

Package ‘miscset’

February 24, 2017

Type Package

Title Miscellaneous Tools Set

Version 1.1.0

Date 2017-02-24

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Description A collection of miscellaneous methods to simplify various tasks, including plotting, data.frame and matrix transformations, environment functions, regular expression methods, and string and logical operations, as well as numerical and statistical tools. Most of the methods are simple but useful wrappers of common base R functions, which extend S3 generics or provide default values for important parameters.

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Imports data.table, devtools, ggplot2, gridExtra, parallel, Rcpp (>= 0.11.1), stats, xtable

LinkingTo Rcpp

Suggests knitr, rmarkdown, stringr

VignetteBuilder knitr

URL <https://github.com/setempler/miscset>

BugReports <https://github.com/setempler/miscset/issues>

RoxygenNote 6.0.1

NeedsCompilation yes

Repository CRAN

Date/Publication 2017-02-24 16:46:57

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miscset-package	<i>Miscellaneous R Tools</i>
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Description

A collection of miscellaneous methods to simplify various tasks, including plotting, data.frame and matrix transformations, environment functions, regular expression methods, and string and logical operations, as well as numerical and statistical tools. Most of the methods are simple but useful wrappers of common base R functions, which extend S3 generics or provide default values for important parameters.

Details

The package vignette provides a comprehensive overview and examples for the usage of all available functions in the package. View with `vignette("miscset")`.

Author(s)

Sven E. Templer

ciplot

Barplot with Confidence Intervals

Description

Create barplots of a list of numeric values and error bars according to the confidence interval, standard deviation, interquartile range, etc.

Usage

```
ciplot(x, ...)

## Default S3 method:
ciplot(x, ..., ylim, height.fun = mean,
       height.args = list(), error.fun = confint, error.args = list(),
       arrows.args = list(code = 3, angle = 90), na.rm = TRUE)
```

Arguments

x	List of numeric values
...	Arguments forwarded to barplot in default method.
ylim	A range for the y-axis limits.
height.fun	Function to apply on each list object to calculate the height of the bars from.
height.args	Arguments forwarded to height.fun, as a named list.
error.fun	Function to calculate the error size. See also details.
error.args	Arguments forwarded to error.fun, as a named list.
arrows.args	Arguments forwarded to arrows, as a named list.
na.rm	Logical, remove missing values.

Details

Example for quantiles:

```
interquartile <- function(x) {quartile(x,.75)-mean(x)}
quantileQ <- function(x, q) {abs(quartile(x,q[1])-mean(x))}
```

Author(s)

Sven E. Templer

collapse*Collapse objects***Description**

Collapse objects as in the `paste` function.

Usage

```
collapse(x, sep, ...)

## Default S3 method:
collapse(x, sep = "", ..., .unique = FALSE,
        .sort = FALSE, .decreasing = FALSE)

## S3 method for class 'data.frame'
collapse(x, sep = "", by = names(x), ...,
         .unique = FALSE, .sort = FALSE, .decreasing = FALSE, .unlist = FALSE,
         .sortby = FALSE)
```

Arguments

<code>x</code>	Any R object.
<code>sep</code>	A character string to separate value columns. <code>NULL</code> retains a vector.
<code>...</code>	Forwarded to or from other methods.
<code>.unique</code>	Logical, return only unique values.
<code>.sort</code>	Logical, sort the values.
<code>.decreasing</code>	Logical, if sorting, then by decreasing values.
<code>by</code>	Column names to split data frame by, before applying collapse on each remaining column within each piece. Using the default (all columns), then <code>unique(x)</code> is returned. Columns can be specified by names or integer with the column numbers. Using <code>0</code> or <code>NULL</code> collapses all columns.
<code>.unlist</code>	Logical, if value columns need to be unlisted before collapsing.
<code>.sortby</code>	Logical, sort the output on the <code>by</code> columns. This applies , If <code>x</code> was a <code>data.table</code> , then the keys are set as the <code>by</code> values.

