Package 'scipub'

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Title Summarize Data for Scientific Publication

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Description Create and format tables and APA statistics for scientific publication. This includes making a 'Table 1' to summarize demographics across groups, correlation tables with significance indicated by stars, and extracting formatted statistical summarizes from simple tests for in-text notation. The package also includes functions for Winsorizing data based on a Z-statistic cutoff.
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apastat

Format simple statistic test results for scientific publication

Description

The apastat function summarizes statistic test results scientific publication. This currently will take stats::t.test, stats::cor.test, or stats::lm results as input. The output is intended to be included as in-text parenthetical statistics in publication.

Usage

```
apastat(test, roundN = 2, es = c(TRUE, FALSE), ci = c(TRUE, FALSE), var = NULL)
```

Arguments

test	The stats::t.test, stats::cor.test, or stats::lm object to be formatted.
roundN	The number of decimal places to round all output to (default=2).
es	Include effect side (Cohen's d for t-test or 2-level factor lm variable), default to TRUE.
ci	Include confidence interval of estimate, default to TRUE.
var	Only for lm object, select name of variable to summarize (default=NULL), if NULL, will summarize overall model fit.

Value

Output formatted statistics

```
apastat(stats::cor.test(psydat$Age, psydat$Height))
apastat(stats::t.test(Height ~ Sex, data = psydat))
apastat(stats::lm(data = psydat, Height ~ Age + Sex))
apastat(stats::lm(data = psydat, Height ~ Age + Sex), var = "Age")
```

correltable 3

correltable	Create correlation table (with stars for significance) for scientific publication
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Description

The correltable function can be used to create correlation table (with stars for significance) for scientific publication This is intended to summarize correlations between (vars) from an input dataset (data). Correlations are based on stats::cor, use and method follow from that function. Stars indicate significance: *p<.05, **p<.01, ***p<.001 For formatting, variables can be renamed, numbers can be rounded, upper or lower triangle only can be selected (or whole matrix), and empty columns/rows can be dropped if using triangles. For more compact columns, variable names can be numbered in the rows and column names will be corresponding numbers. If only cross-correlation between two sets of variables is desired (no correlations within a set of variables), vars2 and var_names can be used. This function will drop any non-numeric variables by default. Requires tidyverse and stats libraries.

Usage

```
correltable(
  data,
  vars = NULL,
  var_names = vars,
  vars2 = NULL,
  var_names2 = vars2,
  method = c("pearson", "spearman"),
  use = c("pairwise", "complete"),
  round_n = 2,
  tri = c("upper", "lower", "all"),
  cutempty = c(FALSE, TRUE),
  colnum = c(FALSE, TRUE),
  html = c(FALSE, TRUE)
)
```

Arguments

data	The input dataset.
vars	A list of the names of variables to correlate, e.g. c("Age","height","WASI"), if NULL, all variables in data will be used.
var_names	An optional list to rename the vars colnames in the output table, e.g. c("Age (years)","Height (inches)","IQ"). Must match vars in length. If not supplied, vars will be printed as is.
vars2	If cross-correlation between two sets of variables is desired, add a second list of variables to correlate with vars; Overrides tri, cutempty, and colnum.
var_names2	An optional list to rename the vars2 colnames in the output table If not supplied, vars2 will be printed as is.
method	Type of correlation to calculate $c("pearson", "spearman")$, based on stats::cor, $default = "pearson"$.
use	Use pairwise.complete.obs or restrict to complete cases c("pairwise", "complete"), based on stats::cor, default = "pairwise".

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round_n The number of decimal places to round all output to (default=2). Select output formatting c("upper", "lower", "all"); KEEP the upper triangle, tri lower triangle, or all values, default ="upper. If keeping only upper/lower triangle with tri, cut empty row/column, default=FALSE. cutempty For more concise column names, number row names and just use corresponding colnum numbers as column names, default=FALSE, if TRUE overrides cutempty. html Format as html in viewer or not (default=F, print in console), needs library(htmlTable) installed.

