Package 'skMisc'

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Title Sercan Kahveci's Miscellaneous Functions

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Description	n Contains a wide range of functions.
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AnovaTable

 $Compare\ multilevel\ models$

Description

Compare multilevel models

Usage

```
AnovaTable(
    ...,
    fullmodel,
    models,
    serial = F,
    suppress = c("AIC", "deviance", "logLik")
)

## S3 method for class 'AnovaTable'
print(x)
```

Arguments

Model objects to be compared
 A model to which all other models are to be compared; only use if ... is not specified.
 Models to compare to fullmodel. Only use if ... is not specified.
 If TRUE, models are compared serially; if false, all models will be compared to the first.
 Suppress
 Character vector of column names to suppress in printed output.

Value

A data frame containing model fit metrics such as AIC, BIC, marginal R-squared (the effect size of fixed effects only), conditional R-squared (the effect size of all model terms), loglikelihood, deviance, and a likelihood ratio test.

clamp 3

clamp clamp

Description

clamp

Usage

```
clamp(val, minval, maxval)
```

Arguments

val The vector/matrix to clamp

minval Minimum value; all lower values are clamped to this value

maxval Maximum value; all higher values are clamped to this value

Value

Clamped vector.

Examples

clamp(0:10,2,8)

coerce

coerce a vector to contain only TRUE and FALSE

Description

coerce a vector to contain only TRUE and FALSE

Usage

```
coerce(x, default = FALSE)
```

Arguments

x Numeric/logical vector/matrix to coerce into TRUE/FALSE

default default returned value if NULL or NA is encountered

Value

logical vector or matrix with only T and F

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Examples

```
coerce(NULL)
# FALSE

coerce(c(T,F,NA,NA,T))
# T F F F T

coerce(matrix(c(T,T,F,F,NA,NA),nrow=2))
# [,1] [,2] [,3]
#[1,] TRUE FALSE FALSE
#[2,] TRUE FALSE FALSE
```

colVars

Compute column and row variances

Description

Compute column and row variances

Usage

```
colVars(x, na.rm = T)
rowVars(x, na.rm = T)
```

Arguments

x an input matrix of data.frame

na.rm Logical indicating whether NA values should be omitted before variance

computation

combobulate

Get all possible combinations of strings

Description

combobulate() returns all possible combinations of the provided character strings, each combination merged into a single string.

Usage

```
combobulate(...)
```

Arguments

... Character vectors to combobulate.

Value

A character vector.

compcorr 5

Examples

```
combobulate("Hello ",c("Sir","Madam"),", ",c("may I take your order?","what shall it be?"))
# [1] "Hello Sir, may I take your order?"
# [2] "Hello Madam, may I take your order?"
# [3] "Hello Sir, what shall it be?"
# [4] "Hello Madam, what shall it be?"
```

compcorr

Test if two correlation coefficients significantly differ

Description

Uses Fisher's r to z transformation, then performs a z-test on the resulting z-scores

Usage

```
compcorr(cor1, cor2, n1, n2)
```

Arguments

cor1, cor2 Correlation values being compared

n1, n2 Sample sizes of the correlation coefficients

Value

List containing the z-score and p-value

References

http://vassarstats.net/rdiff.html

CorrCrunch

Analyse the robustness of a correlation

Description

CorrCrunch() computes the minimum number of cases that need to be removed from a dataset to flip the sign of a correlation coefficient. This can be useful in distinguishing genuine correlations from spurious findings that hinge on one or two outliers. Cases are removed iteratively; in each iteration the case that maximally shrinks the correlation coefficient is removed.

Usage

```
CorrCrunch(x, y, verbose = F)
```

Arguments

x, y Numeric vectors to correlate. verbose if TRUE, prints verbose output. CorTable

Value

A list containing the number of cases that need to be removed to flip the sign of the correlation coefficient; the proportion removed cases in the data; and a data.frame without these cases.

Examples

```
CorrCrunch(mtcars$mpg,mtcars$wt)
#Holdout needed to flip the sign: 19 (63.33%)
#Final r: 0.01181141
```

CorTable

Create a Correlation Table

Description

Create a Correlation Table

Usage

```
CorTable(df, rowids, columnids, rowdf, columndf)
```

Arguments

df

A data.frame.

rowids, columnids

character vectors containing column names from df that need to be correlated.

rowdf, columndf

data.frames whose columns need to be correlated. Either df,rowids,& columnids or rowdf & columndf are required.