Details

For the `data.frame` method, `x` is converted to a `data.table` before applying the piece- and column-wise collapses. If the input is already inheriting from `data.table`, then the class is retained. `.sortby` is causing `setkeyv(x, by)` to be applied to `x` after converting to a `data.table`.

Author(s)

Sven E. Templer

Examples

```

#
## some data

set.seed(12)
s <- s2 <- sample(LETTERS[1:4], 9, replace = TRUE)
s2[1:2] <- rev(s2[1:2])
d <- data.frame(group = rep(letters[c(3,1,2)], each = 3),
                 value = s,
                 level = factor(s2),
                 stringsAsFactors = FALSE)

## collapse vectors

collapse(letters)
collapse(1:3)           # coerced to character
collapse(LETTERS[1:5], "-") # separated by '-'

## collapse data.frames

# by all columns (same as unique)
collapse(d)
# by a grouping column
collapse(d, by = 1)
# by multiple, but not all columns
collapse(d, by = c("group", "value"))
# return single row
collapse(d, by = 0)
# return single row, unique and sorted values
collapse(d, by = 0, .unique = TRUE, .sort = TRUE)

#

```

Description

Calculate confidence intervals for values of a numeric vector.

Usage

```

## S3 method for class 'numeric'
confint(object, parm = qnorm, level = 0.95, ...,
        na.rm = TRUE, ret.attr = TRUE)

```

Arguments

<code>object</code>	A numeric vector.
<code>parm</code>	Function for quantile calculation. e.g. <code>qnorm</code> , <code>qt</code>
<code>level</code>	Size of confidence ($0 < \text{size} < 1$).
<code>...</code>	Unused.
<code>na.rm</code>	Logical, remove missing values for <code>sd</code> and <code>mean</code> .
<code>ret.attr</code>	Logical, to include the mean value and function arguments as attributes of the returned object.

Value

Returns a numeric vector with the lower and upper range of the confidence interval.

Author(s)

Sven E. Templer

Examples

```
#  
confint(1:3)  
confint(1:3, ret.attr = FALSE)  
#
```

do.rbind

Bind data.frames in a List by Rows

Description

Same as `do.call(rbind, x)`, but adding a column with the name of each table. Missing names are replaced by integers.

Usage

```
do.rbind(x, idcol = "Name", keep.rownames = FALSE)
```

Arguments

<code>x</code>	List with data.frames. Non data.frame objects are dropped.
<code>idcol</code>	Name for column with ids in output.
<code>keep.rownames</code>	Logical, keep rownames.

Value

Returns a data.frame

Author(s)

Sven E. Templer

duplicates *Determine Duplicates*

Description

Determine duplicates. duplicates returns a logical vector, duplicatei an integer vector.

Usage

```
duplicates(x)  
duplicatei(x, first = TRUE)
```

Arguments

x	A vector or data.frame to search for duplicates.
first	Logical, TRUE to return the index also for the first occurrence of values. Otherwise, a 0 is the index for the first occurrence.

Value

duplicates returns a logical vector as [duplicated](#), but with TRUE values also for the first occurrence of duplicated values.

duplicatei returns the index of the first occurrence of each unique value.

Author(s)

Sven E. Templer

Examples

```
#  
  
x <- c(7, 7, 7, 2, 3, 2)  
data.frame(  
  data = x,  
  duplicated = duplicated(x),  
  duplicates = duplicates(x),  
  duplicatei = duplicatei(x),  
  duplicatei0 = duplicatei(x, FALSE))  
  
#
```

enpaire*Create a Pairwise List from a Matrix*

Description

Transform a `matrix` or `dist` object to a pairwise list.

Usage

```
enpaire(x, ...)

## Default S3 method:
enpaire(x, ...)

## S3 method for class 'dist'
enpaire(x, upper = T, lower = T, ...)

## S3 method for class 'matrix'
enpaire(x, upper = T, lower = T, ...)
```

Arguments

<code>x</code>	Object of class <code>matrix</code> .
<code>...</code>	Arguments passed to methods.
<code>upper</code>	Logical, return values from upper triangle.
<code>lower</code>	Logical, return values from lower triangle.

