

Package: WJSmisc (via r-universe)

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Title Miscellaneous functions from W. Joel Schneider

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Version 0.3

Description Several functions I find useful.

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URL <https://github.com/wjschne/WJSmisc>

BugReports <https://github.com/wjschne/WJSmisc/issues>

Imports corrr, dplyr, forcats, ggplot2, magrittr, patchwork, psych, rlang ($\geq 0.1.2$), stats, grDevices, stringr, superheat, rstudioapi, shiny, tibble, tidyr, GPArotation, utils, purrr, rlist, scales, readr, tidyselect, ggtext, signs

Encoding UTF-8

RoxygenNote 7.2.3

Suggests roxygen2, testthat ($\geq 2.1.0$)

Config/pak/sysreqs make libicu-dev libjpeg-dev libpng-dev libxml2-dev libssl-dev libx11-dev zlib1g-dev

Repository <https://wjschne.r-universe.dev>

RemoteUrl <https://github.com/wjschne/WJSmisc>

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angle2hjust	<i>Convert angles to ggplot2 hjust</i>
-------------	--

Description

Convert angles to ggplot2 hjust

Usage

```
angle2hjust(theta, multiplier = 1.5, as_degrees = FALSE)
```

Arguments

<code>theta</code>	angle in radians
<code>multiplier</code>	distance from point
<code>as_degrees</code>	use degrees instead of radians

Value

numeric

<code>angle2vjust</code>	<i>Convert angles to ggplot2 vjust</i>
--------------------------	--

Description

Convert angles to ggplot2 vjust

Usage

```
angle2vjust(theta, multiplier = 1.5, as_degrees = FALSE)
```

Arguments

<code>theta</code>	angle in radians
<code>multiplier</code>	distance from point
<code>as_degrees</code>	use degrees instead of radians

Value

numeric

Examples

```
library(tibble)
library(ggplot2)
library(dplyr)
xy_ratio <- pi
tibble(theta = seq(0,2*pi, length.out = 9),
       y = sin(theta),
       slope = cos(theta) * xy_ratio,
       text_angle = atan(slope) + pi / 2) %>%
  ggplot(aes(theta,y)) +
  stat_function(fun = sin) +
  geom_point() +
  geom_label(aes(label = LETTERS[1:9],
                vjust = angle2vjust(text_angle, multiplier = 1.5),
                hjust = angle2hjust(text_angle, multiplier = 1.5)),
            label.size = NA,
            label.padding = unit(1, "mm")) +
```

```

scale_x_continuous(expression(theta),
                    breaks = seq(0,2*pi, length.out = 9),
                    labels = label_parsed(c("0", "frac(pi,4)", "frac(pi,2)",
                                             "frac(3 * pi,4)", "pi", "frac(5*pi,4)",
                                             "frac(3 * pi,2)", "frac(7*pi,4)", "2*pi"))) +
scale_y_continuous(expression(sin(theta))) +
coord_fixed(ratio = xy_ratio, clip = "off") +
theme_minimal()

```

<code>attach_function</code>	<i>Converts the default values of a function's arguments to variables and attaches them to the global environment</i>
------------------------------	---

Description

Converts the default values of a function's arguments to variables and attaches them to the global environment

Usage

```
attach_function(f)
```

Arguments

<code>f</code>	Function
----------------	----------

Value

Attaches function arguments to global environment

Examples

```

my_function <- function(x, y = 2) x + y

# Sets y to 2
attach_function(my_function)

```

<code>cor2pcor</code>	<i>Converts a correlation matrix to a partial correlation matrix</i>
-----------------------	--

Description

Converts a correlation matrix to a partial correlation matrix

Usage

```
cor2pcor(R)
```

`cornames<-`

5

Arguments

`R` correlation matrix

Value

Partial correlation matrix

Examples

```
R <- matrix(.6, nrow = 3, ncol = 3)
diag(R) <- 1
cor2pcor(R)
```

`cornames<-` *Name for square arrays like correlation matrices*

Description

Name for square arrays like correlation matrices

Usage

```
cornames(x) <- value
```

Arguments

`x` a square array, matrix, or data.frame

`value` a vector of names

Value

a named square array, matrix, or data.frame

Examples

```
R <- tri2cor(.5)
cornames(R) <- c("A", "B")
```

cor_ellipse	<i>Generate correlation ellipse data</i>
-------------	--

Description

Generate correlation ellipse data

Usage

```
cor_ellipse(  
  r = 0,  
  mean = c(0, 0),  
  sd = c(1, 1),  
  p = 0.95,  
  split_x = NULL,  
  split_y = NULL,  
  n_points = 10000  
)
```

