwiselinterp](#)

Other algebra: [bilinear](#), [cubicspline](#), [division](#), [fibonacci](#), [horner](#), [isPrime](#), [linterp](#), [nthroot](#), [pwiselinterp](#), [quadratic](#)

**Examples**

```
x <- c(1, 2, 3)
y <- x^2 + 5 * x - 3
f <- polyinterp(x, y)
```



---

pwiselinterp

*Piecewise linear interpolation*

---

## Description

Finds a piecewise linear function that interpolates the data points

## Usage

```
pwiselinterp(x, y)
```

## Arguments

x	a vector of x values
y	a vector of y values

## Details

pwiselinterp finds a piecewise linear function that interpolates the data points. For each x-y ordered pair, there function finds the unique line interpolating them. The function will return a data.frame with three columns.

The column x is the upper bound of the domain for the given piece. The columns m and b represent the coefficients from the y-intercept form of the linear equation,  $y = mx + b$ .

The matrix will contain length(x) rows with the first row having m and b of NA.

## Value

a matrix with the linear function components

## See Also

Other interp: [bezier](#), [bilinear](#), [cubicspline](#), [linterp](#), [nn](#), [polyinterp](#)

Other algebra: [bilinear](#), [cubicspline](#), [division](#), [fibonacci](#), [horner](#), [isPrime](#), [linterp](#), [nthroot](#), [polyinterp](#), [quadratic](#)

## Examples

```
x <- c(5, 0, 3)
y <- c(4, 0, 3)
f <- pwiselinterp(x, y)
```

---

`quadratic`*The quadratic equation.*

---

**Description**

Find the zeros of a quadratic equation.

**Usage**

```
quadratic(b2, b1, b0)
```

```
quadratic2(b2, b1, b0)
```

**Arguments**

<code>b2</code>	the coefficient of the $x^2$ term
<code>b1</code>	the coefficient of the $x$ term
<code>b0</code>	the constant term

**Details**

`quadratic` and `quadratic2` implement the quadratic equation from standard algebra in two different ways. The `quadratic` function is susceptible to cascading numerical error and the `quadratic2` has reduced potential error.

**Value**

numeric vector of solutions to the equation

**See Also**

Other algebra: [bilinear](#), [cubicspline](#), [division](#), [fibonacci](#), [horner](#), [isPrime](#), [linterp](#), [nthroot](#), [polyinterp](#), [pwiselinterp](#)

**Examples**

```
quadratic(1, 0, -1)
quadratic(4, -4, 1)
quadratic2(1, 0, -1)
quadratic2(4, -4, 1)
```

---

`refmatrix`*Matrix to Row Echelon Form*

---

**Description**

Transform a matrix to row echelon form.

**Usage**`refmatrix(m)``rrefmatrix(m)``solvematrix(A, b)`**Arguments**

`m` a matrix

`A` a square matrix representing the coefficients of a linear system in `solvematrix`

`b` a vector representing the right-hand side of the linear system in `solvematrix`

**Details**

`refmatrix` reduces a matrix to row echelon form. This is not a reduced row echelon form, though that can be easily calculated from the diagonal. This function works on non-square matrices.

`rrefmatrix` returns the reduced row echelon matrix.

`solvematrix` solves a linear system using `rrefmatrix`.

**Value**

the modified matrix

**See Also**

Other linear: [choleskymatrix](#), [detmatrix](#), [gdls](#), [invmatrix](#), [iterativematrix](#), [lumatrix](#), [rowops](#), [tridiagmatrix](#), [vecnorm](#)

**Examples**

```
A <- matrix(c(1, 2, -7, -1, -1, 1, 2, 1, 5), 3)
refmatrix(A)
```

---

resizeImage	<i>Image resizing</i>
-------------	-----------------------

---

**Description**

Resize images using nearest neighbor and

**Usage**

```
resizeImageNN(imx, width, height)
```

```
resizeImageBL(imx, width, height)
```

**Arguments**

imx	a 3-dimensional array containing image data
width	the new width
height	the new height

**Details**

The *resizeImageNN* function uses the nearest neighbor method to resize the image. Also, *resizeImageBL* uses bilinear interpolation to resize the image.

