

Package ‘modelssummary’

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Type Package

Title Summary Tables and Plots for Statistical Models and Data: Beautiful, Customizable, and Publication-Ready

Description Create beautiful and customizable tables to summarize several statistical models side-by-side. Draw coefficient plots, multi-level cross-tabs, dataset summaries, balance tables (a.k.a. “Table 1s”), and correlation matrices. This package supports dozens of statistical models, and it can produce tables in HTML, LaTeX, Word, Markdown, PDF, PowerPoint, Excel, RTF, JPG, or PNG. Tables can easily be embedded in ‘Rmarkdown’ or ‘knitr’ dynamic documents.

Version 0.7.0

URL <https://vincentarelbundock.github.io/modelssummary/>

BugReports <https://github.com/vincentarelbundock/modelssummary/issues/>

Depends R (>= 3.5.0)

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kableExtra (>= 1.2.1),
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performance,
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tidyr,
tidyselect

Suggests broom.mixed,
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<code>coef_rename</code>	<i>Rename model terms</i>
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Description

A convenience function which can be passed to the `coef_rename` argument of the `modelsummary` function.

Usage

```
coef_rename(
  x,
  factor = TRUE,
  factor_name = TRUE,
  backticks = TRUE,
  titlecase = TRUE,
  underscore = TRUE,
  asis = TRUE
)
```

Arguments

<code>x</code>	character vector of term names to transform
<code>factor</code>	boolean remove the "factor()" label
<code>factor_name</code>	boolean remove the "factor()" label and the name of the variable
<code>backticks</code>	boolean remove backticks
<code>titlecase</code>	boolean convert to title case
<code>underscore</code>	boolean replace underscores by spaces
<code>asis</code>	boolean remove the I from as-is formula calls

<code>datasummary</code>	<i>Summary tables using 2-sided formulae: crosstabs, frequencies, table 1s and more.</i>
--------------------------	--

Description

`datasummary` can use any summary function which produces one numeric or character value per variable. The examples section of this documentation shows how to define custom summary functions. The package also ships with several shortcut summary functions: `Min`, `Max`, `Mean`, `Median`, `Var`, `SD`, `NPercent`, `NUnique`, `Ncol`, `P0`, `P25`, `P50`, `P75`, `P100`.

Usage

```

datasummary(
  formula,
  data,
  output = "default",
  fmt = 2,
  title = NULL,
  notes = NULL,
  align = NULL,
  add_columns = NULL,
  add_rows = NULL,
  sparse_header = TRUE
)

```

Arguments

formula	A two-sided formula to describe the table: rows ~ columns. See the Examples section for a mini-tutorial and the Details section for more resources.
data	A data.frame (or tibble)
output	filename or object type (character string) <ul style="list-style-type: none"> • Supported filename extensions: .html, .tex, .md, .txt, .png, .jpg. • Supported object types: "default", "html", "markdown", "latex", "latex_tabular", "data.frame", "modelsummary_list", "gt", "kableExtra", "huxtable", "flextable". • To change the default output format, type <code>options(modelsummary_default = "latex")</code>, where <code>latex</code> can be any of the valid object types listed above. • Warning: the output argument <i>cannot</i> be used when customizing tables with external packages. • See the 'Details' section below for more information.
fmt	determines how to format numeric values <ul style="list-style-type: none"> • integer: the number of digits to keep after the period format(<code>round(x, fmt), nsmall=fmt</code>) • character: passed to the <code>sprintf</code> function (e.g., <code>'\</code> • function: returns a formatted character string.
title	string
notes	list or vector of notes to append to the bottom of the table.
align	A character string of length equal to the number of columns in the table. "lcr" means that the first column will be left-aligned, the 2nd column center-aligned, and the 3rd column right-aligned.
add_columns	a data.frame (or tibble) with the same number of rows as your main table.
add_rows	a data.frame (or tibble) with the same number of columns as your main table. By default, rows are appended to the bottom of the table. You can define a "position" attribute of integers to set the row positions. See Examples section below.
sparse_header	TRUE or FALSE. TRUE eliminates column headers which have a unique label across all columns, except for the row immediately above the data. FALSE keeps all headers. The order in which terms are entered in the formula determines the order in which headers appear. For example, <code>x~mean*z</code> will print the mean-related header above the z-related header.'

Details

Visit the 'modelssummary' website for more usage examples: <https://vincentarelbundock.github.io/modelssummary>

The 'datasummary' function is a thin wrapper around the 'tabular' function from the 'tables' package. More details about table-making formulas can be found in the 'tables' package documentation: [?tables::tabular](#)

Hierarchical or "nested" column labels are only available for these output formats: kableExtra, gt, html, rtf, and LaTeX. When saving tables to other formats, nested labels will be combined to a "flat" header.