Arguments

<code>r</code>	Correlation coefficient
<code>mean</code>	Vector of 2 means
<code>sd</code>	Vector of 2 standard deviations
<code>p</code>	Proportion of data ellipse covers
<code>split_x</code>	Split ellipse at x value
<code>split_y</code>	Split ellipse at y value
<code>n_points</code>	Number of points

Value

data.frame

Examples

```
cor_ellipse(r = .75)
```

`cor_heat`*Correlation plot*

Description

Correlation plot

Usage

```
cor_heat(  
  d,  
  test_names = colnames(d),  
  margins = 0.55,  
  text_size = 4,  
  dendrograms = TRUE,  
  palette_col = c("firebrick", "white", "royalblue"),  
  x_label_angle = 90,  
  reorder_vars = TRUE,  
  heat.lim = c(-1, 1),  
  heat.pal.values = seq(0, 1, 1/(length(palette_col) - 1)),  
  ...  
)
```

Arguments

<code>d</code>	Data or correlation matrix
<code>test_names</code>	Vector of names of variables
<code>margins</code>	Width of margins for labels
<code>text_size</code>	Size of text
<code>dendrograms</code>	If TRUE, add dendrograms
<code>palette_col</code>	A vector of three colors for heatmap
<code>x_label_angle</code>	Angle of x-axis label
<code>reorder_vars</code>	If TRUE, reorder variables based on cluster analysis
<code>heat.lim</code>	Vector of the lower and upper bounds on heat map
<code>heat.pal.values</code>	Vector of values of where on the scale each color in <code>palette_col</code> falls. Defaults to even spacing.
<code>...</code>	Additional parameters passed to <code>superheat::superheat</code>

<code>cor_text</code>	<i>Format text of a correlation matrix</i>
-----------------------	--

Description

Format text of a correlation matrix

Usage

```
cor_text(r, digits = 2, remove_leading_zero = TRUE, remove_diagonal = FALSE)
```

Arguments

<code>r</code>	a matrix of correlations
<code>digits</code>	Number of digits to round
<code>remove_leading_zero</code>	If TRUE, remove leading zero from all correlations
<code>remove_diagonal</code>	If TRUE, remove diagonal of ones

Value

a matrix of correlations as text

Examples

```
cor_text(matrix(.5,2,2))
```

<code>cross_vectors</code>	<i>Create unique combinations of vectors</i>
----------------------------	--

Description

Create unique combinations of vectors

Usage

```
cross_vectors(..., sep = "_")
```

Arguments

<code>...</code>	vectors
<code>sep</code>	Separate character

Value

A character vector

Examples

```
cross_vectors(c("a", "b"), c(1,2))
```

<code>df2matrix</code>	<i>Convert data.frame and tibbles to matrices with named rows and columns</i>
------------------------	---

Description

Convert data.frame and tibbles to matrices with named rows and columns

Usage

```
df2matrix(d, first_col_is_row_names = TRUE)
```

Arguments

<code>d</code>	data.frame or tibble
<code>first_col_is_row_names</code>	TRUE if first column has row names

Value

matrix

Examples

```
d <- data.frame(rname = c("x1", "x2"), x1 = c(1,2), x2 = c(3,4))
df2matrix(d)
```

<code>geom_richlabel</code>	<i>A wrapper for ggtext::geom_richtext</i>
-----------------------------	--

Description

A wrapper for ggtext::geom_richtext

Usage

```
geom_richlabel(
  mapping,
  label.margin = unit(2, "mm"),
  label.padding = unit(0.5, "mm"),
  label.color = NA,
  fill = "white",
  text_size = 12,
  ...
)
```

