

# Package ‘rje’

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**Type** Package

**Title** Miscellaneous Useful Functions for Statistics

**Version** 1.10.10

**Description** A series of functions in some way considered useful to the author. These include functions for subsetting tables and generating indices for arrays, conditioning and intervening in probability distributions, generating combinations and more...

**Depends** R (>= 2.0.0),

**License** GPL (>= 2)

**LazyLoad** yes

**Imports** knitr

**Suggests** testthat, rmarkdown

**VignetteBuilder** knitr

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## R topics documented:

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rje-package

*Miscellaneous useful functions*

---

## Description

A series of useful functions, some available in different forms in other packages, but which have been extended, sped up, or otherwise modified in some way considered useful to the author.

## Details

|           |            |
|-----------|------------|
| Package:  | rje        |
| Type:     | Package    |
| Version:  | 1.10.7     |
| Date:     | 2017-08-14 |
| License:  | GPL (>= 2) |
| LazyLoad: | yes        |

## Author(s)

Robin Evans

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Maintainer: Robin Evans <evans@stats.ox.ac.uk>

---

`and0`*Fast pairwise logical operators*

---

**Description**

Fast but loose implementations of AND and OR logical operators.

**Usage**

```
and0(x, y)
```

**Arguments**

`x, y`                    logical or numerical vectors

**Details**

Returns pairwise application of logical operators AND and OR. Vectors are recycled as usual.

**Value**

A logical vector of length  $\max(\text{length}(x), \text{length}(y))$  with entries `x[1] & x[2]` etc.; each entry of `x` or `y` is TRUE if it is non-zero.

**Note**

These functions should only be used with well understood vectors, and may not deal with unusual cases correctly.

**Examples**

```
and0(c(0,1,0), c(1,1,0))
## Not run:
set.seed(1234)
x = rbinom(5000, 1, 0.5)
y = rbinom(5000, 1, 0.5)

# 3 to 4 times improvement over `&`
system.time(for (i in 1:5000) and0(x,y))
system.time(for (i in 1:5000) x & y)

## End(Not run)
```

**Description**

Allows use of an Armijo rule or coarse line search as part of minimisation (or maximisation) of a differentiable function of multiple arguments (via gradient descent or similar). Repeated application of one of these rules should (hopefully) lead to a local minimum.

**Usage**

```
armijo(fun, x, dx, beta = 3, sigma = 0.5, grad, maximise = FALSE,
       searchup = TRUE, adj.start = 1, ...)
```

```
coarseLine(fun, x, dx, beta = 3, maximise = FALSE, ...)
```

**Arguments**

|           |                                                                                                                                        |
|-----------|----------------------------------------------------------------------------------------------------------------------------------------|
| fun       | a function whose first argument is a numeric vector                                                                                    |
| x         | a starting value to be passed to fun                                                                                                   |
| dx        | numeric vector containing feasible direction for search; defaults to $-\text{grad}$ for ordinary gradient descent                      |
| beta      | numeric value (greater than 1) giving factor by which to adjust step size                                                              |
| sigma     | numeric value (less than 1) giving steepness criterion for move                                                                        |
| grad      | numeric gradient of $f$ at $x$ (will be estimated if not provided)                                                                     |
| maximise  | logical: if set to TRUE search is for a maximum rather than a minimum.                                                                 |
| searchup  | logical: if set to TRUE method will try to find largest move satisfying Armijo criterion, rather than just accepting the first it sees |
| adj.start | an initial adjustment factor for the step size.                                                                                        |
| ...       | other arguments to be passed to fun                                                                                                    |

**Details**

coarseLine performs a stepwise search and tries to find the integer  $k$  minimising  $f(x_k)$  where

$$x_k = x + \beta^k dx.$$

Note  $k$  may be negative. This is generally quicker and dirtier than the Armijo rule.

armijo implements an Armijo rule for moving, which is to say that

$$f(x_k) - f(x) < -\sigma \beta^k dx \cdot \nabla_x f.$$

This has better convergence guarantees than a simple line search, but may be slower in practice. See Bertsekas (1999) for theory underlying the Armijo rule.

Each of these rules should be applied repeatedly to achieve convergence (see example below).

**Value**

A list comprising

|      |                                                                                                                                                                                                                                                                                                                                             |
|------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| best | the value of the function at the final point of evaluation                                                                                                                                                                                                                                                                                  |
| adj  | the constant in the step, i.e. $\beta^n$                                                                                                                                                                                                                                                                                                    |
| move | the final move; i.e. $\beta^n dx$                                                                                                                                                                                                                                                                                                           |
| code | an integer indicating the result of the function; 0 = returned OK, 1 = very small move suggested, may be at minimum already, 2 = failed to find minimum: function evaluated to NA or was always larger than $f(x)$ (direction might be infeasible), 3 = failed to find minimum: stepsize became too small or large without satisfying rule. |

**Functions**

- coarseLine: Coarse line search

**Author(s)**

Robin Evans

**References**

Bertsekas, D.P. *Nonlinear programming*, 2nd Edition. Athena, 1999.