**Value**

a three-dimensional array containing the resized image.

---

revolution-solid	<i>Volumes of solids of revolution</i>
------------------	--

---

**Description**

Find the volume of a solid of revolution

**Usage**

```
shellmethod(f, a, b)
```

```
discmethod(f, a, b)
```

**Arguments**

f	function of revolution
a	lower-bound of the solid
b	upper-bound of the solid

**Details**

The functions `discmethod` and `shellmethod` implement the algorithms for finding the volume of solids of revolution. The `discmethod` function is suitable for volumes revolved around the x-axis and the `shellmethod` function is suitable for volumes revolved around the y-axis.

**Value**

the volume of the solid

**See Also**

Other integration: [adaptint](#), [gaussint](#), [giniquintile](#), [mcint](#), [midpt](#), [romberg](#), [simp38](#), [simp](#), [trap](#)

**Examples**

```
f <- function(x) { x^2 }
shellmethod(f, 1, 2)
discmethod(f, 1, 2)
```

---

romberg

*Romberg Integration*


---

**Description**

Romberg's adaptive integration

**Usage**

```
romberg(f, a, b, m, tab = FALSE)
```

**Arguments**

<code>f</code>	function to integrate
<code>a</code>	the lowerbound of integration
<code>b</code>	the upperbound of integration
<code>m</code>	the maximum number of iterations
<code>tab</code>	if TRUE, return the table of values

**Details**

The `romberg` function uses Romberg's rule to calculate the integral of the function `f` over the interval from `a` to `b`. The parameter `m` sets the number of intervals to use when evaluating. Additional options are passed to the function `f` when evaluating.

**Value**

the value of the integral

**See Also**

Other integration: [adaptint](#), [gaussint](#), [giniquintile](#), [mcint](#), [midpt](#), [revolution-solid](#), [simp38](#), [simp](#), [trap](#)

Other newton-cotes: [adaptint](#), [giniquintile](#), [midpt](#), [simp38](#), [simp](#), [trap](#)

**Examples**

```
f <- function(x) { sin(x)^2 + log(x) }
romberg(f, 1, 10, m = 3)
romberg(f, 1, 10, m = 5)
romberg(f, 1, 10, m = 10)
```

---

rowops

*Elementary row operations*

---

**Description**

These are elementary operations for a matrix. They do not presume a square matrix and will work on any matrix. They use R's internal row addressing to function.

**Usage**

```
swaprows(m, row1, row2)
```

```
replacero(m, row1, row2, k)
```

```
scalero(m, row, k)
```

**Arguments**

m	a matrix
row1	a source row
row2	a destination row
k	a scaling factor
row	a row to modify

**Details**

replacero replaces one row with the sum of itself and the multiple of another row. swaprows swap two rows in the matrix. scalero scales all entries in a row by a constant.

**Value**

the modified matrix

**See Also**

Other linear: [choleskymatrix](#), [detmatrix](#), [gdls](#), [invmatrix](#), [iterativematrix](#), [lumatrix](#), [refmatrix](#), [tridiagmatrix](#), [vecnorm](#)

**Examples**

```
n <- 5
A <- matrix(sample.int(10, n^2, TRUE) - 1, n)
A <- swaprows(A, 2, 4)
A <- replacerow(A, 1, 3, 2)
A <- scalerow(A, 5, 10)
```

---

sa

*Simulated annealing*

---

**Description**

Use simulated annealing to find the global minimum

**Usage**

```
sa(f, x, temp = 10000, rate = 1e-04)
```

```
tspsa(x, temp = 100, rate = 1e-04)
```

**Arguments**

f	function representing f
x	an initial estimate of the minimum
temp	the initial temperature
rate	the cooling rate

**Details**

Simulated annealing finds a global minimum by mimicing the metallurgical process of annealing.