Arguments

<code>mapping</code>	<code>aes()</code>
<code>label.margin</code>	grid unit margin
<code>label.padding</code>	grid unit margin
<code>label.color</code>	color
<code>fill</code>	color
<code>text_size</code>	text size in point units
<code>...</code>	additional parameters passed to <code>ggtext::geom_richtext</code>

Value

`ggtext::geom_richtext`

Examples

```
library(tibble)
library(ggplot2)
library(dplyr)
library(WJSmisc)
tibble(x = 0, y = 0, l = "A") %>%
  ggplot() +
  geom_richtlabel(aes(x,y,label = l))
```

`get_quote`

Retrieve text within a span or div with a named id

Description

Retrieve text within a span or div with a named id

Usage

```
get_quote(id, file, blockquote = TRUE)
```

Arguments

<code>id</code>	the id string (without #).
<code>file</code>	file name
<code>blockquote</code>	return as block quote (defaults to TRUE)

Value

a character vector of length 1

`ggsvg` *Save ggplot as .pdf, then convert to .svg via pdf2svg*

Description

Save ggplot as .pdf, then convert to .svg via pdf2svg

Usage

```
ggsvg(f, width = 6.5, height = 6.5, ...)
```

Arguments

<code>f</code>	Filename of the svg file. Omit the ".svg" at the end.
<code>width</code>	width passed to ggplot2::ggsave
<code>height</code>	width passed to ggplot2::ggsave
<code>...</code>	Additional parameters passed to ggplot2::ggsave

`ggtext_size` *Convert ggplot theme font size to geom_text size*

Description

Convert ggplot theme font size to geom_text size

Usage

```
ggtext_size(base_size, ratio = 0.8)
```

Arguments

<code>base_size</code>	theme font size
<code>ratio</code>	ratio of text size to theme font size. Defaults to .8 so that geom text will be the same size as default sized axis labels.

Value

numeric vector

Examples

```
ggtext_size(16)
```

`insert_latex_color` *Insert latex colors into a matrix*

Description

Insert latex colors into a matrix

Usage

```
insert_latex_color(  
  m,  
  color_cells,  
  color_rownames = NULL,  
  color_colnames = NULL  
)
```

Arguments

<code>m</code>	matrix of values
<code>color_cells</code>	matrix of latex colors for matrix cells
<code>color_rownames</code>	vector of latex colors for row names
<code>color_colnames</code>	vector of latex colors for column names

Value

matrix

Examples

```
# A matrix of zeros  
m <- matrix(0, nrow = 2, ncol = 2)  
  
# A matrix of NA values the same size as m  
latex_colors <- m * NA  
  
# Make the top row red  
latex_colors[1,] <- "red"  
  
# Insert colors into m  
insert_latex_color(m, latex_colors)
```

irt_plot_app	<i>IRT Plot Shiny App</i>
--------------	---------------------------

Description

IRT Plot Shiny App

Usage

```
irt_plot_app()
```

latexarray	<i>Make latex array</i>
------------	-------------------------

Description

Make latex array

Usage

```
latexarray(  
  M,  
  left = "",  
  right = "",  
  env = "array",  
  includenames = TRUE,  
  align = NULL,  
  lines = TRUE  
)
```

Arguments

M	A matrix
left	left delimiter options: [(
right	right delimiter options:)]
env	Array environment
includenames	Include column and row names
align	Column alignment. For a three column matrix, alignment defaults to centering columns("ccc"). If there are row labels, the default would be "r ccc" to right-align the row labels and separate them with vertical line.
lines	Include lines separating column and row names

Value

character string

Examples

```
M <- diag(3)
colnames(M) <- LETTERS[1:3]
rownames(M) <- LETTERS[1:3]
latexarray(M)
```

lm_matrix*Regression from correlation matrix*

Description

Regression from correlation matrix

Usage

```
lm_matrix(R, ind, dep)
```

Arguments

R	correlation matrix
ind	independent variables
dep	dependent variable

Value

list of coefficients and R2

make_indicators*Make latent variable indicators with rbeta_ms function*

Description

Make latent variable indicators with rbeta_ms function

Usage

```
make_indicators(
  latent,
  indicators = NULL,
  mu = 0.8,
  sigma = 0.05,
  k = 3,
  digits = 3
)
```

