

# Package ‘statnet.common’

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**Title** Common R Scripts and Utilities Used by the Statnet Project Software

**Description** Non-statistical utilities used by the software developed by the Statnet Project. They may also be of use to others.

**Imports** utils, methods

**BugReports** <https://github.com/statnet/statnet.common/issues>

**License** GPL-3 + file LICENSE

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all_identical	<i>Test if all items in a vector or a list are identical.</i>
---------------	---

---

**Description**

Test if all items in a vector or a list are identical.

**Usage**

```
all_identical(x)
```

**Arguments**

x                    a vector or a list

**Value**

TRUE if all elements of x are identical to each other.

**See Also**

[base::identical\(\)](#)

**Examples**

```
stopifnot(!all_identical(1:3))
stopifnot(all_identical(list("a", "a", "a")))
```

---

append.rhs.formula	<i>Functions for Querying, Validating and Extracting from ERGM Formulas</i>
--------------------	---

---

**Description**

append.rhs.formula appends a list of terms to the RHS of a formula. If the formula is one-sided, the RHS becomes the LHS, if keep.onesided==FALSE (the default).

**Usage**

```
append.rhs.formula(object, newterms, keep.onesided = FALSE)
nonsimp.update.formula(object, new, ..., from.new = FALSE)
term.list.formula(rhs, sign = +1)
```

**Arguments**

object	formula object to be updated
newterms	list of terms (names) to append to the formula, or a formula whose RHS terms will be used
keep.onesided	if the initial formula is one-sided, keep it whether to keep it one-sided or whether to make the initial formula the new LHS
new	new formula to be used in updating
...	Additional arguments. Currently unused.
from.new	logical or character vector of variable names. controls how environment of formula gets updated.
rhs	a formula-style call containing the right hand side of formula, obtained by <code>fm1a[[3]]</code> for a two-sided formula and <code>fm1a[[2]]</code> for a one-sided formula.
sign	an internal parameter used by <code>term.list.formula</code> when calling itself recursively.

**Value**

`append.rhs.formula` each return an updated formula object

`nonsimp.update.formula` each return an updated formula object

`terms.list.formula` returns a list of formula terms, with an additional numerical vector attribute "sign" with of the same length, giving the corresponding term's sign as +1 or -1.

**Functions**

- `nonsimp.update.formula`: `nonsimp.update.formula` is a reimplementation of [update.formula](#) that does not simplify. Note that the resulting formula's environment is set as follows. If `from.new==FALSE`, it is set to that of `object`. Otherwise, a new sub-environment of `object`, containing, in addition, variables in `new` listed in `from.new` (if a character vector) or all of `new` (if `TRUE`).
- `term.list.formula`: `term.list.formula` returns a list containing terms in a given formula, handling + and - operators and parentheses, and keeping track of whether a term has a plus or a minus sign.

**Examples**

```
## append.rhs.formula

(f1 <- append.rhs.formula(y~x,list(as.name("z1"),as.name("z2"))))
(f2 <- append.rhs.formula(~y,list(as.name("z"))))
(f3 <- append.rhs.formula(~y+x,list(as.name("z"))))
(f4 <- append.rhs.formula(~y,list(as.name("z")),TRUE))
(f5 <- append.rhs.formula(y~x,~z1+z2))
```

---

check.control.class	<i>Check if the class of the control list is one of those that can be used by the calling function</i>
---------------------	--

---

### Description

This function can be called to check that the control list passed is appropriate for the function to be controlled. It does so by looking up the class of the control argument (defaulting to the control variable in the calling function) and checking if it matches a list of acceptable classes.

### Usage

```
check.control.class(OKnames = {      sc <- sys.calls()
  as.character(sc[[length(sc) - 1]][[1]]) }, myname = {      sc <- sys.calls()
  as.character(sc[[length(sc) - 1]][[1]]) }, control = get("control", pos =
  parent.frame()))
```

### Arguments

OKnames	List of control function names which are acceptable.
myname	Name of the calling function (used in the error message).
control	The control list. Defaults to the control variable in the calling function.