**Examples**

```
# minimisation of simple function of three variables
x = c(0,-1,4)
f = function(x) ((x[1]-3)^2 + sin(x[2])^2 + exp(x[3]) - x[3])

tol = .Machine$double.eps
mv = 1

while (mv > tol) {
  # or replace with coarseLine()
  out = armijo(f, x, sigma=0.1)
  x = out$x
  mv = sum(out$move^2)
}

# correct solution is c(3,0,0) (or c(3,k*pi,0) for any integer k)
x
```

---

`combinations`*Combinations of Integers*

---

**Description**

Returns a matrix containing each possible combination of one entry from vectors of the lengths provided.

**Usage**

```
combinations(p)
powerSetMat(n)
```

**Arguments**

|                |                                  |
|----------------|----------------------------------|
| <code>p</code> | vector of non-negative integers. |
| <code>n</code> | non-negative integer.            |

**Details**

Returns a matrix, each row being one possible combination of integers from the vectors  $(0, 1, \dots, p_i - 1)$ , for  $i$  between 1 and `length(p)`.

Based on `bincombinations` from package `e1071`, which provides the binary case.

`powerSetMat` is just a wrapper for `combinations(rep(2, n))`.

**Value**

A matrix with number of columns equal to the length of `p`, and number of rows equal to  $p_1 \times \dots \times p_k$ , each row corresponding to a different combination. Ordering is reverse-lexographic.

**Author(s)**

Robin Evans

**Examples**

```
combinations(c(2, 3, 3))
powerSetMat(3)
```

---

|                 |                                           |
|-----------------|-------------------------------------------|
| conditionMatrix | <i>Find conditional probability table</i> |
|-----------------|-------------------------------------------|

---

### Description

Given a numeric array or matrix (of probabilities), calculates margins of some dimensions conditional on particular values of others.

### Usage

```
conditionMatrix(x, variables, condition = NULL, condition.value = NULL,
  dim = NULL, incol = FALSE, undef = NaN)
```

```
conditionTable(x, variables, condition = NULL, condition.value = NULL,
  undef = NaN, order = TRUE)
```

```
conditionTable2(x, variables, condition, undef = NaN)
```

### Arguments

|                 |                                                                                                                                                                  |
|-----------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| x               | A numeric array.                                                                                                                                                 |
| variables       | An integer vector containing the margins of interest from x.                                                                                                     |
| condition       | An integer vector containing the dimensions of x to condition on.                                                                                                |
| condition.value | An integer vector or list of the same length as condition, containing the values to condition with. If NULL, then the full conditional distribution is returned. |
| dim             | Integer vector containing dimensions of variables. Assumed all binary if not specified.                                                                          |
| incol           | Logical specifying whether not the distributions are stored as the columns in the matrix; assumed to be rows by default.                                         |
| undef           | if conditional probability is undefined, what should the value be given as                                                                                       |
| order           | logical - if TRUE conditioned variables come last, if FALSE variables are in original order.                                                                     |

### Details

conditionTable calculates the marginal distribution over the dimensions in variables for each specified value of the dimensions in condition. Single or multiple values of each dimension in condition may be specified in condition.value; in the case of multiple values, condition.value must be a list.

The sum over the dimensions in variables is normalized to 1 for each value of condition.

conditionTable2 is just a wrapper which returns the conditional distribution as an array of the same dimensions and ordering as the original x. Values are repeated as necessary.

`conditionMatrix` takes a matrix whose rows (or columns if `incols = TRUE`) each represent a separate multivariate probability distribution and finds the relevant conditional distribution in each case. These are then returned in the same format. The order of the variables under `conditionMatrix` is always as in the original distribution, unlike for `conditionTable` above.

The probabilities are assumed in reverse lexicographic order, as in a flattened R array: i.e. the first value changes fastest: (1,1,1), (2,1,1), (1,2,1), ..., (2,2,2).

`condition.table` and `condition.table2` are identical to `conditionTable` and `conditionTable2`.

### Value

`conditionTable` returns an array whose first `length(variables)` corresponds to the dimensions in `variables`, and the remainder (if any) to dimensions in `condition` with a corresponding entry in `condition.value` of length > 1.

`conditionTable2` always returns an array of the same dimensions as `x`, with the variables in the same order.

### Functions

- `conditionMatrix`: Conditioning in matrix of distributions
- `conditionTable2`: Conditioning whilst preserving all dimensions

### Author(s)

Mathias Drton, Robin Evans

### See Also

[marginTable](#), [margin.table](#), [interventionTable](#)

### Examples

```
x = array(1:16, rep(2,4))
x = x/sum(x) # probability distribution on 4 binary variables x1, x2, x3, x4.

# distribution of x2, x3 given x1 = 1 and x4=2.
conditionTable(x, c(2,3), c(1,4), c(1,2))
# x2, x3 given x1 = 1,2 and x4 = 2.
conditionTable(x, c(2,3), c(1,4), list(1:2,2))

# complete conditional of x2, x3 given x1, x4
conditionTable(x, c(2,3), c(1,4))

# conditionTable2 leaves dimensions unchanged
tmp = conditionTable2(x, c(2,3), c(1,4))
aperm(tmp, c(2,3,1,4))

####
set.seed(2314)
# set of 10 2x2x2 probability distributions
```



```
x = rdirichlet(10, rep(1,8))  
  
conditionMatrix(x, 3, 1)  
conditionMatrix(x, 3, 1, 2)
```

---

cubeHelix

*Cube Helix colour palette*

---

### Description

Cube Helix is a colour scheme designed to be appropriate for screen display of intensity images. The scheme is intended to be monotonically increasing in brightness when displayed in greyscale. This might also provide improved visualisation for colour blindness sufferers.