Value

Returns a `data.frame`. The first and second column represent the dimension names for a value in `x`. The following columns contain the values for the upper or lower triangle.

Author(s)

Sven E. Templer

See Also

[squarematrix](#)

Examples

```
# 

m <- matrix(letters[1:9], 3, 3, dimnames = list(1:3,1:3))
enpaire(m)
enpaire(m, lower = FALSE)
```

#

factorNA*Create a Factor with NA as Level*

Description

Create a [factor](#) with NA values included and positioned as last level.

Usage

```
factorNA(x, ...)
```

Arguments

x	A vector coerced to character.
...	Forwarded to factor . x and levels are defined.

Author(s)

Sven E. Templer

gghcl*HTML Colours Like ggplot2*

Description

Calculate HTML colour code from a palette like ggplot2 uses.

Usage

```
gghcl(n, sub = 1:n, h = c(0, 360) + 15, c = 100, l = 65, ...)
```

Arguments

n	Numeric value to determine size of palette.
sub	Numeric vector with values within range from 1 to n to subset palette.
h	Hue of the colour. Within range of a circle's degrees.
c	Chroma of the colour.
l	Luminance of the colour. Within range from 1 to 100.
...	Further arguments passed to function hcl.

Details

See [?hcl](#) for explanation of h, c and l.

Value

Returns a character vector containing HTML colour code of the standard ggplot colour palette.

Author(s)

Sven E. Templer

See Also

[hcl](#)

Examples

```
#  
  
# Plot some palettes:  
par(mfrow = c(3,1), mai = c(.1,.1,1,.1))  
p <- matrix(1:10, 10)  
image(p, col = gghcl(5), axes = FALSE, main ="gghcl(5)")  
image(p, col = gghcl(10), axes = FALSE, main = "gghcl(10)")  
image(p, col = gghcl(10, 1:5), axes = FALSE, main ="gghcl(10, 1:5)")  
# dev.off() # to reset \code{par}  
  
#
```

ggplotGrid

Arrange a List of ggplots

Description

Arrange a list of ggplots with [grid.arrange](#) and output on local graphic device or as pdf/png when a path is supplied. `ggplotGridA4` writes the plots to a DIN A4 (8 x 11 inches) pdf file directly.

Usage

```
ggplotGrid(l, path, ncol = 1, nrow = 1, width = 8, height = 11,  
          res = 300, pdf.cairo = TRUE, onefile = TRUE, ...)  
  
ggplotGridA4(l, path, ncol = 2, nrow = 1, wide = TRUE)  
  
ggplotlist(x, ncol = 1, path, width = 11, height = 8)
```

Arguments

l	List with ggplot objects.
path	Plot to file of type pdf or png. Determine type by path ending (.pdf or .png). Optional in ggplotlist: A character string that gives the path to export the plot to a file, ending with 'pdf' or 'png' (case insensitive). If missing, then the grid is returned to the current graphic device.
ncol	Number of columns.
nrow	Number of rows per page, only for pdfs.
width	For pdfs/pngs the width in inches, else ignored.
height	For pdfs/pngs the height in inches, else ignored.
res	Resolution in dpi for pngs.
pdf.cairo	Use cairo_pdf (or cairo_ps, svg) instead of pdf
onefile	Create one file, see cairo_pdf .
...	Forwarded to cairo_pdf
wide	Wide format pdf pages (11x8 inches).
x	A list containing at least one ggplot object of class gg.

Author(s)

Sven E. Templer

Examples

```
# 

## Not run:
library(ggplot2)
d <- data.frame(a=1:5,b=1:5)
x <- list(
  ggplot(d, aes(x=a,y=b,col=b)) + geom_line(),
  ggplot(d, aes(x=a,y=b,shape=factor(b))) + geom_point())
ggplotlist(x, 2)
## End(Not run)

#
```

Description

Function to extract a certain index from gregexpr().