### Note

In earlier versions, OKnames and myname were autodetected. This capability has been deprecated and results in a warning issued once per session. They now need to be set explicitly.

### See Also

set.control.class, print.control.list

---

compress.data.frame	<i>"Compress" a data frame.</i>
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---

### Description

compress.data.frame "compresses" a data frame, returning unique rows and a tally of the number of times each row is repeated, as well as a permutation vector that can reconstruct the original data frame. decompress.data.frame reconstructs the original data frame.

### Usage

```
compress.data.frame(x)
```

```
decompress.data.frame(x)
```

**Arguments**

`x` For `compress.data.frame` a [data.frame](#) to be compressed. For `decompress.data.frame` a [list](#) as returned by `compress.data.frame`.

**Value**

For `compress.data.frame`, a [list](#) with three elements:

<code>rows</code>	Unique rows of <code>x</code>
<code>frequencies</code>	A vector of the same length as the number of rows, giving the number of times the corresponding row is repeated
<code>ordering</code>	A vector such that if <code>c</code> is the compressed data frame, <code>c\$rows[c\$ordering, , drop=FALSE]</code> equals the original data frame, except for row names
<code>rownames</code>	Row names of <code>x</code>

For `decompress.data.frame`, the original data frame.

**See Also**

[data.frame](#)

**Examples**

```
(x <- data.frame(V1=sample.int(3,30,replace=TRUE),
                V2=sample.int(2,30,replace=TRUE),
                V3=sample.int(4,30,replace=TRUE)))

(c <- compress.data.frame(x))

stopifnot(all(decompress.data.frame(c)==x))
```

---

compress\_rows

*A generic function to compress a row-weighted table*

---

**Description**

Compress a matrix or a data frame with duplicated rows, updating row weights to reflect frequencies, or reverse the process, reconstructing a matrix like the one compressed (subject to permutation of rows and weights not adding up to an integer).

**Usage**

```
compress_rows(x, ...)
```

```
decompress_rows(x, target.nrows = NULL, ...)
```

**Arguments**

x	a weighted matrix or data frame.
...	extra arguments for methods.
target.nrows	the approximate number of rows the uncompressed matrix should have; if not achievable exactly while respecting proportionality, a matrix with a slightly different number of rows will be constructed.

**Value**

For `compress_rows` A weighted matrix or data frame of the same type with duplicated rows removed and weights updated appropriately.

---

`control.list.accessor` *Named element accessor for ergm control lists*

---

**Description**

Utility method that overrides the standard '\$' list accessor to disable partial matching for ergm `control.list` objects

**Usage**

```
## S3 method for class 'control.list'
object$name
```

**Arguments**

object	list-coerceable object with elements to be searched
name	literal character name of list element to search for and return

**Details**

Executes `getElement` instead of `$` so that element names must match exactly to be returned and partially matching names will not return the wrong object.

**Value**

Returns the named list element exactly matching name, or NULL if no matching elements found

**Author(s)**

Pavel N. Krivitsky

**See Also**

see `getElement`

---

control.remap	<i>Overwrite control parameters of one configuration with another.</i>
---------------	--

---

### Description

Given a `control.list`, and two prefixes, `from` and `to`, overwrite the elements starting with `to` with the corresponding elements starting with `from`.

### Usage

```
control.remap(control, from, to)
```

### Arguments

<code>control</code>	An object of class <code>control.list</code> .
<code>from</code>	Prefix of the source of control parameters.
<code>to</code>	Prefix of the destination of control parameters.

### Value

An `control.list` object.

### Author(s)

Pavel N. Krivitsky

### See Also

[print.control.list](#)

### Examples

```
(l <- set.control.class("test", list(a.x=1, a.y=2)))
control.remap(l, "a", "b")
```

---

ERRVL	<i>Return the first argument passed (out of any number) that is not a try-error (result of <a href="#">try</a> encountering an error).</i>
-------	--

---

### Description

This function is inspired by [NVL](#), and simply returns the first argument that is not a try-error, raising an error if all arguments are try-errors.

### Usage

```
ERRVL(...)
```

**Arguments**

... Expressions to be tested; usually outputs of `try`.