Value

Output Table 1

Examples

```
correltable(data = psydat)
correltable(
  data = psydat, vars = c("Age", "Height", "iq"),
  tri = "lower", html = TRUE
correltable(
  data = psydat, vars = c("Age", "Height", "iq"),
var_names = c("Age (months)", "Height (inches)", "IQ"),
  tri = "upper", colnum = TRUE, html = TRUE
correltable(
  data = psydat, vars = c("Age", "Height", "iq"),
  var_names = c("Age (months)", "Height (inches)", "IQ"),
  vars2 = c("depressT", "anxT"),
  var_names2 = c("Depression T", "Anxiety T"), html = TRUE
)
```

FullTable1

Create Table 1 of group summary with stats for scientific publication

Description

The FullTable1 function can be used to create a Table1 for scientific publication. This is intended to summarize demographic and other variables (vars) split by a grouping variable (strata) from an input dataset (data). Continuous variables will be summarized as mean (SD) and tested across groups using t-test or ANOVA (for 3+ level strata). Categorical variables will be summarized as N (%) and tested across groups as chi-squared. Effect sizes for group differences will be calculated as Cohen's d, partial eta-squared, Odds Ratio, Cramer's V depending on the test. Requires tidyverse and stats libraries.

Usage

```
FullTable1(
 data,
  strata = NULL,
  vars = NULL,
```

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```
var_names = vars,
factor_vars = NULL,
round_n = 2,
es_col = c(TRUE, FALSE),
p_col = c(TRUE, FALSE),
stars = c("col", "name", "stat", "none"),
html = c(FALSE, TRUE)
)
```

Arguments

data The input dataset (will be converted to tibble).

strata The grouping variable of interest (converted to factor), if NULL will make one

column table.

vars A list of variables to summarize, e.g. c("Age", "sex", "WASI").

var_names An optional list to rename the variable colnames in the output table, e.g. c("Age

(years)", "Sex", "IQ"). Must match vars in length. If not supplied, vars will be

printed as is.

factor_vars An optional list of variables from vars to use as class factor, e.g. c("sex").

Note that any character, factor, or logical class variables will be summarized as

categorical by default.

round_n The number of decimal places to round output to (default=2).

es_col Include a column for effect size of group difference? (default=T).

p_col Include a column for p-value of group difference? (default=TRUE).

stars Where to include stars indicating significance of group differences. Options:

"col"=separate column (default), "name"= append to variable name, "stat"= ap-

pend to group difference statistic, "none" for no stars.

html Format as html in viewer or not (default=FALSE, print in console), needs li-

brary(htmlTable) installed.

Value

Output Table 1

```
FullTable1(
  data = psydat,
  vars = c("Age", "Height", "depressT"), strata = "Sex"
)
FullTable1(
  data = psydat,
  vars = c("Age", "Height", "depressT"), strata = "Sex"
)
FullTable1(
  data = psydat, vars = c("Age", "Sex", "Height", "depressT"),
  var_names = c("Age (months)", "Sex", "Height (inches)", "Depression T"),
  strata = "Income", stars = "name", p_col = FALSE
)
```

6 partial_correltable

Description

The partial_correltable function can be used to create partial correlation table (with stars for significance) for scientific publication This is intended to summarize partial correlations between (vars) from an input dataset (data), residualizing all vars by partialvars. This function allows for numeric, binary, and factor variables as partialvars, but only numeric vars are used and any non-numeric vars will be dropped. All other flags follow from scipub::correltable. Correlations are based on stats::cor, use and method follow from that function. Stars indicate significance: *p<.05, **p<.01, ***p<.001 For formatting, variables can be renamed, numbers can be rounded, upper or lower triangle only can be selected (or whole matrix), and empty columns/rows can be dropped if using triangles. For more compact columns, variable names can be numbered in the rows and column names will be corresponding numbers. Requires tidyverse and stats libraries.

Usage