Usage

```
gregexprind(pattern, text, n, ...)
```

Arguments

<code>pattern</code>	Character string containing a regular expression to be searched in <code>text</code> .
<code>text</code>	Character vector where the search is performed.
<code>n</code>	Numeric value or character string "last" to extract nth or last position of <code>pattern</code> in each value of <code>text</code> .
...	Arguments passed to function <code>gregexpr()</code> .

Value

Numeric vector of length `length(text)`.

Author(s)

Sven E. Templer

See Also

See [gregexpr](#) for further information on arguments.

See [regex](#) for the use of regular expressions.

Examples

```
#  
gregexprind(c("a"),c("ababa","ab","xyz",NA), 1)  
gregexprind(c("a"),c("ababa","ab","xyz",NA), 2)  
gregexprind(c("a"),c("ababa","ab","xyz",NA), "last")  
#
```

Description

Given a package name or string, start the package help index page in a browser.

Usage

```
help.index(pkg, browser = NULL)
```

Arguments

pkg	A character string or expression with the name of a package.
browser	The browser to display. text and pdf don't use a browser, but builtin text/pdf (help_type). Otherwise a character string for the browser program binary to call or function.

Author(s)

Sven E. Templer

info

Print enhanced session information

Description

Based on and enhancing devtools::session_info.

Usage

```
info(..., width = 120)
```

Arguments

...	Forwarded to other methods.
width	Console width in columns.

Author(s)

Sven E. Templer

See Also

[session_info](#)

Examples

```
info()  
devtools::session_info()  
sessionInfo()
```

leading0*Numeric to Character with Leading Zero(s)***Description**

Transform numeric values to character string prepending leading zero(s).

Usage

```
leading0(num, digits = 2)
```

Arguments

<code>num</code>	Numeric vector (character also possible) to transform.
<code>digits</code>	Numeric value of minimum length of output strings.

Value

Character vector with same length of strings of each value. Original "string" is prepended by zero(s). String length is at least `max(nchar(as.character(num)))`.

Author(s)

Sven E. Templer <sven.templer@gmail.com>

Examples

```
#  
# use with paste to generate strings of equal size:  
paste0("observation", leading0(1:10, 3))  
#
```

lload*Load RData Objects to a List***Description**

Load multiple .RData files and return a (simplified) list.

Usage

```
lload(path = ".", pattern = ".RData", recursive = FALSE,  
      simplify = TRUE, verbose = TRUE)
```

Arguments

path	Character string with the path, as used in list.files .
pattern	A regular expression for file name patterns, as used in list.files .
recursive	Logical. Search the path recursive.
simplify	Logical, unlist when there are only unique object names.
verbose	Logical. Print information on screen about loading process.

Value

Returns a list of length n, when there are n data files loaded. All objects are stored in sublists. Names are according to files, and names of sublists to objects per file. If simplified, the list is of length m, when there are m objects in total loaded.

Author(s)

Sven E. Templer

See Also

[load](#)

lsall

List Object Details

Description

Return a data.frame with a list of all objects of a specified environment.

Usage

`lsall(envir = .GlobalEnv, ...)`

Arguments

envir	An environment where to look for objects.
...	Arguments forwarded to <code>ls</code> .

Value

Returns a data.frame with object names, lengths, classes, modes and sizes or NULL if the environment is empty.

Author(s)

Sven E. Templer

See Also[ls](#)**Examples**

```
#  
  
lsall()  
obj1 <- 1:3  
obj2 <- data.frame(1:3)  
obj3 <- list(1:3)  
lsall()  
  
#
```

mgrep1*Multiple Pattern Matching and Replacement***Description**

`mgrep1` allows multiple patterns search in character vectors, offering multicore support to parallelize search over all patterns using [mclapply](#).