Arguments

<code>latent</code>	name of latent variable
<code>indicators</code>	vector of indicator names (assigned automatically if left NULL)
<code>mu</code>	mean of standardized coefficients
<code>sigma</code>	sd of standardized coefficients
<code>k</code>	number of indicator variables
<code>digits</code>	number of digits to round coefficients

Value

lavaan code for latent variable assignment

Examples

```
make_indicators("depression", mu = 0.8, sigma = 0.05, k = 4)
```

<code>modregplot</code>	<i>Simple slopes plot</i>
-------------------------	---------------------------

Description

generates simple slopes plot from moderated regression equation

Usage

```
modregplot(
  predictor_range = c(-4, 4),
  moderator_values = c(-1, 0, 1),
  intercept = 0,
  predictor_coef = 0,
  moderator_coef = 0,
  interaction_coef = 0,
  predictor_label = "X",
  criterion_label = "Y",
  moderator_label = "Moderator"
)
```

Arguments

<code>predictor_range</code>	a length 2 vector of the range of values to be plotted on the predictor variable, Default: c(-4, 4)
<code>moderator_values</code>	a vector of moderator values to be plotted, Default: c(-1, 0, 1)
<code>intercept</code>	the intercept of the regression equation, Default: 0

```

predictor_coef      the regression coefficient for the predictor variable, Default: 0
moderator_coef      the regression coefficient for the moderator variable, Default: 0
interaction_coef     the interaction term coefficient, Default: 0
predictor_label      the label for the predictor variable, Default: 'X'
criterion_label      the label for the moderator variable, Default: 'Y'
moderator_label      PARAM_DESCRIPTION, Default: 'Moderator'

```

Value

a ggplot of the simple slopes

Examples

```

modregplot(
  predictor_range = c(-2, 2),
  moderator_values = c(Low = -1, High = 1),
  intercept = 6,
  predictor_coef = 2,
  moderator_coef = 0,
  interaction_coef = 1,
  predictor_label = "Psychopathy",
  criterion_label = "Aggression",
  moderator_label = "Impulsivity"
)

```

multivariate_ci

General a multivariate confidence interval for a set of scores

Description

General a multivariate confidence interval for a set of scores

Usage

```
multivariate_ci(x, rxx, mu, sigma, ci = 0.95, v_names = names(x))
```

Arguments

```

x          a vector of scores
rxx        a vector reliability coefficients
mu         a vector means
sigma      a covariance matrix
ci         confidence level
v_names    a vector of names

```


Value

data.frame

Examples

```
x_wisc <- c(
  vci = 130,
  vsi = 130,
  fri = 70,
  wmi = 130,
  psi = 130
)
rxx_wisc <- c(
  vci = .92,
  vsi = .92,
  fri = .93,
  wmi = .92,
  psi = .88
)
R_wisc <- ("
index vci vsi fri wmi psi
vci 1.00 0.59 0.59 0.53 0.30
vsi 0.59 1.00 0.62 0.50 0.36
fri 0.59 0.62 1.00 0.53 0.31
wmi 0.53 0.50 0.53 1.00 0.36
psi 0.30 0.36 0.31 0.36 1.00") |>
  readr::read_tsv() |>
  tibble::column_to_rownames("index") |>
  as.matrix()
multivariate_ci(
  x = x_wisc,
  rxx = rxx_wisc,
  mu = rep(100, 5),
  sigma = R_wisc * 225
)
```

parallel_analysis *ggplot of parallel analysis from the psych package*

Description

ggplot of parallel analysis from the psych package

Usage

```
parallel_analysis(
  d,
  fm = "pa",
  factor_based = TRUE,
```

```
vcolors = c("firebrick", "royalblue"),  
font_family = "sans",  
...  
)
```

Arguments

<code>d</code>	data to be analyzed
<code>fm</code>	factor method passed to <code>psych::fa.parallel</code>
<code>factor_based</code>	TRUE is factor-based and FALSE is principal component-based
<code>vcolors</code>	vector of 2 colors for lines
<code>font_family</code>	Name of font
<code>...</code>	parameters passed to <code>psych::fa.parallel</code>

Examples

```
d <- psych::bfi[,1:25]  
parallel_analysis(d)
```

```
paste_matrix_from_clipboard  
Paste matrix code from clipboard
```