**Value**

The first argument that is not a try-error. Stops with an error if all are.

**See Also**

`try`, `inherits`

**Examples**

```
print(ERRVL(1,2,3)) # 1
print(ERRVL(try(solve(0)),2,3)) # 2
```

---

logspace.utils

*Utilities for performing calculations on logarithmic scale.*

---

**Description**

A small suite of functions to compute sums, means, and weighted means on logarithmic scale, minimizing loss of precision.

**Usage**

```
log_sum_exp(logx, use_ldouble = FALSE)
```

```
log_mean_exp(logx, use_ldouble = FALSE)
```

```
lweighted.mean(x, logw)
```

```
lweighted.var(x, logw)
```

**Arguments**

`logx` Numeric vector of  $\log(x)$ , the natural logarithms of the values to be summed or averaged.

`use_ldouble` Whether to use long double precision in the calculation. If TRUE, 's C built-in `logspace_sum()` is used. If FALSE, the package's own implementation based on it is used, using double precision, which is (on most systems) several times faster, at the cost of precision.

`x` Numeric vector of  $x$ , the (raw) values to be summed or averaged. For `lweighted.mean`, `x` may also be a matrix, in which case the weighted mean will be computed for each column of `x`.

`logw` Numeric vector of  $\log(w)$ , the natural logarithms of the weights.



**Value**

The functions return the equivalents of the following R expressions, but faster and with less loss of precision:

```
log_sum_exp(logx) log(sum(exp(logx)))
```

```
log_mean_exp(logx) log(mean(exp(logx)))
```

```
lweighted.mean(x, logw) sum(x*exp(logw))/sum(exp(logw)) for x scalar and colSums(x*exp(logw))/sum(exp(logw))
for x matrix
```

```
lweighted.var(x, logw) crossprod(x*exp(logw/2))/sum(exp(logw))
```

**Author(s)**

Pavel N. Krivitsky

**Examples**

```
logx <- rnorm(1000)
stopifnot(all.equal(log(sum(exp(logx))), log_sum_exp(logx)))
stopifnot(all.equal(log(mean(exp(logx))), log_mean_exp(logx)))

x <- rnorm(1000)
logw <- rnorm(1000)
stopifnot(all.equal(m <- sum(x*exp(logw))/sum(exp(logw)), lweighted.mean(x, logw)))
stopifnot(all.equal(sum((x-m)^2*exp(logw))/sum(exp(logw)),
                      lweighted.var(x, logw), check.attributes=FALSE))

x <- cbind(x, rnorm(1000))
stopifnot(all.equal(m <- colSums(x*exp(logw))/sum(exp(logw)),
                    lweighted.mean(x, logw), check.attributes=FALSE))
stopifnot(all.equal(crossprod(t(t(x)-m)*exp(logw/2))/sum(exp(logw)),
                    lweighted.var(x, logw), check.attributes=FALSE))
```

---

NVL

*Convenience functions for handling [NULL](#) objects.*

---

**Description**

Convenience functions for handling [NULL](#) objects.

**Usage**

```
NVL(...)
```

```
NVL(x) <- value
```

**Arguments**

...	Expressions to be tested.
x	an object to be overwritten if <a href="#">NULL</a> .
value	new value for x.

## Functions

- `NVL`: Inspired by SQL function `NVL`, `NVL(...)` returns the first argument that is not `NULL`, or `NULL` if all arguments are `NULL`.
- `NVL<-`: Assigning to `NVL` overwrites its first argument if that argument is `NULL`. Note that it will *always* return the right-hand-side of the assignment (`value`), regardless of what `x` is.

## See Also

`NULL`, `is.null`, `if`

## Examples

```
a <- NULL

print(a) # NULL
print(NVL(a,0)) # 0

b <- 1

print(b) # 1
print(NVL(b,0)) # 1

# Also,
print(NVL(NULL,1,0)) # 1
print(NVL(NULL,0,1)) # 0
print(NVL(NULL,NULL,0)) # 0
print(NVL(NULL,NULL,NULL)) # NULL

NVL(a) <- 2
a # 2
NVL(b) <- 2
b # still 1
```

---

opttest

*Optionally test code depending on environment variable.*

---

## Description

A convenience wrapper to run code based on whether an environment variable is defined.