**Value**

the x value of the minimum found

**See Also**

Other optimz: [bisection](#), [goldsect](#), [gradient](#), [hillclimbing](#), [newton](#), [secant](#)

**Examples**

```
f <- function(x) { x^6 - 4 * x^5 - 7 * x^4 + 22 * x^3 + 24 * x^2 + 2 }
sa(f, 0)
```

```
f <- function(x) { (x[1] - 1)^2 + (x[2] - 1)^2 }
sa(f, c(0, 0), 0.05)
```

---

 secant

*Secant Method*


---

**Description**

The secant method for root finding

**Usage**

```
secant(f, x, tol = 0.001, m = 100)
```

**Arguments**

f	function to integrate
x	an initial estimate of the root
tol	the error tolerance
m	the maximum number of iterations

**Details**

The secant method for root finding extends Newton's method to estimate the derivative. It will return when the interval between them is less than `tol`, the error tolerance. However, this implementation also stop if after `m` iterations.

**Value**

the real root found

**See Also**

Other optimz: [bisection](#), [goldsect](#), [gradient](#), [hillclimbing](#), [newton](#), [sa](#)

**Examples**

```
f <- function(x) { x^3 - 2 * x^2 - 159 * x - 540 }
secant(f, 1)
```



---

`simp`*Simpson's rule*

---

**Description**

Use Simpson's rule to integrate a function

**Usage**

```
simp(f, a, b, m = 100)
```

**Arguments**

<code>f</code>	function to integrate
<code>a</code>	the a-bound of integration
<code>b</code>	the b-bound of integration
<code>m</code>	the number of subintervals to calculate

**Details**

The `simp` function uses Simpson's rule to calculate the integral of the function `f` over the interval from `a` to `b`. The parameter `m` sets the number of intervals to use when evaluating. Additional options are passed to the function `f` when evaluating.

**Value**

the value of the integral

**See Also**

Other integration: [adaptint](#), [gaussint](#), [giniquintile](#), [mcint](#), [midpt](#), [revolution-solid](#), [romberg](#), [simp38](#), [trap](#)

Other newton-cotes: [adaptint](#), [giniquintile](#), [midpt](#), [romberg](#), [simp38](#), [trap](#)

**Examples**

```
f <- function(x) { sin(x)^2 + cos(x)^2 }
simp(f, -pi, pi, m = 10)
simp(f, -pi, pi, m = 100)
simp(f, -pi, pi, m = 1000)
```

---

`simp38`*Simpson's 3/8 rule*

---

**Description**

Use Simpson's 3/8 rule to integrate a function

**Usage**

```
simp38(f, a, b, m = 100)
```

**Arguments**

<code>f</code>	function to integrate
<code>a</code>	the a-bound of integration
<code>b</code>	the b-bound of integration
<code>m</code>	the number of subintervals to calculate

**Details**

The `simp38` function uses Simpson's 3/8 rule to calculate the integral of the function `f` over the interval from `a` to `b`. The parameter `m` sets the number of intervals to use when evaluating. Additional options are passed to the function `f` when evaluating.

**Value**

the value of the integral

**See Also**

Other integration: [adaptint](#), [gaussint](#), [giniquintile](#), [mcint](#), [midpt](#), [revolution-solid](#), [romberg](#), [simp](#), [trap](#)

Other newton-cotes: [adaptint](#), [giniquintile](#), [midpt](#), [romberg](#), [simp](#), [trap](#)

**Examples**

```
f <- function(x) { sin(x)^2 + log(x) }
simp38(f, 1, 10, m = 10)
simp38(f, 1, 10, m = 100)
simp38(f, 1, 10, m = 1000)
```

---

`summation`*Two summing algorithms*

---

**Description**

Find the sum of a vector

**Usage**`naivesum(x)``kahansum(x)``pwisum(x)`**Arguments**

`x` a vector of numbers to be summed

**Details**

`naivesum` calculates the sum of a vector by keeping a counter and repeatedly adding the next value to the interim sum. `kahansum` uses Kahan's algorithm to capture the low-order precision loss and ensure that the loss is reintegrated into the final sum. `pwisum` is a recursive implementation of the piecewise summation algorithm that divides the vector in two and adds the individual vector sums for a result.