### Usage

```
cubeHelix(n, start = 0.5, r = -1.5, hue = 1, gamma = 1)
```

### Arguments

|       |                                                                                        |
|-------|----------------------------------------------------------------------------------------|
| n     | integer giving the number of colours in the scale                                      |
| start | numeric: start gives the initial angle (in radians) of the helix                       |
| r     | numeric: number of rotations of the helix over the scale; can be negative              |
| hue   | numeric controlling the saturation of colour: 0 gives pure greyscale, defaults to 1    |
| gamma | numeric which can be used to emphasise lower or higher intensity values, defaults to 1 |

### Details

The function evaluates a helix which moves through the RGB "cube", beginning at black (0,0,0) and finishing at white (1,1,1). Evenly spaced points on this helix in the cube are returned as RGB colours. This provides a colour palette in which intensity increases monotonically, which makes for good transfer to greyscale displays or printouts. This also may have advantages for colour blindness sufferers. See references for further details.

### Value

Vector of RGB colours (strings) of length n.

### Author(s)

Dave Green  
Robin Evans

## References

Green, D. A., 2011, A colour scheme for the display of astronomical intensity images. *Bulletin of the Astronomical Society of India*, 39, 289. <http://adsabs.harvard.edu/abs/2011BASI...39..289G>

See Dave Green's page at <http://www.mrao.cam.ac.uk/~dag/CUBEHELIX/> for other details.

## See Also

[rainbow](#) (for other colour palettes).

## Examples

```
cubeHelix(21)

## Not run:
cols = cubeHelix(101)

plot.new()
plot.window(xlim=c(0,1), ylim=c(0,1))
axis(side=1)
for (i in 1:101) {
  rect((i-1)/101,0,(i+0.1)/101,1, col=cols[i], lwd=0)
}

## End(Not run)

## Not run:
require(grDevices)
# comparison with other palettes
n = 101
cols = cubeHelix(n)
heat = heat.colors(n)
rain = rainbow(n)
terr = terrain.colors(n)

plot.new()
plot.window(xlim=c(-0.5,1), ylim=c(0,4))
axis(side=1, at=c(0,1))
axis(side=2, at=1:4-0.5, labels=1:4, pos=0)
for (i in 1:n) {
  rect((i-1)/n,3,(i+0.1)/n,3.9, col=cols[i], lwd=0)
  rect((i-1)/n,2,(i+0.1)/n,2.9, col=heat[i], lwd=0)
  rect((i-1)/n,1,(i+0.1)/n,1.9, col=rain[i], lwd=0)
  rect((i-1)/n,0,(i+0.1)/n,0.9, col=terr[i], lwd=0)
}
legend(-0.6,4,legend=c("4. cube helix", "3. heat", "2. rainbow", "1. terrain"), box.lwd=0)

## End(Not run)
```

---

|              |                                 |
|--------------|---------------------------------|
| designMatrix | <i>Orthogonal Design Matrix</i> |
|--------------|---------------------------------|

---

**Description**

Produces a matrix whose rows correspond to an orthogonal binary design matrix.

**Usage**

```
designMatrix(n)
```

**Arguments**

n integer containing the number of elements in the set.

**Value**

An integer matrix of dimension  $2^n$  by  $2^n$  containing 1 and -1.

**Note**

The output matrix has orthogonal columns and is symmetric, so (up to a constant) is its own inverse. Operations with this matrix can be performed more efficiently using the fast Hadamard transform.

**Author(s)**

Robin Evans

**See Also**

[combinations](#), [subsetMatrix](#).

**Examples**

```
designMatrix(3)
```

**Description**

Density function and random generation for Dirichlet distribution with parameter vector alpha.

**Usage**

```
ddirichlet(x, alpha, log = FALSE, tol = 1e-10)
rdirichlet(n, alpha)
```

**Arguments**

|       |                                                             |
|-------|-------------------------------------------------------------|
| x     | vector (or matrix) of points in sample space.               |
| alpha | vector of Dirichlet hyper parameters.                       |
| log   | logical; if TRUE, natural logarithm of density is returned. |
| tol   | tolerance of vectors not summing to 1 and negative values.  |
| n     | number of random variables to be generated.                 |

**Details**

If x is a matrix, each row is taken to be a different point whose density is to be evaluated. If the number of columns in (or length of, in the alpha, the vector sum to 1.

The  $k$ -dimensional Dirichlet distribution has density

$$\frac{\Gamma(\sum_i \alpha_i)}{\prod_i \Gamma(\alpha_i)} \prod_{i=1}^k x_i^{\alpha_i - 1}$$

assuming that  $x_i > 0$  and  $\sum_i x_i = 1$ , and zero otherwise.