Description

Paste matrix code from clipboard

Usage

```
paste_matrix_from_clipboard(digits = 2, as_matrix = TRUE)
```

Arguments

<code>digits</code>	Number of digits to round
<code>as_matrix</code>	Convert to matrix. Defaults to 'TRUE'

pdf2svg	<i>Function that converts pdf files to svg. Must have the pdf2svg program installed (https://github.com/dawbarton/pdf2svg)</i>
---------	---

Description

Function that converts pdf files to svg. Must have the pdf2svg program installed (<https://github.com/dawbarton/pdf2svg>)

Usage

```
pdf2svg(f)
```

Arguments

f	Filename of the pdf file. Omit the ".pdf" at the end.
---	---

plotnorm	<i>Plot a normal distribution shading below x</i>
----------	---

Description

Plot a normal distribution shading below x

Usage

```
plotnorm(  
  x = 0,  
  mu = 0,  
  sigma = 1,  
  below = TRUE,  
  show_proportion = TRUE,  
  show_x = TRUE,  
  show_param = TRUE,  
  text_size = 14,  
  font_family = "sans",  
  shade_fill = "royalblue"  
)
```

Arguments

x	number to divide normal distribution
mu	mean of normal distribution
sigma	standard deviation of normal distribution
below	If TRUE, shade lower portion of normal distribution

<code>show_proportion</code>	If TRUE, display proportions
<code>show_x</code>	If TRUE, display x value
<code>show_param</code>	If TRUE, display mean and standard deviation
<code>text_size</code>	Base text size
<code>font_family</code>	Name of font
<code>shade_fill</code>	Color of shaded region

Examples

```
plotnorm(90, 100, 15)
```

<code>plot_loading</code>	<i>Creates a plot of factor loadings</i>
---------------------------	--

Description

Creates a plot of factor loadings

Usage

```
plot_loading(
  f,
  font_family = "sans",
  font_size = 14,
  loading_text_size = font_size * 0.8,
  factor_names = sort(colnames(f$loadings)),
  nudge_loadings = 0.05
)
```

Arguments

<code>f</code>	output of a factor analysis from the <code>psych::fa</code> function
<code>font_family</code>	Name of font
<code>font_size</code>	Size of font
<code>loading_text_size</code>	size of loading font,
<code>factor_names</code>	names of the factors #'
<code>nudge_loadings</code>	nudge loadings on x dimension

Examples

```
library(GPARotation)
library(psych)
fit <- fa(psych::bfi[,1:25], nfactors = 5)
plot_loading(fit)
```

prob_label	<i>Format numeric probabilities as text labels</i>
------------	--

Description

Format numeric probabilities as text labels

Usage

```
prob_label(  
  p,  
  accuracy = 0.01,  
  digits = NULL,  
  max_digits = NULL,  
  remove_leading_zero = TRUE,  
  round_zero_one = TRUE  
)
```

Arguments

<code>p</code>	numeric vector of probabilities
<code>accuracy</code>	accuracy of rounding
<code>digits</code>	Optional. Number of digits to round. Overrides accuracy parameter
<code>max_digits</code>	Optional. Maximum zeros or nines before rounding to 0 or 1
<code>remove_leading_zero</code>	Removes leading zero from probability
<code>round_zero_one</code>	Apply rounding to 0 and 1

Value

character vector

Examples

```
prob_label(seq(0,1, 0.1))
```

```
proportion2percentile
```

Rounds proportions to significant digits both near 0 and 1, then converts to percentiles

Description

Rounds proportions to significant digits both near 0 and 1, then converts to percentiles

Usage

```
proportion2percentile(  
  p,  
  digits = 2,  
  remove_leading_zero = TRUE,  
  add_percent_character = FALSE  
)
```

Arguments

p	probability
digits	rounding digits
remove_leading_zero	If TRUE, remove leading zero
add_percent_character	If TRUE, add percent character

Value

character vector

Examples

```
proportion2percentile(0.011111)
```

```
proportion_round
```

Rounds proportions to significant digits both near 0 and 1

Description

Rounds proportions to significant digits both near 0 and 1

Usage

```
proportion_round(p, digits = 2)
```

Arguments

`p` probability
`digits` rounding digits

Value

numeric vector

Examples

```
proportion_round(0.01111)
```

<code>pthreshold</code>	<i>Probability a true score will be below a threshold</i>
-------------------------	---

