

Package ‘ipmisc’

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Type Package

Title Miscellaneous Functions for Data Cleaning and Analysis

Version 1.1.0

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Description Provides functions needed for data cleaning and formatting and forms data cleaning and wrangling backend for the following packages: 'ggstatsplot', 'groupedstats', 'pairwiseComparisons', and 'statsExpressions'.

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URL <https://indrajeetpatil.github.io/ipmisc/>,
<https://github.com/IndrajeetPatil/ipmisc>

BugReports <https://github.com/IndrajeetPatil/ipmisc/issues>

Depends R (>= 3.5.0)

Imports broomExtra, crayon, dplyr (>= 0.8.3), forcats, rlang (>= 0.4.2), rstudioapi, tibble (>= 2.0.1), tidyr (>= 1.0.0), zeallot

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R topics documented:

bartlett_message	2
bugs_long	3
easystats_to_tidy_names	4
ipmisc	5
iris_long	5
long_to_wide_converter	6
movies_long	7
movies_wide	8
normality_message	9
outlier_df	10
set_cwd	11
signif_column	11
sort_xy	12
specify_decimal_p	13
Index	14

bartlett_message	<i>Display homogeneity of variance test as a message</i>
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Description

A note to the user about the validity of assumptions for the default linear model.

Usage

```
bartlett_message(data, x, y, lab = NULL, k = 2, output = "message", ...)
```

Arguments

data	A dataframe (or a tibble) from which variables specified are to be taken. A matrix or tables will not be accepted.
x	The grouping variable from the dataframe data.
y	The response (a.k.a. outcome or dependent) variable from the dataframe data.
lab	A character describing label for the variable. If NULL, variable name will be used.
k	Number of digits after decimal point (should be an integer) (Default: k = 3).
output	What output is desired: "message" (default) or "stats" (or "tidy") objects.
...	Currently ignored.

Value

A list of class "htest" containing the following components:

statistic	Bartlett's K-squared test statistic.
parameter	the degrees of freedom of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Bartlett test of homogeneity of variances".
data.name	a character string giving the names of the data.

Examples

```
# getting message
bartlett_message(
  data = iris,
  x = Species,
  y = Sepal.Length,
  lab = "Iris Species"
)

# getting results from the test
bartlett_message(
  data = mtcars,
  x = am,
  y = wt,
  output = "tidy"
)
```

bugs_long

Tidy version of the "Bugs" dataset.

Description

Tidy version of the "Bugs" dataset.

Usage

```
bugs_long
```

Format

A data frame with 372 rows and 6 variables

- subject. Dummy identity number for each participant.
- gender. Participant's gender (Female, Male).
- region. Region of the world the participant was from.

- education. Level of education.
- condition. Condition of the experiment the participant gave rating for (**LDLF**: low frighteningness and low disgustingness; **LFHD**: low frighteningness and high disgustingness; **HFHD**: high frighteningness and low disgustingness; **HFHD**: high frighteningness and high disgustingness).
- desire. The desire to kill an arthropod was indicated on a scale from 0 to 10.

Details

This data set, "Bugs", provides the extent to which men and women want to kill arthropods that vary in frighteningness (low, high) and disgustingness (low, high). Each participant rates their attitudes towards all arthropods. Subset of the data reported by Ryan et al. (2013).

Source

<https://www.sciencedirect.com/science/article/pii/S0747563213000277>

Examples

```
dim(bugs_long)
head(bugs_long)
dplyr::glimpse(bugs_long)
```

easystats_to_tidy_names

Convert easystats package outputs to tidymodels conventions.

Description

Both broom package from tidymodels universe and parameters package from easystats universe can provide model summaries for a large number of model objects. This is a convenience function that converts naming conventions adopted in easystats to the ones adopted in the broom package.

Usage

```
easystats_to_tidy_names(x)
```

Arguments

x A statistical model object

Examples

```
# example model object
mod <- stats::lm(formula = wt ~ am * cyl, data = mtcars)

# `tidy`-fied output
easystats_to_tidy_names(parameters::model_parameters(mod))
```

ipmisc	ipmisc
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Description

Collection of functions to help with certain aspects of data-wrangling and data analysis that are not covered in the existing R packages.

Details

For more documentation, see [README](#) on GitHub.

iris_long	<i>Edgar Anderson's Iris Data in long format.</i>
-----------	---

Description

Edgar Anderson's Iris Data in long format.

Usage

```
iris_long
```

Format

A data frame with 600 rows and 5 variables

- id. Dummy identity number for each flower (150 flowers in total).
- Species. The species are *Iris setosa*, *versicolor*, and *virginica*.
- condition. Factor giving a detailed description of the attribute (Four levels: "Petal.Length", "Petal.Width", "Sepal.Length", "Sepal.Width").
- attribute. What attribute is being measured ("Sepal" or "Petal").
- measure. What aspect of the attribute is being measured ("Length" or "Width").
- value. Value of the measurement.