## Usage

```
opttest(expr, testname = NULL, testvar = "ENABLE_statnet_TESTS",
  yesvals = c("y", "yes", "t", "true", "1"), lowercase = TRUE)
```

## Arguments

<code>expr</code>	An expression to be evaluated only if <code>testvar</code> is set to a non-empty value.
<code>testname</code>	Optional name of the test. If given, and the test is skipped, will print a message to that end, including the name of the test, and instructions on how to enable it.

testvar	Environment variable name. If set to one of the yesvals, expr is run. Otherwise, an optional message is printed.
yesvals	A character vector of strings considered affirmative values for testvar.
lowercase	Whether to convert the value of testvar to lower case before comparing it to yesvals.

---

order	<i>Implement the <a href="#">sort</a> and <a href="#">order</a> methods for <code>data.frame</code> and <code>matrix</code>, sorting it in lexicographic order.</i>
-------	---

---

### Description

These function return a data frame sorted in lexicographic order or a permutation that will rearrange it into lexicographic order: first by the first column, ties broken by the second, remaining ties by the third, etc..

### Usage

```
order(..., na.last = TRUE, decreasing = FALSE)

## Default S3 method:
order(..., na.last = TRUE, decreasing = FALSE)

## S3 method for class 'data.frame'
order(..., na.last = TRUE, decreasing = FALSE)

## S3 method for class 'matrix'
order(..., na.last = TRUE, decreasing = FALSE)

## S3 method for class 'data.frame'
sort(x, decreasing = FALSE, ...)
```

### Arguments

...	Ignored for sort. For order, first argument is the data frame to be ordered. (This is needed for compatibility with <a href="#">order</a> .)
na.last	See <a href="#">order</a> documentation.
decreasing	Whether to sort in decreasing order.
x	A <a href="#">data.frame</a> to sort.

### Value

For sort, a data frame, sorted lexicographically. For order, a permutation I (of a vector 1:nrow(x)) such that x[I, , drop=FALSE] equals x ordered lexicographically.

### See Also

[data.frame](#), [sort](#), [order](#), [matrix](#)

**Examples**

```
data(iris)
head(iris)
head(order(iris))
head(sort(iris))
stopifnot(identical(sort(iris),iris[order(iris),]))
```

---

paste.and

*Concatenates the elements of a vector (optionally enclosing them in quotation marks or parentheses) adding appropriate punctuation and unions.*

---

**Description**

A vector `x` becomes "`x[1]`", "`x[1]` and `x[2]`", or "`x[1]`, `x[2]`, and `x[3]`", depending on the length of `x`.

**Usage**

```
paste.and(x, oq = "", cq = "")
```

**Arguments**

<code>x</code>	A vector.
<code>oq</code>	Opening quotation symbol. (Defaults to none.)
<code>cq</code>	Closing quotation symbol. (Defaults to none.)

**Value**

A string with the output.

**See Also**

paste, cat

**Examples**

```
print(paste.and(c()))
print(paste.and(1))
print(paste.and(1:2))
print(paste.and(1:3))
print(paste.and(1:4))
```

---

print.control.list      *Pretty print the control list*

---

### Description

This function prints the control list, including what it can control and the elements.

### Usage

```
## S3 method for class 'control.list'  
print(x, ...)
```

### Arguments

x                      A list generated by a control.\* function.  
...                    Unused at this time.

### See Also

[check.control.class](#), [set.control.class](#)

---

rle.utils                *RLE utilities*

---

### Description

Simple utilities for operations on RLE-encoded vectors.