Examples

```
## Not run:

# The left-hand side of the formula describes rows, and the right-hand side
# describes columns. This table uses the "mpg" variable as a row and the "mean"
# function as a column:

datasummary(mpg ~ mean, data = mtcars)

# This table uses the "mean" function as a row and the "mpg" variable as a column:

datasummary(mean ~ mpg, data = mtcars)

# Display several variables or functions of the data using the "+"
# concatenation operator. This table has 2 rows and 2 columns:

datasummary(hp + mpg ~ mean + sd, data = mtcars)

# Nest variables or statistics inside a "factor" variable using the "*" nesting
# operator. This table shows the mean of "hp" and "mpg" for each value of
# "cyl":

mtcars$cyl <- as.factor(mtcars$cyl)
datasummary(hp + mpg ~ cyl * mean, data = mtcars)

# If you don't want to convert your original data
# to factors, you can use the 'Factor()'
# function inside 'datasummary' to obtain an identical result:

datasummary(hp + mpg ~ Factor(cyl) * mean, data = mtcars)

# You can nest several variables or statistics inside a factor by using
# parentheses. This table shows the mean and the standard deviation for each
# subset of "cyl":

datasummary(hp + mpg ~ cyl * (mean + sd), data = mtcars)

# Summarize all numeric variables with 'All()'
datasummary(All(mtcars) ~ mean + sd, data = mtcars)

# Define custom summary statistics. Your custom function should accept a vector
# of numeric values and return a single numeric or string value:

minmax <- function(x) sprintf("[%.2f, %.2f]", min(x), max(x))
mean_na <- function(x) mean(x, na.rm = TRUE)
```

```

datasummary(hp + mpg ~ minmax + mean_na, data = mtcars)

# To handle missing values, you can pass arguments to your functions using
# '*Arguments()'

datasummary(hp + mpg ~ mean * Arguments(na.rm = TRUE), data = mtcars)

# For convenience, 'modelsummary' supplies several convenience functions
# with the argument `na.rm=TRUE` by default: Mean, Median, Min, Max, SD, Var,
# P0, P25, P50, P75, P100, NUnique, Histogram

datasummary(hp + mpg ~ Mean + SD + Histogram, data = mtcars)

# These functions also accept a 'fmt' argument which allows you to
# round/format the results

datasummary(hp + mpg ~ Mean * Arguments(fmt = "%.3f") + SD * Arguments(fmt = "%.1f"), data = mtcars)

# Save your tables to a variety of output formats:
f <- hp + mpg ~ Mean + SD
datasummary(f, data = mtcars, output = 'table.html')
datasummary(f, data = mtcars, output = 'table.tex')
datasummary(f, data = mtcars, output = 'table.md')
datasummary(f, data = mtcars, output = 'table.docx')
datasummary(f, data = mtcars, output = 'table.pptx')
datasummary(f, data = mtcars, output = 'table.jpg')
datasummary(f, data = mtcars, output = 'table.png')

# Display human-readable code
datasummary(f, data = mtcars, output = 'html')
datasummary(f, data = mtcars, output = 'markdown')
datasummary(f, data = mtcars, output = 'latex')

# Return a table object to customize using a table-making package
datasummary(f, data = mtcars, output = 'gt')
datasummary(f, data = mtcars, output = 'kableExtra')
datasummary(f, data = mtcars, output = 'flextable')
datasummary(f, data = mtcars, output = 'huxtable')

# add_rows
new_rows <- data.frame(a = 1:2, b = 2:3, c = 4:5)
attr(new_rows, 'position') <- c(1, 3)
datasummary(mpg + hp ~ mean + sd, data = mtcars, add_rows = new_rows)

## End(Not run)

```

datasummary_balance *Balance table: Summary statistics for different subsets of the data (e.g., control and treatment groups)*

Description

Balance table: Summary statistics for different subsets of the data (e.g., control and treatment groups)

Usage

```

datasummary_balance(
  formula,
  data,
  output = "default",
  fmt = 1,
  title = NULL,
  notes = NULL,
  align = NULL,
  add_columns = NULL,
  add_rows = NULL,
  dinm = TRUE,
  dinm_statistic = "std.error",
  ...
)

```

Arguments

formula	a one-sided formula with the "condition" or "column" variable on the right-hand side.
data	A data.frame (or tibble). If this data includes columns called "blocks", "clusters", and/or "weights", the "estimatr" package will consider them when calculating the difference in means.
output	filename or object type (character string) <ul style="list-style-type: none"> • Supported filename extensions: .html, .tex, .md, .txt, .png, .jpg. • Supported object types: "default", "html", "markdown", "latex", "latex_tabular", "data.frame", "modelsummary_list", "gt", "kableExtra", "huxtable", "flectable". • To change the default output format, type <code>options(modelsummary_default = "latex")</code>, where <code>latex</code> can be any of the valid object types listed above. • Warning: the output argument <i>cannot</i> be used when customizing tables with external packages. • See the 'Details' section below for more information.
fmt	determines how to format numeric values <ul style="list-style-type: none"> • integer: the number of digits to keep after the period format (<code>round(x, fmt), nsmall=fmt</code>) • character: passed to the <code>sprintf</code> function (e.g., <code>\</code>) • function: returns a formatted character string.
title	string
notes	list or vector of notes to append to the bottom of the table.
align	A character string of length equal to the number of columns in the table. "lcr" means that the first column will be left-aligned, the 2nd column center-aligned, and the 3rd column right-aligned.
add_columns	a data.frame (or tibble) with the same number of rows as your main table.
add_rows	a data.frame (or tibble) with the same number of columns as your main table. By default, rows are appended to the bottom of the table. You can define a "position" attribute of integers to set the row positions. See Examples section below.