Details

This famous (Fisher's or Anderson's) iris data set gives the measurements in centimeters of the variables sepal length and width and petal length and width, respectively, for 50 flowers from each of 3 species of iris. The species are *Iris setosa*, *versicolor*, and *virginica*.

This is a modified dataset from datasets package.

Source

<https://stat.ethz.ch/R-manual/R-devel/library/datasets/html/iris.html>

Examples

```
dim(iris_long)
head(iris_long)
dplyr::glimpse(iris_long)
```

long_to_wide_converter

Converts long-format dataframe to wide-format dataframe

Description

This conversion is helpful mostly for repeated measures design.

Usage

```
long_to_wide_converter(data, x, y, paired = TRUE, ...)
```

Arguments

data	A dataframe (or a tibble) from which variables specified are to be taken. A matrix or tables will not be accepted.
x	The grouping variable from the dataframe data.
y	The response (a.k.a. outcome or dependent) variable from the dataframe data.
paired	Logical that decides whether the experimental design is repeated measures/within-subjects or between-subjects. The default is FALSE.
...	Currently ignored.

Value

A dataframe in the wide (or Cartesian) format.

Author(s)

Indrajeet Patil

Examples

```
long_to_wide_converter(
  data = iris_long,
  x = condition,
  y = value,
  paired = TRUE
)
```

movies_long	<i>Movie information and user ratings from IMDB.com (long format).</i>
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Description

Movie information and user ratings from IMDB.com (long format).

Usage

```
movies_long
```

Format

A data frame with 1,579 rows and 8 variables

- title. Title of the movie.
- year. Year of release.
- budget. Total budget (if known) in US dollars
- length. Length in minutes.
- rating. Average IMDB user rating.
- votes. Number of IMDB users who rated this movie.
- mpaa. MPAA rating.
- genre. Different genres of movies (action, animation, comedy, drama, documentary, romance, short).

Details

Modified dataset from ggplot2movies package.

The internet movie database, <http://imdb.com/>, is a website devoted to collecting movie data supplied by studios and fans. It claims to be the biggest movie database on the web and is run by amazon. More about information imdb.com can be found online, http://imdb.com/help/show_leaf?about, including information about the data collection process, http://imdb.com/help/show_leaf?infosource.

Movies were are identical to those selected for inclusion in movies_wide but this dataset has been constructed such that every movie appears in one and only one genre category.

Source

<https://CRAN.R-project.org/package=ggplot2movies>

Examples

```
dim(movies_long)
head(movies_long)
dplyr::glimpse(movies_long)
```

`movies_wide`*Movie information and user ratings from IMDB.com (wide format).*

Description

Movie information and user ratings from IMDB.com (wide format).

Usage

```
movies_wide
```

Format

A data frame with 1,579 rows and 13 variables

- `title`. Title of the movie.
- `year`. Year of release.
- `budget`. Total budget in millions of US dollars
- `length`. Length in minutes.
- `rating`. Average IMDB user rating.
- `votes`. Number of IMDB users who rated this movie.
- `mpaa`. MPAA rating.
- `action`, `animation`, `comedy`, `drama`, `documentary`, `romance`, `short`. Binary variables representing if movie was classified as belonging to that genre.
- `NumGenre`. The number of different genres a film was classified in an integer between one and four

Details

Modified dataset from `ggplot2movies` package.

The internet movie database, <http://imdb.com/>, is a website devoted to collecting movie data supplied by studios and fans. It claims to be the biggest movie database on the web and is run by amazon. More information about `imdb.com` can be found online, http://imdb.com/help/show_leaf?about, including information about the data collection process, http://imdb.com/help/show_leaf?infosource.

Movies were selected for inclusion if they had a known length and had been rated by at least one `imdb` user. Small categories such as documentaries and NC-17 movies were removed.

Source

<https://CRAN.R-project.org/package=ggplot2movies>

Examples

```
dim(movies_wide)
head(movies_wide)
dplyr::glimpse(movies_wide)
```

normality_message *Display normality test result as a message.*

Description

A note to the user about the validity of assumptions for the default linear model.

Usage

```
normality_message(x, lab = NULL, k = 2, output = "message", ...)
```

Arguments

x	A numeric vector.
lab	A character describing label for the variable. If NULL, a generic "x" label will be used.
k	Number of digits after decimal point (should be an integer) (Default: k = 3).
output	What output is desired: "message" (default) or "stats" (or "tidy") objects.
...	Additional arguments (ignored).

Value

A list with class "htest" containing the following components:

statistic	the value of the Shapiro-Wilk statistic.
p.value	an approximate p-value for the test. This is said in Royston (1995) to be adequate for $p.value < 0.1$.
method	the character string "Shapiro-Wilk normality test".
data.name	a character string giving the name(s) of the data.