If the sum of row entries in x differs from 1 by more than tol, is assumed to be

**Value**

rdirichlet returns a matrix, each row of which is an independent draw alpha.

ddirichlet returns a vector, each entry being the density of the corresponding row of x. If x is a vector, then the output will have length 1.

**Author(s)**

Robin Evans

**References**

[http://en.wikipedia.org/wiki/Dirichlet\\_distribution](http://en.wikipedia.org/wiki/Dirichlet_distribution)

**Examples**

```
x = rdirichlet(10, c(1,2,3))
x

# Find densities at random points.
ddirichlet(x, c(1,2,3))
# Last column to be inferred.
ddirichlet(x[,c(1,2)], c(1,2,3))
ddirichlet(x, matrix(c(1,2,3), 10, 3, byrow=TRUE))
```

---

 expit

*Expit and Logit.*


---

**Description**

Functions to take the expit and logit of numerical vectors.

**Usage**

```
expit(x)
```

**Arguments**

`x` vector of real numbers; for `logit` to return a sensible value these should be between 0 and 1.

**Details**

`logit` implements the usual logit function, which is

$$\text{logit}(x) = \log \frac{x}{1-x},$$

and `expit` its inverse:

$$\text{expit}(x) = \frac{e^x}{1+e^x}.$$

It is assumed that  $\text{logit}(0) = -\text{Inf}$  and  $\text{logit}(1) = \text{Inf}$ , and correspondingly for `expit`.

**Value**

A real vector corresponding to the expits or logits of `x`

**Warning**

Choosing very large (positive or negative) values to apply to `expit` may result in inaccurate inversion (see example below).

**Author(s)**

Robin Evans

**Examples**

```
x = c(5, -2, 0.1)
y = expit(x)
logit(y)

# Beware large values!
logit(expit(100))
```

---

`fastHadamard`*Compute fast Hadamard-transform of vector*

---

**Description**

Passes vector through Hadamard orthogonal design matrix. Also known as the Fast Walsh-Hadamard transform.

**Usage**

```
fastHadamard(x, pad = FALSE)
```

**Arguments**

|                  |                                                                                         |
|------------------|-----------------------------------------------------------------------------------------|
| <code>x</code>   | vector of values to be transformed                                                      |
| <code>pad</code> | optional logical asking whether vector not of length $2^k$ should be padded with zeroes |

**Details**

This is equivalent to multiplying by `designMatrix(log2(length(x)))` but should run much faster

**Value**

A vector of the same length as `x`

**Author(s)**

Robin Evans

**See Also**

[designMatrix](#), [subsetMatrix](#).

**Examples**

```
fastHadamard(1:8)
fastHadamard(1:5, pad=TRUE)
```

---

fsapply

*Fast and loose application of function over list.*

---

**Description**

Faster highly stripped down version of sapply()

**Usage**

```
fsapply(x, FUN)
```

**Arguments**

|     |                                                                                                                                   |
|-----|-----------------------------------------------------------------------------------------------------------------------------------|
| x   | a vector (atomic or list) or an expression object.                                                                                |
| FUN | the function to be applied to each element of x. In the case of functions like +, the function name must be backquoted or quoted. |

**Details**

This is just a wrapper for `unlist(lapply(x, FUN))`, which will behave as `sapply` if `FUN` returns an atomic vector of length 1 each time.

Speed up over `sapply` is not dramatic, but can be useful in time critical code.

**Value**

A vector of results of applying `FUN` to `x`.

**Warning**

Very loose version of `sapply` which should really only be used if you're confident about how `FUN` is applied to each entry in `x`.

**Author(s)**

Robin Evans

**Examples**

```
x = list(1:1000)
tmp = fsapply(x, sin)

## Not run:
x = list()
set.seed(142313)
for (i in 1:1000) x[[i]] = rnorm(100)

system.time(for (i in 1:100) sapply(x, function(x) last(x)))
system.time(for (i in 1:100) fsapply(x, function(x) last(x)))

## End(Not run)
```

---

greaterThan

*Comparing numerical values*

---

**Description**

Just a wrapper for comparing numerical values, for use with quicksort.

**Usage**

```
greaterThan(x, y)
```

**Arguments**

|   |                   |
|---|-------------------|
| x | A numeric vector. |
| y | A numeric vector. |

**Details**

Just returns -1 if x is less than y, 1 if x is greater, and 0 if they are equal (according to ==). The vectors wrap as usual if they are of different lengths.

**Value**

An integer vector.

**Author(s)**

Robin Evans

**See Also**

`<` for traditional Boolean operator.



**Examples**

```
greaterThan(4,6)

# Use in sorting algorithm.
quickSort(c(5,2,9,7,6), f=greaterThan)
order(c(5,2,9,7,6))
```

---

|          |                                                 |
|----------|-------------------------------------------------|
| indexBox | <i>Get indices of adjacent entries in array</i> |
|----------|-------------------------------------------------|

---

**Description**

Determines the relative vector positions of entries which are adjacent in an array.