Usage

```
mgrep1(patterns, text, log.fun = all, na.replace = FALSE,  
       use.which = FALSE, cores = 1, ...)
```

Arguments

<code>patterns</code>	A vector or list containing regular expressions (regex) to be searched in <code>text</code> . Coerced to character.
<code>text</code>	Character vector on which the search is performed.
<code>log.fun</code>	A function to apply on the result of matching each pattern on each element of <code>text</code> . Determines the output. See section Value .
<code>na.replace</code>	A single value to replace each NA with in the result.
<code>use.which</code>	A logical value. TRUE to convert result with <code>which</code> . Only if output <code>is.atomic</code> , otherwise ignored. Deprecated.
<code>cores</code>	Numeric value for how many cores to use for computation using <code>mclapply</code> .
<code>...</code>	Further arguments passed to functions grep1 .

Value

Depending on the function defined with `log.fun`, the return value is either

- a vector, e.g. for functions like `any`, `all` or `sum`.
- a matrix is obtained with e.g. `identity` or `as.integer`. Each row holds the result of a single pattern.
- a list is returned for functions which create results of different lengths for each element, such as `which`.

Author(s)

Sven E. Templer

See Also

`grepl`, `mclapply`

Examples

```
#  
  
# strings  
s <- c("ab", "ac", "bc", NA)  
  
# match all patterns (default)  
mgrepl(c("a", "b"), s)  
  
# match any of the patterns  
mgrepl(c("a", "b"), s, any)  
grepl("a|b", s)  
  
# return logical matrix, one column for each pattern  
mgrepl(c("a", "b"), s, identity)  
  
# return count of matches  
mgrepl(c("a", "b"), s, sum)  
  
#
```

Description

Return the series of triangular (/triangle) numbers up to a number of `n` rows of a triangle. The series has the entry number "A000217" at <https://oeis.org/A000217> and starts like this: 0, 1, 3, 6, 10, ...

Usage

```
ntri(n)
```

Arguments

n Positive integer value for sequence length.

Value

Returns an integer vector of length n.

Author(s)

Sven E. Templer (<sven.templer@gmail.com>)

nunique

Amount and Index of Unique Values

Description

Return the index or amount of unique values in a vector.

Usage

```
nunique(x, na = TRUE, ...)
```

```
uniquei(x, na = TRUE, ...)
```

Arguments

x	Numeric vector to transform.
na	Logical, TRUE to include/count NA.
...	Arguments forwarded to unique .

Author(s)

Sven E. Templer

Examples

```
#  
  
v <- c("a", "b", "a", NA)  
nunique(v)  
nunique(v, FALSE)  
uniquei(v)  
uniquei(v, FALSE)
```

```
#
```

p2star

*P Value Significance Level Indicator***Description**

Transform p-values to character (e.g. stars) indicators by significance levels with the function [symnum](#).

Usage

```
p2star(p, breaks = c(0, 0.001, 0.01, 0.05, 0.1, 1), symbols = c("***", "**",
  "*", ".", "n.s."))
```

Arguments

<code>p</code>	Vector with p values
<code>breaks</code>	The breaks from min (0) to max (1).
<code>symbols</code>	Symbols to use for values between breaks from min to max.

Author(s)

Sven E. Templer

Examples

```
#  
p2star(c(1e-5,.1,.9))  
#
```

plotn

*Plot Nothing (but a Plot)***Description**

Create a plot, with empty elements by presetting default parameters.

Usage

```
plotn(x = 0:1, y = NULL, type = "n", xlab = "", ylab = "",
  xaxt = "n", yaxt = "n", frame.plot = F, ...)
```

Arguments

<code>x</code>	Coordinates of the points.
<code>y</code>	Coordinates of the y-axis.
<code>type</code>	Plot type.
<code>xlab, ylab</code>	Axis titles.
<code>xaxt, yaxt</code>	Axis types.
<code>frame.plot</code>	Plot the frame.
<code>...</code>	Forwarded arguments to <code>plot</code> .

Details

For details about the function see [plot](#), which is called from `plotn`. More detailed information in [plot.default](#) and [par](#).

Author(s)

Sven E. Templer

`rmall`

Remove All Objects from Global Environment

Description

Remove all objects from the global environment.