Description

Probability a true score will be below a threshold

Usage

```
pthreshold(x, threshold, rxx, mu = 0, sigma = 1)
```

Arguments

`x` observed score
`threshold` threshold
`rxx` reliability coefficient
`mu` population mean
`sigma` population standard deviation

Value

probability

Examples

```
pthreshold(x = .5, threshold = 1, rxx = 0.9)
```

pvalueAPA	<i>APA p-value rounding</i>
-----------	-----------------------------

Description

APA p-value rounding

Usage

```
pvalueAPA(p, inline = FALSE, mindigits = 2, maxdigits = 3)
```

Arguments

<code>p</code>	probability
<code>inline</code>	to be used in an inline rmarkdown (default is FALSE)
<code>mindigits</code>	minimum rounding digits
<code>maxdigits</code>	maximum rounding digits

Value

character

Examples

```
pvalueAPA(0.01111)
```

rbeta_ms	<i>Random beta distribution with specified mean and sd</i>
----------	--

Description

Random beta distribution with specified mean and sd

Usage

```
rbeta_ms(n = 1, mu = 0.5, sigma = 0.025)
```

Arguments

<code>n</code>	Number of data points
<code>mu</code>	Mean of random beta distribution
<code>sigma</code>	SD of random beta distribution

Value

a vector of numeric values

Examples

```
rbeta_ms(n = 5, mu = 0.8, sigma = 0.1)
```

`remove_leading_zero` *Remove leading zero from numbers*

Description

Remove leading zero from numbers

Usage

```
remove_leading_zero(x, digits = 2, ...)
```

Arguments

<code>x</code>	vector of numbers
<code>digits</code>	rounding digits
<code>...</code>	Arguments passed to <code>formatC</code>

Value

vector of characters

Examples

```
remove_leading_zero(c(0.5,-0.2))
```

`rotate2dmatrix` *Rotate a 2-column matrix*

Description

Rotate a 2-column matrix

Usage

```
rotate2dmatrix(x, theta, degrees = FALSE, origin = c(0, 0))
```

Arguments

<code>x</code>	a 2-column matrix
<code>theta</code>	angle
<code>degrees</code>	if TRUE, theta is in degrees instead of radians
<code>origin</code>	point of rotation

Value

a rotated 2-column matrix

Examples

```
x <- matrix(seq(10), ncol = 2)
rotate2dmatrix(x, pi)
```

signs_centered	<i>centered signed numbers</i>
----------------	--------------------------------

Description

A wrapper function for the signs::signs function. It adds a figure space to negative numbers so that it appear as if the minus sign does not affect the number's centering.

Usage

```
signs_centered(x, space = " ", ...)
```

Arguments

<code>x</code>	a numeric vector
<code>space</code>	a character to be added to negative numbers (defaults to a figure space)
<code>...</code>	parameters passed to signs:signs

Value

a vector of numbers converted to characters

Examples

```
library(ggplot2)
d <- data.frame(x = -4:0, y = -4:0)
# In these 2 plots, Compare the centering of the negative numbers on the x-axis
ggplot(d, aes(x,y))
ggplot(d, aes(x,y)) +
  scale_x_continuous(labels = signs_centered)
```

skewed_axis	<i>Draw skewed axes in ggplot2</i>
-------------	------------------------------------

Description

Draw skewed axes in ggplot2

Usage

```
skewed_axis(  
  theta,  
  axis_title = "X",  
  draw_ticks = TRUE,  
  draw_axis_text = TRUE,  
  remove_origin = TRUE,  
  tick_height = 0.02,  
  lwd = 0.5,  
  text_size = 12,  
  color = NA,  
  family = NA,  
  mu = 0,  
  sigma = 1,  
  tick_label_interval = ifelse(sigma%%3 == 0, 3, 2)  
)
```