**Usage**

```
indexBox(upp, lwr, dim)
```

**Arguments**

|     |                                                                                                                       |
|-----|-----------------------------------------------------------------------------------------------------------------------|
| upp | A vector of non-negative integers, giving the distance in the positive direction from the centre in each co-ordinate. |
| lwr | A vector of non-positive integers, giving the negative distance from the centre.                                      |
| dim | integer vector of array dimensions.                                                                                   |

**Details**

Given a particular cell in an array, which are the entries within (for example) 1 unit in any direction? This function gives the (relative) value of such indices. See examples.

Indices may be repeated if the range exceeds the size of the array in any dimension.

**Value**

An integer vector giving relative positions of the indices.

**Author(s)**

Robin Evans

**See Also**

[arrayInd](#).

**Examples**

```

arr = array(1:144, dim=c(3,4,3,4))
arr[2,2,2,3]
# which are entries within 1 unit each each direction of 2,2,2,3?

inds = 89 + indexBox(1,-1,c(3,4,3,4))
inds = inds[inds > 0 & inds <= 144]
arrayInd(inds, c(3,4,3,4))

# what about just in second dimension?
inds = 89 + indexBox(c(0,1,0,0),c(0,-1,0,0),c(3,4,3,4))
inds = inds[inds > 0 & inds <= 144]
arrayInd(inds, c(3,4,3,4))

```

---

interventionMatrix     *Calculate interventional distributions.*

---

**Description**

Calculate interventional distributions from a probability table or matrix of multivariate probability distributions.

**Usage**

```

interventionMatrix(x, variables, condition, dim = NULL, incol = FALSE)

interventionTable(x, variables, condition)

```

**Arguments**

|           |                                                                                                                          |
|-----------|--------------------------------------------------------------------------------------------------------------------------|
| x         | An array of probabilities.                                                                                               |
| variables | The margin for the intervention.                                                                                         |
| condition | The dimensions to be conditioned upon.                                                                                   |
| dim       | Integer vector containing dimensions of variables. Assumed all binary if not specified.                                  |
| incol     | Logical specifying whether not the distributions are stored as the columns in the matrix; assumed to be rows by default. |

**Details**

This just divides the joint distribution  $p(x)$  by  $p(v|c)$ , where  $v$  is variables and  $c$  is condition.

Under certain causal assumptions this is the interventional distribution  $p(x | do(v))$  (i.e. if the direct causes of  $v$  are precisely  $c$ .)

intervention.table() is identical to interventionTable().

**Value**

A numerical array of the same dimension as  $x$ .

**Functions**

- `interventionMatrix`: Interventions in matrix of distributions

**Author(s)**

Robin Evans

**References**

Pearl, J., *Causality*, 2nd Edition. Cambridge University Press, 2009.

**See Also**

[conditionTable](#), [marginTable](#)

**Examples**

```
set.seed(413)
# matrix of distributions
p = rdirichlet(10, rep(1,16))
interventionMatrix(p, 3, 2)

# take one in an array
ap = array(p[1,], rep(2,4))
interventionTable(ap, 3, 2)
```

---

is.subset

*Check subset inclusion*

---

**Description**

Determines whether one vector contains all the elements of another.

**Usage**

```
is.subset(x, y)
```

**Arguments**

|   |         |
|---|---------|
| x | vector. |
| y | vector. |

**Details**

Determines whether or not every element of `x` is also found in `y`. Returns TRUE if so, and FALSE if not.

**Value**

A logical of length 1.

**Author(s)**

Robin Evans

**See Also**

[setmatch](#).

**Examples**

```
is.subset(1:2, 1:3)
is.subset(1:2, 2:3)
```

---

is.wholenumber

*Determine whether number is integral or not.*

---

**Description**

Checks whether a numeric value is integral, up to machine or other specified precision.

**Usage**

```
is.wholenumber(x, tol = .Machine$double.eps^0.5)
```

**Arguments**

|                  |                              |
|------------------|------------------------------|
| <code>x</code>   | numeric vector to be tested. |
| <code>tol</code> | The desired precision.       |

**Value**

A logical vector of the same length as `x`, containing the results of the test.

**Author(s)**

Robin Evans

**Examples**

```
x = c(0.5, 1, 2L, 1e-20)
is.wholenumber(x)
```

---

|      |                                         |
|------|-----------------------------------------|
| last | <i>Last element of a vector or list</i> |
|------|-----------------------------------------|

---

**Description**

Returns the last element of a list or vector.

**Usage**

```
last(x)
```

**Arguments**

x                    a list or vector.

**Details**

Designed to be faster than using `tail()` or `rev()`, and cleaner than writing `x[length(x)]`.

**Value**

An object of the same type as `x` of length 1 (or empty if `x` is empty).

**Author(s)**

Robin Evans

**See Also**

[tail](#), [rev](#).

**Examples**

```
last(1:10)
```

---

|             |                                         |
|-------------|-----------------------------------------|
| marginTable | <i>Compute margin of a table faster</i> |
|-------------|-----------------------------------------|

---

### Description

Computes the margin of a contingency table given as an array, by summing out over the dimensions not specified.