### Usage

```
## S3 method for class 'rle'  
c(...)  
  
## S3 method for class 'rle'  
!x  
  
binop.rle(e1, e2, FUN)  
  
## S3 method for class 'rle'  
e1 | e2  
  
## S3 method for class 'rle'  
e1 & e2  
  
compact.rle(x)  
  
## S3 method for class 'rle'  
any(..., na.rm = FALSE)
```

```

## S3 method for class 'rle'
all(..., na.rm = FALSE)

## S3 method for class 'rle'
e1 * e2

## S3 method for class 'rle'
e1 / e2

## S3 method for class 'rle'
e1 - e2

## S3 method for class 'rle'
e1 + e2

## S3 method for class 'rle'
e1 ^ e2

## S3 method for class 'rle'
e1 %% e2

## S3 method for class 'rle'
e1 %/% e2

## S3 method for class 'rle'
e1 == e2

## S3 method for class 'rle'
e1 > e2

## S3 method for class 'rle'
e1 < e2

## S3 method for class 'rle'
e1 != e2

## S3 method for class 'rle'
e1 <= e2

## S3 method for class 'rle'
e1 >= e2

## S3 method for class 'rle'
rep(x, ..., scale = c("element", "run"))

```

### Arguments

...	For c, objects to be concatenated. The first object must be of class <code>rle()</code> . For rep, see documentation for <code>rep()</code> .
x, e1, e2	Arguments to unary (x) and binary (e1 and e2) operators.
FUN	A binary function or operator or a name of one. It is assumed to be vectorized: it expects two vectors of equal lengths and outputs a vector of the same length.

na.rm            see documentation for `any()` and `all()`.  
 scale            whether to replicate the elements of the RLE-compressed vector or the runs.

### Value

All functions return an `rle()` object. By default, the functions and the operators do not merge adjacent runs with the same value. This must be done explicitly with `compact.rle()`.

### Functions

- `binop.rle`: Perform an arbitrary binary operation on the pair of vectors represented by the `rle()` objects.
- `compact.rle`: Compact the `rle()` object by merging adjacent runs.

### Note

Since `rle()` stores run lengths as integers, `compact.rle()` will not merge runs that add up to lengths greater than what can be represented by a 32-bit signed integer (2147483647).

The `rep()` method for `rle()` objects is very limited at this time: . Even though the default setting is to replicate elements of the vector, only the run-replicating functionality is implemented at this time.

### Examples

```
x <- rle(as.logical(rbinom(10,1,.7)))
y <- rle(as.logical(rbinom(10,1,.3)))

stopifnot(c(inverse.rle(x),inverse.rle(y))==inverse.rle(c(x,y)))

stopifnot(!inverse.rle(x)==inverse.rle(!x))
stopifnot((inverse.rle(x)|inverse.rle(y))==inverse.rle(x|y))
stopifnot((inverse.rle(x)&inverse.rle(y))==inverse.rle(x&y))
stopifnot(identical(rle(inverse.rle(x)&inverse.rle(y)),compact.rle(x&y)))

big <- structure(list(lengths=as.integer(rep(.Machine$integer.max/4,6)),
                      values=rep(TRUE,6)), class="rle")

stopifnot(all(aggregate(as.numeric(lengths)~values,
                        data=as.data.frame(unclass(big)),FUN=sum)
            ==
            aggregate(as.numeric(lengths)~values,
                        data=as.data.frame(unclass(compact.rle(big))),
                        FUN=sum)))

x <- rle(as.logical(rbinom(10,1,.9)))
y <- rle(as.logical(rbinom(10,1,.1)))

stopifnot(any(x)==any(inverse.rle(x)))
stopifnot(any(y)==any(inverse.rle(y)))

stopifnot(all(x)==all(inverse.rle(x)))
stopifnot(all(y)==all(inverse.rle(y)))
```

```

x <- rle(sample(c(-1,+1), 10, c(.7,.3), replace=TRUE))
y <- rle(sample(c(-1,+1), 10, c(.3,.7), replace=TRUE))

stopifnot((inverse.rle(x)*inverse.rle(y))==inverse.rle(x*y))
stopifnot((inverse.rle(x)/inverse.rle(y))==inverse.rle(x/y))
stopifnot((-inverse.rle(y))==inverse.rle(-y))
stopifnot((inverse.rle(x)-inverse.rle(y))==inverse.rle(x-y))
stopifnot((+inverse.rle(y))==inverse.rle(+y))
stopifnot((inverse.rle(x)+inverse.rle(y))==inverse.rle(x+y))
stopifnot((inverse.rle(x)^inverse.rle(y))==inverse.rle(x^y))
stopifnot((inverse.rle(x)%inverse.rle(y))==inverse.rle(x%y))
stopifnot((inverse.rle(x)/inverse.rle(y))==inverse.rle(x%/y))
stopifnot((inverse.rle(x)==inverse.rle(y))==inverse.rle(x==y))
stopifnot((inverse.rle(x)>inverse.rle(y))==inverse.rle(x>y))
stopifnot((inverse.rle(x)<inverse.rle(y))==inverse.rle(x<y))
stopifnot((inverse.rle(x)!=inverse.rle(y))==inverse.rle(x!=y))
stopifnot((inverse.rle(x)<=inverse.rle(y))==inverse.rle(x<=y))
stopifnot((inverse.rle(x)>=inverse.rle(y))==inverse.rle(x>=y))

x <- rle(sample(c(-1,+1), 10, c(.7,.3), replace=TRUE))
y <- rpois(length(x$lengths), 2)

stopifnot(all(rep(inverse.rle(x), rep(y, x$lengths))==inverse.rle(rep(x, y, scale="run"))))