```
partial_correltable(
  data,
  vars = NULL,
  var_names = vars,
  partialvars = NULL,
  partialvar_names = partialvars,
  method = c("pearson", "spearman"),
  use = c("pairwise", "complete"),
  round_n = 2,
  tri = c("upper", "lower", "all"),
  cutempty = c(FALSE, TRUE),
  colnum = c(FALSE, TRUE),
  html = c(FALSE, TRUE)
)
```

Arguments

data The input dataset.

vars A list of the names of 2+ variables to correlate, e.g. c("Age", "height", "WASI").

All variables must be numeric.

var_names An optional list to rename the vars colnames in the output table, e.g. c("Age

(years)", "Height (inches)", "IQ"). Must match vars in length. If not supplied,

vars will be printed as is.

partialvars A list of the names of 1+ variables to partial out, e.g. c("iq", "Sex", "Income").

Can include numeric, binary, factor variables.

partialvar_names

An optional list to rename the partialvars colnames in the output table, e.g. c("IQ (WASI)", "Sex", "Income"). Must match partialvar_names in length. If not supplied, partialvar_names will be printed as is.

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method	Type of correlation to calculate c("pearson", "spearman"), based on stats::cor, default = "pearson".
use	Use pairwise.complete.obs or restrict to complete cases c("pairwise", "complete"), based on stats::cor, default = "pairwise".
round_n	The number of decimal places to round all output to (default=2).
tri	Select output formatting c("upper", "lower", "all"); KEEP the upper triangle, lower triangle, or all values, default = "upper.
cutempty	If keeping only upper/lower triangle with tri, cut empty row/column, default=FALSE.
colnum	For more concise column names, number row names and just use corresponding numbers as column names, default=FALSE, if TRUE overrides cutempty.
html	Format as html in viewer or not (default=F, print in console), needs library(htmlTable) installed.

Value

Output Table 1

Examples

```
partial_correltable(
  data = psydat, vars = c("Age", "Height", "iq"),
  partialvars = c("Sex", "Income"),
  tri = "lower", html = TRUE
)

partial_correltable(
  data = psydat, vars = c("Age", "Height", "iq"),
  var_names = c("Age (months)", "Height (inches)", "IQ"),
  partialvars = c("Sex", "Income"),
  tri = "upper", colnum = TRUE, html = TRUE
)

partial_correltable(
  data = psydat, vars = c("Age", "Height", "iq"),
  var_names = c("Age (months)", "Height (inches)", "IQ"),
  partialvars = c("anxT"),
  partialvar_names = "Anxiety",
  tri = "all", html = TRUE
)
```

psydat

Sample demographic and clinical data for 5,000 children

Description

An example dataset containing demographic and clinical data for 5,000 children. The variables are as follows:

Usage

```
data(psydat)
```

8 winsorZ

Format

```
A data frame with 5000 rows and 7 variables:
```

```
Age age in months (107.2–136.4)

Sex biological sex, 4 missing value (M, F)

Income reported family income, 404 missing values (<50K, >=100K, >=50K&<100K)

Height height in inches, 7 missing values (36.05–84.51)

iq cognition test, 179 missing values (34.86–222.99)

depressT depression symptom severity T-score, 8 missing values (48.53–91.32)

anxT anxiety symptom severity T-score, 8 missing values (48.76–93,67)
```

winsorZ

Winsorize outliers based on z-score cutoff to next most extreme non-outlier value

Description

The winsorZ function identifies outliers based on Z-score cutoff and replaces with the next most extreme non-outlier value. This involves z-scoring the variable and identifying/replacing any cases beyond the z-score threshold. The winsorZ_find function is an optional companion to flag any Z-score outliers to tally as needed.

Usage

```
winsorZ(x, zbound = 3)
```

Arguments

x The input variable to Winsorize.

zbound The Z-score cutoff (default=3, i.e. outliers are Z>3 | Z<-3).

Value

Output Winsorized variable

```
winsorZ(psydat$iq)
## Not run:
psydat %>%
   dplyr::select(c(iq, anxT)) %>%
   map(winsorZ)
psydat %>% mutate_at(c("iq", "anxT"), list(~ winsorZ(.)))
psydat %>% mutate_if(is.double, list(~ winsorZ(.)))
## End(Not run)
```

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winsorZ_find Identify outliers based on z-score cutoff that are Winsorized by the winsorZ function	winsorZ_find	Identify outliers based on z-score cutoff that are Winsorized by the winsorZ function
--	--------------	---

Description

The winsorZ_find function is an optional companion to the winsorZ function. The winsorZ function identifies Z-score outliers and replaces with the next most extreme non-outlier value. The winsorZ_find function finds/identifies these Z-score outliers (outliers=1, non-outliers=0).

Usage

```
winsorZ_find(x, zbound = 3)
```

Arguments

x The input variable to check for Z-score outliers.

zbound The Z-score cutoff (default=3, i.e. outliers are Z>3 | Z<-3).

Value

Output logical variable of Z-score outliers

```
summary(winsorZ_find(psydat$iq))
## Not run:
psydat %>% mutate_at(c("iq", "anxT"), list(out = ~ winsorZ_find(.)))
## End(Not run)
```

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