### Usage

```
marginTable(x, margin = NULL, order = TRUE)
marginMatrix(x, margin, dim = NULL, incols = FALSE, order = FALSE)
```

### Arguments

|        |                                                                                                                                  |
|--------|----------------------------------------------------------------------------------------------------------------------------------|
| x      | a numeric array                                                                                                                  |
| margin | integer vector giving margin to be calculated (1 for rows, etc.)                                                                 |
| order  | logical - should indices of output be ordered as in the vector margin? Defaults to TRUE for marginTable, FALSE for marginMatrix. |
| dim    | Integer vector containing dimensions of variables. Assumed all binary if not specified.                                          |
| incols | Logical specifying whether not the distributions are stored as the columns in the matrix; assumed to be rows by default.         |

### Details

With `order = TRUE` this is the same as the base function `margin.table()`, but faster.

With `order = FALSE` the function is even faster, but the indices in the margin are returned in their original order, regardless of the way they are specified in `margin`.

`propTable()` returns a renormalized contingency table whose entries sum to 1. It is equivalent to `prop.table()`, but faster.

### Value

The relevant marginal table. The class of `x` is copied to the output table, except in the summation case.

### Note

Original functions are [margin.table](#) and [prop.table](#).

**Examples**

```

m <- matrix(1:4, 2)
marginTable(m, 1)
marginTable(m, 2)

propTable(m, 2)

# 3-way example
m <- array(1:8, rep(2,3))
marginTable(m, c(2,3))
marginTable(m, c(3,2))
marginTable(m, c(3,2), order=FALSE)

#' set.seed(2314)
# set of 10 2x2x2 probability distributions
x = rdirichlet(10, rep(1,8))

marginMatrix(x, c(1,3))
marginMatrix(t(x), c(1,3), incols=TRUE)

```

---

patternRepeat

*Complex repetitions*

---

**Description**

Recreate patterns for collapsed arrays

**Usage**

```
patternRepeat(x, which, n, careful = TRUE, keep.order = FALSE)
```

**Arguments**

|            |                                                                                                                                                                                                                                  |
|------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| x          | A vector to be repeated.                                                                                                                                                                                                         |
| which      | Which indices of the implicit array are given in x.                                                                                                                                                                              |
| n          | Dimensions of implicit array.                                                                                                                                                                                                    |
| careful    | logical indicating whether to check validity of arguments, but therefore slow things down.                                                                                                                                       |
| keep.order | logical indicating whether to respect the ordering of the entries in the vector which, in which case data are permuted before replication. In other words, does x change fastest in which[1], or in the minimal entry for which? |

**Details**

These functions allow for the construction of complex repeating patterns corresponding to those obtained by unwrapping arrays. Consider an array with dimensions  $n$ ; then for each value of the dimensions in which, this function returns a vector which places the corresponding entry of  $x$  into every place which would match this pattern when the full array is unwrapped.

For example, if a full 4-way array has dimensions  $2*2*2*2$  and we consider the margin of variables 2 and 4, then the function returns the pattern  $c(1,1,2,2,1,1,2,2,3,3,4,4,3,3,4,4)$ . The entries 1,2,3,4 correspond to the patterns (0,0), (1,0), (0,1) and (1,1) for the 2nd and 4th indices.

In `patternRepeat()` the argument  $x$  is repeated according to the pattern, while `patternRepeat0()` just returns the indexing pattern. So `patternRepeat(x,which,n)` is effectively equivalent to `x[patternRepeat0(which,n)]`.

The length of  $x$  must be equal to `prod(n[which])`.

**Value**

Both return a vector of length `prod(n)`; `patternRepeat()` one containing suitably repeated and ordered elements of  $x$ , for `patternRepeat0()` it is always the integers from 1 up to `prod(n[which])`.

**Author(s)**

Robin Evans

**See Also**

[rep](#)

**Examples**

```
patternRepeat(1:4, c(1,2), c(2,2,2))
c(array(1:4, c(2,2,2)))

patternRepeat0(c(1,3), c(2,2,2))
patternRepeat0(c(2,3), c(2,2,2))

patternRepeat0(c(3,1), c(2,2,2))
patternRepeat0(c(3,1), c(2,2,2), keep.order=TRUE)

patternRepeat(letters[1:4], c(1,3), c(2,2,2))
```



---

`powerSet`*Power Set*

---

**Description**

Produces the power set of a vector.

**Usage**

```
powerSet(x, m, rev = FALSE)
```

**Arguments**

|                  |                                                             |
|------------------|-------------------------------------------------------------|
| <code>x</code>   | vector of elements (the set).                               |
| <code>m</code>   | maximum cardinality of subsets                              |
| <code>rev</code> | logical indicating whether to reverse the order of subsets. |

**Details**

Creates a list containing every subset of the elements of the vector `x`.

**Value**

A list of vectors of the same type as `x`.

With `rev = FALSE` (the default) the list is ordered such that all subsets containing the last element of `x` come after those which do not, and so on.

**Author(s)**

Robin Evans

**See Also**

[powerSetMat](#).