```

---

set.control.class      *Set the class of the control list*

---

## Description

This function sets the class of the control list, with the default being the name of the calling function.

## Usage

```

set.control.class(myname = {      sc <- sys.calls()
  as.character(sc[[length(sc) - 1]][[1]]) }, control = get("control", pos =
  parent.frame()))

```

## Arguments

myname	Name of the class to set.
control	Control list. Defaults to the control variable in the calling function.

## Value

The control list with class set.

## Note

In earlier versions, OKnames and myname were autodetected. This capability has been deprecated and results in a warning issued once per session. They now need to be set explicitly.



**See Also**

check.control.class, print.control.list

---

statnet.cite

CITATION *file utilities for Statnet packages*

---

**Description**

These functions automate citation generation for Statnet Project packages.

**Usage**

```
statnet.cite.head(pkg)
```

```
statnet.cite.foot(pkg)
```

```
statnet.cite.pkg(pkg)
```

**Arguments**

pkg                    Name of the package whose citation is being generated.

**Value**

For `statnet.cite.head` and `statnet.cite.foot`, an object of type `citationHeader` and `citationFooter`, respectively, understood by the `citation` function, with package name substituted into the template.

For `statnet.cite.pkg`, an object of class `bibentry` containing a 'software manual' citation for the package constructed from the current version and author information in the `DESCRIPTION` and a template.

**See Also**

`citation`, `citHeader`, `citFooter`, `bibentry`

**Examples**

```
statnet.cite.head("statnet.common")
```

```
statnet.cite.pkg("statnet.common")
```

```
statnet.cite.foot("statnet.common")
```

---

statnetStartupMessage *Construct a "standard" startup message to be printed when the package is loaded.*

---

### Description

This function uses information returned by [packageDescription](#) to construct a standard package startup message according to the policy of the Statnet Project. To determine institutional affiliation, it uses a lookup table that maps domain names to institutions. (E.g., \*.uw.edu or \*.washington.edu maps to University of Washington.)

### Usage

```
statnetStartupMessage(pkgname, friends, nofriends)
```

### Arguments

pkgname	Name of the package whose information is used.
friends	<p>This argument is required, but will only be interpreted if the Statnet Project policy makes use of "friendly" package information.</p> <p>A character vector of names of packages whose attribution information incorporates the attribution information of this package, or TRUE. (This may, in the future, lead the package to suppress its own startup message when loaded by a "friendly" package.)</p> <p>If TRUE, the package considers all other packages "friendly". (This may, in the future, lead the package to suppress its own startup message when loaded by another package, but print it when loaded directly by the user.)</p>
nofriends	<p>This argument controls the startup message if the Statnet Project policy does not make use of "friendly" package information but does make use of whether or not the package is being loaded directly or as a dependency.</p> <p>If TRUE, the package is willing to suppress its startup message if loaded as a dependency. If FALSE, it is not.</p>

### Value

A string containing the startup message, to be passed to the [packageStartupMessage](#) call or NULL, if policy prescribes printing 's default startup message. (Thus, if `statnetStartupMessage` returns NULL, the calling package should not call [packageStartupMessage](#) at all.)