Usage

`rmall(...)`

Arguments

<code>...</code>	Arguments forwarded to <code>ls</code> to get all objects.
------------------	--

Author(s)

Sven E. Templer

See Also

[rm](#), [ls](#)

Examples

```
#  
  
a <- b <- letters  
ls()  
rmall()  
ls()  
  
#
```

scale0

Scale Numeric Values to Defined Ranges

Description

Scale numeric values to a range from 0 to 1 with the function `scale0` or to a chosen range with `scaler`.

Usage

```
scale0(x)  
  
scaler(x, r = c(0, 1), b = range(x, na.rm = TRUE))
```

Arguments

x	Numeric vector to transform.
r	Numeric vector of length 2 for range to scale values of x to.
b	Numeric vector of length 2 to define the border of x to use as scaling minimum and maximum.

Author(s)

Sven E. Templer

Examples

```
#  
  
scale0(0:10)  
scale0(-1:3)  
scale0(2:3)  
  
scaler(0:10)  
scaler(0:10, 1:2)  
scaler(0:10, 1:2, c(0, 20))  
  
#
```

<code>sort</code>	<i>Sort data.frame Objects</i>
-------------------	--------------------------------

Description

Sort a data.frame by any column(s).

Usage

```
## S3 method for class 'data.frame'
sort(x, decreasing = FALSE, by = NULL, bye = NULL,
na.last = NA, ...)
```

Arguments

<code>x</code>	A data.frame.
<code>decreasing</code>	Logical, sort in decreasing order. See also sort .
<code>by</code>	Index (integer) or names of columns in <code>x</code> to sort by in that order. If both <code>by</code> and <code>bye</code> are missing, all columns are used to sort in their order.
<code>bye</code>	Unquoted column name or <code>list()</code> or <code>.()</code> with unquoted column names to sort <code>x</code> by. Not evaluated if <code>by</code> is supplied.
<code>na.last</code>	TRUE to put missing values last, FALSE to put first or NA to remove.
<code>...</code>	Ignored for the data.frame method.

Author(s)

Sven E. Templer

Examples

```
# 

d <- data.frame(a=c(1,1,1,2,NA),b=c(2,1,3,1,1),c=5:1)
d
sort(d) # sort by every column (a, then b, then c)
sort(d, TRUE, by="c") # decreasing by column 'c'
sort(d, bye=.(a,c)) # increasing by columns 'a' and then 'c'

#
```

squarematrix*Create a Square Matrix*

Description

Transform any $m \times n$ matrix to a square matrix by column/row names. Stops if no or duplicated dimnames are provided in x.

Usage

```
squarematrix(x)
```

Arguments

x Object of class **matrix**.

Value

Returns a **matrix**.

Author(s)

Sven E. Templer

Examples

```
#  
  
m <- matrix(1:6, 2, dimnames=list(2:3,1:3))  
m  
squarematrix(m)  
  
#
```

strext*Extract a Substring*

Description

This function extracts substring(s) which match a given pattern.

Usage

```
strext(x, pattern, sep = " ", mult = F, unlist = F, cores = 1)
```

Arguments

<code>x</code>	Character vector.
<code>pattern</code>	Regular expression.
<code>sep</code>	Character string which separates the fields.
<code>mult</code>	Logical, if multiple matching fields should be returned, or otherwise NA.
<code>unlist</code>	Logical, unlists multiple results.
<code>cores</code>	Integer for number of computational cores to use.

Details

The function is deprecated and will be removed with `miscset` version 2. It is recommended to use [str_extract](#) or [str_extract_all](#) instead.

Value

A list of character vectors containing the substrings that are matching `pattern` and are separated by `sep` or NA if the pattern could not be found.

Author(s)

Sven E. Templer

Examples

```
#  
library(stringr)  
  
s <- c("A1 B1 C1", "A2 B2", "AA A1", "AA", "B1 A1", "BB AB A1")  
  
strext(s, "[AB][][][:digit:]") # deprecated  
str_extract(s, "[AB][][][:digit:]")  
  
strext(s, "[AB][][][:digit:]", mult = TRUE) # deprecated  
str_extract_all(s, "[AB][][][:digit:]")  
  
strext(s, "[AB][][][:digit:]", mult = TRUE, unlist = TRUE) # deprecated  
unlist(str_extract_all(s, "[AB][][][:digit:]")) # has no <NA> values  
  
strext(s, "[C][][][:digit:]") # deprecated  
str_extract(s, "[C][][][:digit:]")  
  
#
```

str_part	<i>Split String and Return Part</i>
----------	-------------------------------------

Description

Return the nth part of a splitted string.