**Examples**

```
powerSet(1:3)
powerSet(letters[3:5], rev=TRUE)
powerSet(1:5, m=2)
```

---

printPercentage      *Print Percentage of Activity Completed to stdout*

---

### Description

Prints percentage (or alternatively just a count) of loop or similar process which has been completed to the standard output.

### Usage

```
printPercentage(i, n, dp = 0, first = 1, last = n, prev = i - 1)
```

### Arguments

|       |                                                                           |
|-------|---------------------------------------------------------------------------|
| i     | the number of iterations completed.                                       |
| n     | total number of iterations.                                               |
| dp    | number of decimal places to display.                                      |
| first | number of the first iteration for which this percentage was displayed     |
| last  | number of the final iteration for which this percentage will be displayed |
| prev  | number of the previous iteration for which this percentage was displayed  |

### Details

printPercentage will use cat to print the proportion of loops which have been completed (i.e.  $i/n$ ) to the standard output. In doing so it will erase the previous such percentage, except when  $i = \text{first}$ . A new line is added when  $i = \text{last}$ , assuming that the loop is finished.

### Value

NULL

### Warning

This will fail to work nicely if other information is printed to the standard output

### Author(s)

Robin Evans

### Examples

```
x = numeric(100)

for (i in 1:100) {
  x[i] = mean(rnorm(1e5))
  printPercentage(i, 100)
}
```

```
}

i = 0
repeat {
  i = i+1
  if (runif(1) > 0.99) {
    break
  }
  printCount(i)
}
print("\n")
```

---

quickSort

*Quicksort for Partial Orderings*

---

### Description

Implements the quicksort algorithm for partial orderings based on pairwise comparisons.

### Usage

```
quickSort(x, f = greaterThan, random = TRUE)
```

### Arguments

|        |                                                                                                                                                                                |
|--------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| x      | A list or vector of items to be sorted.                                                                                                                                        |
| f      | A function on two arguments for comparing elements of x. Returns -1 if the first argument is less than the second, 1 for the reverse, and 0 if they are equal or incomparable. |
| random | logical - should a random pivot be chosen? (this is recommended) Otherwise middle element is used.                                                                             |

### Details

Implements the usual quicksort algorithm, but may return the same positions for items which are incomparable (or equal). Does not test the validity of f as a partial order.

If x is a numeric vector with distinct entries, this behaves just like [order](#).

### Value

Returns an integer vector giving each element's position in the order (minimal element(s) is 1, etc).

**Warning**

Output may not be consistent for certain partial orderings (using random pivot), see example below. All results will be consistent with a total ordering which is itself consistent with the true partial ordering.

f is not checked to see that it returns a legitimate partial order, so results may be meaningless if it is not.

**Author(s)**

Robin Evans

**References**

<http://en.wikipedia.org/wiki/Quicksort>.

**See Also**

[order](#).

**Examples**

```
set.seed(1)
quickSort(powerSet(1:3), f=subsetOrder)
quickSort(powerSet(1:3), f=subsetOrder)
# slightly different answers, but both corresponding
# to a legitimate total ordering.
```

---

rowMins

*Row-wise minima and maxima*

---

**Description**

Row-wise minima and maxima

**Usage**

```
rowMins(x)
rowMaxs(x)
```

**Arguments**

x                    a numeric (or logical) matrix or data frame

**Details**

The function coerces x to be a data frame and then uses pmin (pmax) on it. This is the same as apply(x, 1, min) but generally faster if the number of rows is large.

**Value**

numeric vector of length `nrow(x)` giving the row-wise minima (or maxima) of `x`.

---

rprobdist

*Generate a joint (or conditional) probability distribution*

---

**Description**

Wrapper functions to quickly generate discrete joint (or conditional) distributions using Dirichlets

**Usage**

```
rprobdist(dim, d, cond, alpha = 1)
```

**Arguments**

|                    |                                                          |
|--------------------|----------------------------------------------------------|
| <code>dim</code>   | the joint dimension of the probability table             |
| <code>d</code>     | number of dimensions                                     |
| <code>cond</code>  | optionally, vertices to condition upon                   |
| <code>alpha</code> | Dirichlet hyper parameter, defaults to 1 (flat density). |

**Details**

`rprobdist` gives an array of dimension `dim` (recycled as necessary to have length `d`, if this is supplied) whose entries are probabilities drawn from a Dirichlet distribution whose parameter vector has entries equal to `alpha` (appropriately recycled).

**Value**

an array of appropriate dimensions

**Side Effects**

Uses as many gamma random variables as cells in the table, so will alter the random seed accordingly.

**Author(s)**

Robin Evans

**Examples**

```
rprobdist(2, 4)      # 2x2x2x2 table
rprobdist(c(2,3,2)) # 2x3x2 table

rprobdist(2, 4, alpha=1/16)  # using unit information prior

# get variables 2 and 4 conditioned upon
rprobdist(2, 4, cond=c(2,4), alpha=1/16)
```

---

 setmatch

*Set Operations*


---

**Description**

Series of functions extending existing vector operations to lists of vectors.