Note that arguments to `friends` and `nofriends` are merely requests, to be interpreted (or ignored) by the `statnetStartupMessage` according to the Statnet Project policy.

### See Also

[packageDescription](#)

**Examples**

```
## Not run:
.onAttach <- function(lib, pkg){
  sm <- statnetStartupMessage("ergm", friends=c("statnet", "ergm.count", "tergm"), nofriends=FALSE)
  if(!is.null(sm)) packageStartupMessage(sm)
}

## End(Not run)
```

---

sweep_cols.matrix	<i>Subtract a elements of a vector from respective columns of a matrix</i>
-------------------	--

---

**Description**

An optimized function equivalent to `sweep(x, 2, STATS)` for a matrix `x`.

**Usage**

```
sweep_cols.matrix(x, STATS, disable_checks = FALSE)
```

**Arguments**

`x` a numeric matrix;

`STATS` a numeric vector whose length equals to the number of columns of `x`.

`disable_checks` if TRUE, do not check that `x` is a numeric matrix and its number of columns matches the length of `STATS`; set in production code for a significant speed-up.

**Value**

A matrix of the same attributes as `x`.

**Examples**

```
x <- matrix(runif(1000), ncol=4)
s <- 1:4

stopifnot(all.equal(sweep_cols.matrix(x, s), sweep(x, 2, s)))
```

---

<code>vector.namesmatch</code>	<i>reorder vector v into order determined by matching the names of its elements to a vector of names</i>
--------------------------------	--

---

### Description

A helper function to reorder vector `v` (if named) into order specified by matching its names to the argument names

### Usage

```
vector.namesmatch(v, names, errname = NULL)
```

### Arguments

<code>v</code>	a vector (or list) with named elements, to be reordered
<code>names</code>	a character vector of element names, corresponding to names of <code>v</code> , specifying desired ordering of <code>v</code>
<code>errname</code>	optional, name to be reported in any error messages. default to <code>deparse(substitute(v))</code>

### Details

does some checking of appropriateness of arguments, and reorders `v` by matching its names to character vector names

### Value

returns `v`, with elements reordered

### Note

earlier versions of this function did not order as advertised

### Examples

```
test<-list(c=1,b=2,a=3)
vector.namesmatch(test,names=c('a','c','b'))
```

---

<code>wmatrix</code>	<i>A data matrix with row weights</i>
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---

### Description

A representation of a numeric matrix with row weights, represented on either linear (`linwmatrix`) or logarithmic (`logwmatrix`) scale.

**Usage**

```
logwmatrix(data = NA, nrow = 1, ncol = 1, byrow = FALSE,
            dimnames = NULL, w = NULL)

linwmatrix(data = NA, nrow = 1, ncol = 1, byrow = FALSE,
            dimnames = NULL, w = NULL)

is.wmatrix(x)

is.logwmatrix(x)

is.linwmatrix(x)

as.linwmatrix(x, ...)

as.logwmatrix(x, ...)

## S3 method for class 'linwmatrix'
as.linwmatrix(x, ...)

## S3 method for class 'logwmatrix'
as.linwmatrix(x, ...)

## S3 method for class 'logwmatrix'
as.logwmatrix(x, ...)

## S3 method for class 'linwmatrix'
as.logwmatrix(x, ...)

## S3 method for class 'matrix'
as.linwmatrix(x, w = NULL, ...)

## S3 method for class 'matrix'
as.logwmatrix(x, w = NULL, ...)

## S3 method for class 'wmatrix'
print(x, ...)

## S3 method for class 'logwmatrix'
print(x, ...)

## S3 method for class 'linwmatrix'
print(x, ...)

## S3 method for class 'logwmatrix'
compress_rows(x, ...)

## S3 method for class 'linwmatrix'
compress_rows(x, ...)

## S3 method for class 'wmatrix'
decompress_rows(x, target.nrows = NULL, ...)
```

```
## S3 method for class 'wmatrix'
x[i, j, ..., drop = FALSE]

## S3 replacement method for class 'wmatrix'
x[i, j, ...] <- value
```

### Arguments

data, nrow, ncol, byrow, dimnames  
passed to `matrix()`.

w  
row weights on the appropriate scale.

x  
an object to be coerced or tested.