Value

A formatted markdown table containing correlation coefficients, p-values, and the number and percentage of cases that need to be removed to flip the sign of each correlation coefficient.

Examples

```
CorTable(mtcars,rowids=c("mpg","disp","hp"),columnids=c("drat","wt","qsec"))
CorTable(rowdf=mtcars[,c(1,3,4)],columndf=mtcars[,5:7])
```

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df.init

Initiate an empty data frame

Description

Initiate an empty data frame

Usage

```
df.init(namelist)
```

Arguments

namelist

A character vector of column names.

Value

A data frame with 0 rows.

ExpandFormula

 $Parse\ a\ lme4\ formula\ and\ return\ all\ main\ effects\ and\ interactions$ as separate terms

Description

Parse a lme4 formula and return all main effects and interactions as separate terms

Usage

```
ExpandFormula(form)
```

Arguments

form

Value

The same formula, but with all interactions and mai neffects as separate terms

Examples

```
ExpandFormula(rt ~ pull * target + (pull * target | subjectid))
#rt ~ pull + target + pull:target + (pull + target + pull:target | subjectid)
```

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ExtractRandomTerms

Extract random terms from a lme4 formula

Description

Extract random terms from a lme4 formula

Usage

```
ExtractRandomTerms(form)
```

Arguments

form

A formula

Value

A named list containing character vectors with random terms; names are group variables.

Examples

FindTopTerms

Find all model terms that are not moderated by a higher-order interaction

Description

Find all model terms that are not moderated by a higher-order interaction

Usage

```
FindTopTerms(form)
```

Arguments

form

a formula

Value

A character vector containing all model terms that are not moderated by a higher-order interaction.

Examples

```
FindTopTerms(speed ~ skill + weight * friction)
#[1] "skill" "weight:friction"
```

LevenshteinDistance 9

 ${\tt Levenshtein Distance} \quad \quad Levenshtein \ distance$

Description

Counts the number of single character deletions, insertions, and substitutions that need to be performed to turn the source string into the target string.

Usage

```
LevenshteinDistance(source, target)
```

Arguments

```
source, target Strings to be compared.
```

Value

The Levenshtein distance between the two strings.

Examples

```
LevenshteinDistance("Yoghurt", "Youtube")
```

logit.weightfun

Downweight outliers

Description

Computes weights; trials within certain bounds of the mean receive the maximum weight while trials outside these bounds are downweighted to 0 or an optional minimum.

Usage

```
logit.weightfun(
    x,
    mean = mean(x),
    s = sd(x),
    sdist = 3,
    taper = 10,
    scale = c("max", "norm"),
    min = 0
)
```

OLcrunch

Arguments

X	A numeric vector
mean	An optional mean of the vector
S	An optional standard deviation of the vector
sdist	The number of standard deviations beyond which values should be down-weighted $$
taper	A number indicating how strongly values exceeding the standard deviation should taper off $$
scale	How the weight vector should be scaled: "norm" sets the sum to 1, "max" sets the maximum to 1.
min	A minimum weight.

Value

A numeric vector of weights

	OLcrunch	Crunch Outliers	
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Description

Crunch Outliers

Usage

```
OLcrunch(x, DS = 3, hardlimit = NULL)
```

Arguments

X	Numeric vector to remove outliers from
DS	A positive numeric value. If value exceeds this many standard deviations, it is counted as an outlier
hardlimit	A numeric vector with two values. If set, values below the first value and above the second will be counted as outliers, and means/standard deviations will be computed from values within these bounds only.

Value

Vector with outlying values set to NA

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pair

Create unique pairs

Description

Combines vectors such that unique unordered sets are derived from the vectors' cross sections.

Usage

```
pair(...)
```

Arguments

... two or more vectors of equal length

Value

a character vector consisting of all input vectors concatenated term-by-term and in alphabetic order.

Examples

```
pair(1:4,4:1)
#[1] "1-4" "2-3" "2-3" "1-4"
```

read.csv.folder

Read and merge all .csv files in a folder

Description

Read and merge all .csv files in a folder

Usage

```
read.csv.folder(
  folder = "./",
  readfunc = list(read.csv, read.csv2, read.table)
)
```

Arguments

folder path to a folder

readfunc list of functions that will be used to read the files; if the first function

fails, the second function will be used, etc.