**Usage**

```
setmatch(x, y, nomatch = NA_integer_)
```

**Arguments**

|         |                                                                                              |
|---------|----------------------------------------------------------------------------------------------|
| x       | list of vectors.                                                                             |
| y       | list of vectors.                                                                             |
| nomatch | value to be returned in the case when no match is found. Note that it is coerced to integer. |

**Details**

setmatch checks whether each vector in the list x is also contained in the list y, and if so returns position of the first such vector in y. The ordering of the elements of the vector is irrelevant, as they are considered to be sets.

subsetmatch is similar to setmatch, except vectors in x are searched to see if they are subsets of vectors in y.

setsetdiff is a setwise version of setdiff, and setsetequal a setwise version of setequal.

**Value**

setmatch and subsetmatch return a vector of integers of length the same as the list x.

setsetdiff returns a sublist x.

setsetequal returns a logical of length 1.

**Note**

These functions are not recursive, in the sense that they cannot be used to test lists of lists. They also do not reduce to the vector case.

**Author(s)**

Robin Evans

**See Also**[match](#), [setequal](#), [setdiff](#)**Examples**

```
x = list(1:2, 1:3)
y = list(1:4, 1:3)
setmatch(x, y)
subsetmatch(x, y)
setsetdiff(x, y)

x = list(1:3, 1:2)
y = list(2:1, c(2,1,3))
setsetequal(x, y)
```

---

`subsetMatrix`*Matrix of Subset Indicators*

---

**Description**

Produces a matrix whose rows indicate what subsets of a set are included in which other subsets.

**Usage**

```
subsetMatrix(n)
```

**Arguments**

`n` integer containing the number of elements in the set.

**Details**

This function returns a matrix, with each row and column corresponding to a subset of a hypothetical set of size  $n$ , ordered lexicographically. The entry in row  $i$ , column  $j$  corresponds to whether or not the subset associated with  $i$  is a superset of that associated with  $j$ .

A 1 or -1 indicates that  $i$  is a superset of  $j$ , with the sign referring to the number of fewer elements in  $j$ . 0 indicates that  $i$  is not a superset of  $j$ .

**Value**

An integer matrix of dimension  $2^n$  by  $2^n$ .

**Note**

The inverse of the output matrix is just `abs(subsetMatrix(n))`.

**Author(s)**

Robin Evans

**See Also**

[combinations](#), [powerSet](#), [designMatrix](#).

**Examples**

```
subsetMatrix(3)
```

---

subsetOrder

*Compare sets for inclusion.*

---

**Description**

A wrapper for `is.subset` which returns set inclusions.

**Usage**

```
subsetOrder(x, y)
```

**Arguments**

`x`                    A vector.  
`y`                    A vector of the same type as `x`.

**Details**

If `x` is a subset of `y`, returns `-1`, for the reverse returns `1`. If sets are equal or incomparable, it returns `0`.

**Value**

A single integer, `0`, `-1` or `1`.

**Author(s)**

Robin Evans

**See Also**

[is.subset](#).



**Examples**

```
subsetOrder(2:4, 1:4)
subsetOrder(2:4, 3:5)
```

---

|          |                        |
|----------|------------------------|
| subtable | <i>Subset an array</i> |
|----------|------------------------|

---

**Description**

More flexible calls of `[]` on an array.

**Usage**

```
subtable(x, variables, levels, drop = TRUE)
```

```
subarray(x, levels, drop = TRUE)
```

```
subtable(x, variables, levels) <- value
```

```
subarray(x, levels) <- value
```

**Arguments**

|                        |                                                                                                 |
|------------------------|-------------------------------------------------------------------------------------------------|
| <code>x</code>         | An array.                                                                                       |
| <code>variables</code> | An integer vector containing the dimensions of <code>x</code> to subset.                        |
| <code>levels</code>    | A list or vector containing values to retain.                                                   |
| <code>drop</code>      | Logical indicating whether dimensions with only 1 retained should be dropped. Defaults to TRUE. |
| <code>value</code>     | Value to assign to entries in table.                                                            |

**Details**

Essentially just allows more flexible calls of `[]` on an array.

`subarray` requires the values for each dimension should be specified, so for a  $2 \times 2 \times 2$  array `x`, `subarray(x, list(1, 2, 1:2))` is just `x[1, 2, 1:2]`.

`subtable` allows unspecified dimensions to be retained automatically. Thus, for example `subtable(x, c(2, 3), list(1, 1:2))` is `x[, 1, 1:2]`.

**Value**

Returns an array of dimension `sapply(value, length)` if `drop=TRUE`, otherwise *specified* dimensions of size 1 are dropped. Dimensions which are unspecified in `subtable` are never dropped.

**Functions**

- `subarray`: Flexible subsetting
- `subtable<-`: Assignment in a table
- `subarray<-`: Assignment in an array

**Author(s)**

Mathias Drton, Robin Evans

**See Also**

[Extract](#)

**Examples**

```
x = array(1:8, rep(2,3))
subarray(x, c(2,1,2)) == x[2,1,2]

x[2,1:2,2,drop=FALSE]
subarray(x, list(2,1:2,2), drop=FALSE)

subtable(x, c(2,3), list(1, 1:2))
```

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