...  
extra arguments, currently unused.

target.nrows  
see `decompress_rows()`.

i, j, value  
rows and columns and values for extraction or replacement; as `matrix()`.

drop  
Used for consistency with the generic. Ignored, and always treated as FALSE.

### Value

An object of class `linwmatrix`/`logwmatrix` and `wmatrix`, which is a `matrix()` but also has an attribute `w` containing row weights on the linear or the natural-log-transformed scale.

### Note

Note that `wmatrix` itself is an "abstract" class: you cannot instantiate it.

Note that at this time, `wmatrix` is designed as, first and foremost, as class for storing compressed data matrices, so most methods that operate on matrices may not handle the weights correctly and may even cause them to be lost.

### See Also

`rowweights()`, `lrowweights()`, `compress_rows()`

### Examples

```
(m <- matrix(1:3, 2, 3, byrow=TRUE))
(m <- rbind(m, 3*m, 2*m, m))
(mlog <- as.logwmatrix(m))
(mlin <- as.linwmatrix(m))
(cmlog <- compress_rows(mlog))
(cmlin <- compress_rows(mlin))

stopifnot(all.equal(as.linwmatrix(cmlog), cmlin))

cmlog[2,] <- 1:3
(cmlog <- compress_rows(cmlog))
stopifnot(sum(rowweights(cmlog))==nrow(m))

(m3 <- matrix(c(1:3, (1:3)*2, (1:3)*3), 3, 3, byrow=TRUE))
(rowweights(m3) <- c(4, 2, 2))

stopifnot(all.equal(compress_rows(as.logwmatrix(m)), as.logwmatrix(m3), check.attributes=FALSE))
```

```
stopifnot(all.equal(rowweights(compress_rows(as.logwmatrix(m))),
                    rowweights(as.logwmatrix(m3),check.attributes=FALSE))
```

---

wmatrix\_weights      *Set or extract weighted matrix row weights*

---

## Description

Set or extract weighted matrix row weights

## Usage

```
rowweights(x, ...)

## S3 method for class 'linwmatrix'
rowweights(x, ...)

## S3 method for class 'logwmatrix'
rowweights(x, ...)

lrowweights(x, ...)

## S3 method for class 'logwmatrix'
lrowweights(x, ...)

## S3 method for class 'linwmatrix'
lrowweights(x, ...)

rowweights(x, ...) <- value

## S3 replacement method for class 'linwmatrix'
rowweights(x, update = TRUE, ...) <- value

## S3 replacement method for class 'logwmatrix'
rowweights(x, update = TRUE, ...) <- value

lrowweights(x, ...) <- value

## S3 replacement method for class 'linwmatrix'
lrowweights(x, update = TRUE, ...) <- value

## S3 replacement method for class 'logwmatrix'
lrowweights(x, update = TRUE, ...) <- value

## S3 replacement method for class 'matrix'
rowweights(x, ...) <- value

## S3 replacement method for class 'matrix'
lrowweights(x, ...) <- value
```

**Arguments**

x	a <code>linwmatrix()</code> , a <code>logwmatrix()</code> , or a <code>matrix()</code> ; a <code>matrix()</code> is coerced to a weighted matrix of an appropriate type.
...	extra arguments for methods.
value	weights to set, on the appropriate scale.
update	if TRUE (the default), the old weights are updated with the new weights (i.e., corresponding weights are multiplied on linear scale or added on on log scale); otherwise, they are overwritten.

**Value**

For the accessor functions, the row weights or the row log-weights; otherwise, a weighted matrix with modified weights. The type of weight (linear or logarithmic) is converted to the required type and the type of weighting of the matrix is preserved.



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