**Value**

the sum

**Examples**

```
k <- 1:10^6
n <- sample(k, 1)
bound <- sample(k, 2)
bound.upper <- max(bound) - 10^6 / 2
bound.lower <- min(bound) - 10^6 / 2
x <- runif(n, bound.lower, bound.upper)
naivesum(x)
kahansum(x)
pwisum(x)
```

---

trap

*Trapezoid method*

---

### Description

Use the trapezoid method to integrate a function

### Usage

```
trap(f, a, b, m = 100)
```

### Arguments

f	function to integrate
a	the a-bound of integration
b	the b-bound of integration
m	the number of subintervals to calculate

### Details

The trap function uses the trapezoid method to calculate the integral of the function  $f$  over the interval from  $a$  to  $b$ . The parameter  $m$  sets the number of intervals to use when evaluating the trapezoids. Additional options are passed to the function  $f$  when evaluating.

### Value

the value of the integral

### See Also

Other integration: [adaptint](#), [gaussint](#), [giniquintile](#), [mcint](#), [midpt](#), [revolution-solid](#), [romberg](#), [simp38](#), [simp](#)

Other newton-cotes: [adaptint](#), [giniquintile](#), [midpt](#), [romberg](#), [simp38](#), [simp](#)

### Examples

```
f <- function(x) { sin(x)^2 + cos(x)^2 }  
trap(f, -pi, pi, m = 10)  
trap(f, -pi, pi, m = 100)  
trap(f, -pi, pi, m = 1000)
```

---

tridiagmatrix	<i>Solve a tridiagonal matrix</i>
---------------	-----------------------------------

---

**Description**

use the tridiagonal matrix algorithm to solve a tridiagonal matrix

**Usage**

```
tridiagmatrix(L, D, U, b)
```

**Arguments**

L	vector of entries below the main diagonal
D	vector of entries on the main diagonal
U	vector of entries above the main diagonal
b	vector of the right-hand side of the linear system

**Details**

tridiagmatrix uses the tridiagonal matrix algorithm to solve a tridiagonal matrix.

**Value**

the solution vector

**See Also**

Other linear: [choleskymatrix](#), [detmatrix](#), [gdls](#), [invmatrix](#), [iterativematrix](#), [lumatrix](#), [refmatrix](#), [rowops](#), [vecnorm](#)

---

vecnorm	<i>Norm of a vector</i>
---------	-------------------------

---

**Description**

Find the norm of a vector

**Usage**

```
vecnorm(b)
```

**Arguments**

b	a vector
---	----------

**Details**

Find the norm of a vector

**Value**

the norm

**See Also**

Other linear: [choleskymatrix](#), [detmatrix](#), [gdls](#), [invmatrix](#), [iterativematrix](#), [lumatrix](#), [refmatrix](#), [rowops](#), [tridiagmatrix](#)

**Examples**

```
x <- c(1, 2, 3)
vecnorm(x)
```

---

wave

*Wave Equation using*

---

**Description**

solve heat equation via forward-time central-space method

**Usage**

```
wave(u, alpha, xdelta, tdelta, n)
```

**Arguments**

u	the initial values of u
alpha	the thermal diffusivity coefficient
xdelta	the change in x at each step in u
tdelta	the time step
n	the number of steps to take

**Details**

The heat solves the heat equation using the forward-time central-space method in one-dimension.

**Value**

a matrix of u values at each time step

**Examples**

```
speed <- 2
x0 <- 0
xdelta <- .05
x <- seq(x0, 1, xdelta)
m <- length(x)
u <- sin(x * pi * 2)
u[11:21] <- 0
tdelta <- .02
n <- 40
z <- wave(u, speed, xdelta, tdelta, n)
```

---

wilkinson

*Wilkinson's Polynomial*

---

**Description**

Wilkinson's polynomial

**Usage**

```
wilkinson(x, w = 20)
```

**Arguments**

x	the x-value
w	the number of terms in the polynomial

**Details**

Wilkinson's polynomial is a terrible joke played on numerical analysis. By tradition, the function is  $f(x) = (x - 1)(x - 2)\dots(x - 20)$ , giving a function with real roots at each integer from 1 to 20. This function is generalized and allows for  $n$  and the function value is  $f(x) = (x - 1)(x - 2)\dots(x - n)$ . The default of  $n$  is 20.

**Value**

the value of the function at  $x$

**Examples**

```
wilkinson(0)
```

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