Examples

```
# message
normality_message(
  x = anscombe$x1,
  lab = "x1",
  k = 3
)

# statistical test object
normality_message(
  x = anscombe$x2,
  output = "tidy"
)
```

`outlier_df`*Adding a column to dataframe describing outlier status*

Description

Adding a column to dataframe describing outlier status

Usage

```
outlier_df(data, x, y, outlier.label, outlier.coef = 1.5, ...)
```

Arguments

<code>data</code>	A dataframe (or a tibble) from which variables specified are to be taken. A matrix or tables will not be accepted.
<code>x</code>	The grouping variable from the dataframe <code>data</code> .
<code>y</code>	The response (a.k.a. outcome or dependent) variable from the dataframe <code>data</code> .
<code>outlier.label</code>	Label to put on the outliers that have been tagged. This can't be the same as <code>x</code> argument.
<code>outlier.coef</code>	Coefficient for outlier detection using Tukey's method. With Tukey's method, outliers are below (1st Quartile) or above (3rd Quartile) <code>coef</code> times the Inter-Quartile Range (IQR) (Default: 1.5).
<code>...</code>	Additional arguments.

Value

The dataframe entered as `data` argument is returned with two additional columns: `isanoutlier` and `outlier` denoting which observation are outliers and their corresponding labels.

Examples

```
# adding column for outlier and a label for that outlier
outlier_df(
  data = morley,
  x = Expt,
  y = Speed,
  outlier.label = Run,
  outlier.coef = 2
) %>%
  dplyr::arrange(outlier)
```

set_cwd	<i>Setting Working Directory in RStudio to where the R Script is.</i>
---------	---

Description

This function will change the current working directory to whichever directory the R script you are currently working on is located. This preempts the trouble of setting the working directory manually.

Usage

```
set_cwd()
```

Value

Path to changed working directory.

Note

This function will work **only with RStudio IDE**. Reference: <https://eranraviv.com/r-tips-and-tricks-working-directory/>

signif_column	<i>Creating a new column with significance labels</i>
---------------	---

Description

This function will add a new column with significance labels to a dataframe containing p -values.

Usage

```
signif_column(data, p, ...)
```

Arguments

data	Data frame from which variables specified are preferentially to be taken.
p	The column containing p -values.
...	Currently ignored.

Value

Returns the dataframe in tibble format with an additional column corresponding to APA-format statistical significance labels.

Author(s)

Indrajeet Patil

Examples

```
# preparing a new dataframe
df <- cbind.data.frame(
  x = 1:5,
  y = 1,
  p.value = c(0.1, 0.5, 0.00001, 0.05, 0.01)
)

# dataframe with significance column
signif_column(data = df, p = p.value)
```

 sort_xy

Sorting y column in data by x.

Description

Sorting y column in data by x.

Usage

```
sort_xy(data, x, y, sort = "none", .fun = mean, ...)
```

Arguments

data	A dataframe (or a tibble) from which variables specified are to be taken. A matrix or tables will not be accepted.
x	The grouping variable from the dataframe data.
y	The response (a.k.a. outcome or dependent) variable from the dataframe data.
sort	If "ascending" (default), x-variable factor levels will be sorted based on increasing values of y-variable. If "descending", the opposite. If "none", no sorting will happen.
.fun	n summary function. It should take one vector for fct_reorder, and two vectors for fct_reorder2, and return a single value.
...	Currently ignored.

Examples

```
sort_xy(ggplot2::msleep, vore, brainwt, sort = "ascending")
```

specify_decimal_p *Formatting numeric (p-)values*

Description

Function to format an R object for pretty printing with a specified (k) number of decimal places. The function also allows really small p -values to be denoted as " $p < 0.001$ " rather than " $p = 0.000$ ". Note that if `p.value` is set to `TRUE`, the minimum value of k allowed is 3. If k is set to less than 3, the function will ignore entered k value and use $k = 3$ instead. **Important:** This function is not vectorized.

Usage

```
specify_decimal_p(x, k = 3, p.value = FALSE)
```

Arguments

<code>x</code>	A numeric value.
<code>k</code>	Number of digits after decimal point (should be an integer) (Default: $k = 3$).
<code>p.value</code>	Decides whether the number is a p -value (Default: <code>FALSE</code>).

Value

Formatted numeric value.

Author(s)

Indrajeet Patil

Examples

```
specify_decimal_p(x = 0.00001, k = 2, p.value = TRUE)
specify_decimal_p(x = 0.008, k = 2, p.value = TRUE)
specify_decimal_p(x = 0.008, k = 3, p.value = FALSE)
```

Index

*Topic **datasets**

bugs_long, [3](#)

iris_long, [5](#)

movies_long, [7](#)

movies_wide, [8](#)

bartlett_message, [2](#)

bugs_long, [3](#)

easystats_to_tidy_names, [4](#)

ipmisc, [5](#)

iris_long, [5](#)

long_to_wide_converter, [6](#)

movies_long, [7](#)

movies_wide, [8](#)

normality_message, [9](#)

outlier_df, [10](#)

set_cwd, [11](#)

signif_column, [11](#)

sort_xy, [12](#)

specify_decimal_p, [13](#)