Usage

```
str_part(x, split, n, ..., roll = F)
```

```
strpart(x, split, n, ..., roll = F)
```

Arguments

x	Character vector.
split	Regular expression splitting strings.
n	Number of part to extract.
...	Arguments passed to <code>strsplit</code> .
roll	Logical, if to use the last when less than maximum parts.

Value

A character vector of the extracted parts.

Author(s)

Sven E. Templer

See Also

[strsplit](#)

Examples

```
#  
  
s <- c("abc", "abcd", "abc")  
  
str_part(s, "", 4)  
str_part(s, "", 4, roll=TRUE)  
  
#
```

str_rev*Reverse Text Strings*

Description

Create a reverse version of strings.

Usage

```
str_rev(x)  
strrev(x)
```

Arguments

x vector with strings. Is coerced to character.

Value

Returns a character vector with reversed strings.

Author(s)

Sven E. Templer

See Also

[rev](#)

Examples

```
#  
s <- c("abc", "asdf")  
str_rev(s)  
#
```

textable

Table to Latex

Description

This function enhances `xtable`: It can write the latex code of the table directly to a file and optionally adds a header/footer for the document structure. Also a system command can be given to convert the tex file to a pdf document, for example.

Usage

```
textable(d, file, caption = NULL, label = NULL, align = NULL,
        rownames = FALSE, topcapt = TRUE, digits = NULL, as.document = FALSE,
        landscape = FALSE, margin = 2, pt.size = 10, cmd = NULL, ...)
```

Arguments

<code>d</code>	Object (will be coerced to <code>data.frame</code>) to transform to a latex table.
<code>file</code>	Character string with output file name. If missing or "", the output is printed to the screen.
<code>caption</code>	Character vector with title of table.
<code>label</code>	Character vector with the latex label or HTML anchor.
<code>align</code>	Character vector with 'l', 'c', 'r' for aligning the columns left, centered or right. Length is either one or 1 (for <code>rownames</code> column) + number of columns in <code>d</code> (even if <code>rownames = FALSE</code>)
<code>rownames</code>	Logical, include row names of <code>d</code> .
<code>topcapt</code>	Logical, put caption and label before 'tabular'.
<code>digits</code>	Number of digits to print from numeric values.
<code>as.document</code>	Logical. TRUE to add the document definition to the output. The document class is an article and the package <code>a4paper</code> is included.
<code>landscape</code>	Logical, use a landscape format for wider tables. Only with <code>as.document=TRUE</code> .
<code>margin</code>	Margin between table and page border in cm. Only with <code>as.document=TRUE</code> .
<code>pt.size</code>	Integer from 10 to 13 for the size of the characters. Only with <code>as.document=TRUE</code> .
<code>cmd</code>	A character vector with the system command to apply on the output file. Only if <code>file</code> is given and <code>as.document</code> is TRUE. NULL or an empty string <code>system</code> is not called.
...	Forwarded arguments to print.xtable .

Details

Example for a system call:

```
cmd = "pdflatex -output-directory /path/to/files/"
```

Value

Returns a character vector invisible. If file is set, then the content is written to a file. Else it is printed to the console.

Author(s)

Sven E. Templer

See Also

[xtable](#)

Examples

```
#  
  
## Not run:  
d <- head(trees)  
dc <- 'R "trees" dataset.'  
textable(d, rownames=TRUE, digits=4, caption=dc)  
textable(d, '/tmp/trees.tex', caption=dc, as.document=TRUE,  
cmd='pdflatex --output-directory /tmp')  
  
## End(Not run)  
  
#
```

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