Arguments

<code>theta</code>	angle in radians
<code>axis_title</code>	character
<code>draw_ticks</code>	logical
<code>draw_axis_text</code>	logical
<code>remove_origin</code>	logical
<code>tick_height</code>	height of ticks
<code>lwd</code>	line width
<code>text_size</code>	size of text
<code>color</code>	color of lines
<code>family</code>	font family
<code>mu</code>	mean of variable
<code>sigma</code>	standard deviation of variable
<code>tick_label_interval</code>	interval of ticks

Value

plot

Examples

```
library(ggplot2)
ggplot(data.frame(x = c(0,1), y = c(4,4))) +
  skewed_axis(pi / 2, color = "black") +
  skewed_axis(pi / 4, color = "black") +
  skewed_axis(0, color = "black") +
  theme_void() +
  coord_equal()
```

snake2subscript

Convert snake case to subscript

Description

Convert snake case to subscript

Usage

```
snake2subscript(
  x,
  sep = "_",
  prefix = "<sub>",
  suffix = "</sub>",
  collapse = NULL,
  recycle0 = FALSE
)
```

Arguments

x	character
sep	separator defaults to "_"
prefix	character prefix
suffix	character suffix
collapse	character parameter passed to paste0
recycle0	parameter passed to paste0

Value

a character

Examples

```
snake2subscript("x_1")
```

str_wrap_equal	<i>Wrap sentence strings into lines of roughly equal width</i>
----------------	--

Description

Wrap sentence strings into lines of roughly equal width

Usage

```
str_wrap_equal(x, max_width = 30L, sep = "\n")
```

Arguments

x	a character vector
max_width	the maximum number of characters in a line (unless a word is longer than 'max_width')
sep	character string that separates text lines

Value

a character vector

Examples

```
library(ggplot2)
library(stringr)
library(dplyr)
d <- data.frame(
  Item = c(
    "Lorem ipsum dolor sit amet, consectetur adipiscing elit.",
    "Duis pretium arcu quis nibh elementum, sed aliquam enim dignissim.",
    "Nullam et ornare enim, et egestas odio.",
    "Aliquam posuere ante quis magna rutrum, id elementum nulla sodales.",
    "Interdum et malesuada fames ac ante ipsum primis in faucibus.",
    "Aenean rutrum lorem at metus pretium, malesuada porta tellus facilisis.",
    "Vestibulum at convallis enim.",
    "Nam malesuada bibendum rutrum.",
    "Donec risus sapien, pulvinar vitae porttitor non, lobortis ac felis."
  ),
  Proportion = seq(.1,.9,.1)
) |>
mutate(Item = forcats::fct_inorder(Item))

# Axis labels with stringr::str_wrap
ggplot(d, aes(Proportion, Item)) +
  geom_col() +
  scale_y_discrete(NULL, labels = \(x) str_wrap(x, width = 25L))

# Axis labels with WJSmisc::str_wrap_equal
```

```
ggplot(d, aes(Proportion, Item)) +
  geom_col() +
  scale_y_discrete(NULL, labels = \(x) str_wrap_equal(x, max_width = 25L))
```

tri2cor *Create square correlation matrix from lower triangle*

Description

Create square correlation matrix from lower triangle

Usage

```
tri2cor(x, variable_names = NULL)
```

Arguments

x vector of correlations
variable_names a vector of variable names

Value

square matrix

Examples

```
tri2cor(c(.2,.3,.4))
```

vector_angle *Compute the angle of a vector*

Description

Compute the angle of a vector

Usage

```
vector_angle(x, origin = c(0, 0), degrees = FALSE, allow_negative = FALSE)
```

Arguments

x A length-2 vector
origin A length-2 vector
degrees If TRUE, returns angles in degrees instead of radians
allow_negative If TRUE, returns angles between $-\pi$ and π (-180 and +180). If FALSE, returns angles between 0 and $2 * \pi$ (0 and 360)

Value

A length-1 vector

Examples

```
vector_angle(c(1,1))  
vector_angle(c(1,1), degrees = TRUE)
```

x2z	<i>Convert x to a z-score</i>
-----	-------------------------------

Description

Convert x to a z-score

Usage

```
x2z(x, mu = mean(x, na.rm = T), sigma = stats::sd(x, na.rm = T))
```

Arguments

x	a numeric vector
mu	mean
sigma	standard deviation

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