Value

A data.frame containing all merged .csv files

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RemoveTopTerms

 $Remove\ all\ possible\ models\ with\ one\ unmoderated\ term\ removed$

Description

Remove all possible models with one unmoderated term removed

Usage

```
RemoveTopTerms(form, randeff = "")
```

Arguments

form A formula

randeff The name of the group from which unmoderated terms should be removed.

To remove from fixed effects, use "" (the default).

Value

A list of formulas which have one unmoderated term removed each. The name of each list item is the term which was removed.

Examples

```
RemoveTopTerms(a ~ b * c + d + (1|e))

#$d

#a ~ b + c + b:c + (1 | e)

#$`b:c`

#a ~ b + c + d + (1 | e)
```

retype

Change classes of columns in a data.frame

Description

retype() changes the class of specific columns; retype_all() changes the class of all columns of a given class.

Usage

```
retype(df, ...)
retype_all(df, from, to)
```

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Arguments

Examples

```
sapply(ToothGrowth, class)
      len
               supp
                         dose
#"numeric" "factor" "numeric"
NewToothGrowth <- retype(ToothGrowth, supp = character(), dose = factor())</pre>
sapply(NewToothGrowth,class)
                             dose
      len.
                 supp
#"numeric" "character"
                         "factor"
sapply(mtcars,class)
       mpg
                 cyl
                          disp
                                      hp
                                             drat
# "numeric" "numeric" "numeric" "numeric" "numeric" "numeric"
      qsec
                 ٧S
                            am
                                    gear
                                              carb
# "numeric" "numeric" "numeric" "numeric" "numeric"
newmtcars <- retype_all(mtcars, "numeric", "character")</pre>
sapply(newmtcars,class)
                                disp
                                             hn
                    cyl
         mpg
  "character" "character" "character" "character"
         wt
                  qsec
                                VS
                                            am
                                                       gear
# "character" "character" "character" "character" "character"
```

smoothvect

Smooth a numeric vector using a moving window algorithm

Description

Smooth a numeric vector using a moving window algorithm

Usage

```
smoothvect(vect, width = 2, both.sides = T, alg = c("mean", "gauss"))
```

Arguments

vect

width Over how many values should the vector be averaged?

both.sides If TRUE (default), takes the mean of width values before and after the current index. If FALSE, only takes values ahead of the current index.

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Value

Smoothed numeric vector

Examples

```
temp<- smoothvect(beaver1$temp)
plot(temp,type="1")</pre>
```

splitColumn

Split a character column into multiple values

Description

Split a character column into multiple values

Usage

```
splitColumn(x, sep = ";")
```

Arguments

x a character vector to split into columnssep a caracter separating the different values

Value

a data.frame of boolean values, with rows representing the unpacked vector entries and columns indicating whether the specific value

 $theme_pecher$

Pecher theme for ggplot Based on the plot design style of prof. Diane Pecher.

Description

Pecher theme for ggplot Based on the plot design style of prof. Diane Pecher.

Usage

```
theme_pecher()
```

tokens_compound_stepwise

Compound tokens without overflowing memory and crashing R

Description

A wrapper around tokens_compound that processes your tokens in chunks, set by argument stepsize. See tokens_compound for more info.

Usage

```
tokens_compound_stepwise(
    x,
    pattern,
    stepsize = 100,
    concatenator = "_",
    valuetype = c("glob", "regex", "fixed"),
    case_insensitive = TRUE,
    join = TRUE
)
```

TransformPlots

Title

Description

Visualize how different transformations of the data will fit to a normal distribution.

Usage

TransformPlots(x)

Arguments

Х

A numeric vector.

Examples

TransformPlots(mtcars\$disp)

verify_types

trypackages

 $In stall\ packages\ if\ necessary,\ then\ load\ them.$

Description

Install packages if neccesary, then load them.

Usage

```
trypackages(...)
```

Arguments

... Unquoted names of packages to try loading, and if unable, install and

Examples

```
trypackages(stats,utils,compiler)
```

 $verify_types$

Verify variable types in bulk

Description

Verify variable types in bulk

Usage

```
verify_types(...)
```

Arguments

Named arguments, where the argument is the object to be checked and the name of the argument is the mode (numeric, list, character, etc)

Value

Returns true on success, causes error if not.

wtd.median 17

${\tt wtd.median} \qquad \qquad Weighted Median$

Description

Weighted Median

Usage

```
wtd.median(x, wts, na.rm = T)
```

Arguments

x an input vector
wts a vector of weights

na.rm Logical indicating whether NA values in the input and weight vectors

should be stripped.

Value

A weighted median of the input values and weights.

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