`dinm` TRUE calculates a difference in means with uncertainty estimates. This option is only available if the `estimatr` package is installed. If data includes columns named "blocks", "clusters", or "weights", this information will be taken into account automatically by `estimatr::difference_in_means`.

`dinm_statistic` string: "std.error" or "p.value"

... all other arguments are passed through to the extractor and table-making functions. This allows users to pass arguments directly to `modelsummary` in order to affect the behavior of other functions behind the scenes. Examples include:

- `broom::tidy(exponentiate=TRUE)` to exponentiate logistic regression
- `kableExtra::kbl(escape=FALSE)` to avoid escaping math characters in `kableExtra` tables.
- `performance::model_performance(metrics="RMSE")` to select goodness-of-fit statistics to extract using the `performance` package (must have set `options(modelsummary_get="easystats")` first).

Examples

```
## Not run:
datasummary_balance(~am, mtcars)

## End(Not run)
```

`datasummary_correlation`

Generate a correlation table for all numeric variables in your dataset.

Description

The names of the variables displayed in the correlation table are the names of the columns in the data. You can rename those columns (with or without spaces) to produce a table of human-readable variables.

Usage

```
datasummary_correlation(
  data,
  output = "default",
  fmt = 2,
  title = NULL,
  notes = NULL,
  method = "pearson",
  ...
)
```

Arguments

`data` A `data.frame` (or `tibble`)

`output` filename or object type (character string)

- Supported filename extensions: `.html`, `.tex`, `.md`, `.txt`, `.png`, `.jpg`.

- Supported object types: "default", "html", "markdown", "latex", "latex_tabular", "data.frame", "modelsummary_list", "gt", "kableExtra", "huxtable", "flextable".
 - To change the default output format, type `options(modelsummary_default = "latex")`, where `latex` can be any of the valid object types listed above.
 - Warning: the output argument *cannot* be used when customizing tables with external packages.
 - See the 'Details' section below for more information.
- `fmt` determines how to format numeric values
- integer: the number of digits to keep after the period format(`round(x, fmt)`, `nsmall=fmt`)
 - character: passed to the `sprintf` function (e.g., `\`)
 - function: returns a formatted character string.
- `title` string
- `notes` list or vector of notes to append to the bottom of the table.
- `method` character or function
- character: "pearson", "kendall", "spearman", or "pearspear" (Pearson correlations above and Spearman correlations below the diagonal)
 - function: takes a `data.frame` with numeric columns and returns a square matrix or `data.frame` with unique `row.names` and `colnames` corresponding to variable names. Note that the `datasummary_correlation_format` can often be useful for formatting the output of custom correlation functions.
- ... other parameters are passed through to the table-making packages. This can be used, for example, to pass arguments such as `escape=FALSE` to `kableExtra`.

Examples

```
## Not run:
library(modelsummary)

# clean variable names (base R)
dat <- mtcars[, c("mpg", "hp")]
colnames(dat) <- c("Miles / Gallon", "Horse Power")
datasummary_correlation(dat)

# clean variable names (tidyverse)
library(tidyverse)
dat <- mtcars %>%
  select(`Miles / Gallon` = mpg,
         `Horse Power` = hp)
datasummary_correlation(dat)

# alternative methods
datasummary_correlation(dat, method = "pearspear")

# custom function
cor_fun <- function(x) cor(x, method = "kendall")
datasummary_correlation(dat, method = cor_fun)

# rename columns alphabetically and include a footnote for reference
note <- sprintf("(%s) %s", letters[1:ncol(dat)], colnames(dat))
note <- paste(note, collapse = "; ")

colnames(dat) <- sprintf("(%s)", letters[1:ncol(dat)])
```

```

datasummary_correlation(dat, notes = note)

# `datasummary_correlation_format`: custom function with formatting
dat <- mtcars[, c("mpg", "hp", "disp")]

cor_fun <- function(x) {
  out <- cor(x, method = "kendall")
  datasummary_correlation_format(
    out,
    fmt = 2,
    upper_triangle = "x",
    diagonal = ".")
}

datasummary_correlation(dat, method = cor_fun)

# use kableExtra and psych to color significant cells
library(psych)
library(kableExtra)

dat <- mtcars[, c("vs", "hp", "gear")]

cor_fun <- function(dat) {
  # compute correlations and format them
  correlations <- data.frame(cor(dat))
  correlations <- datasummary_correlation_format(correlations, fmt = 2)

  # calculate pvalues using the `psych` package
  pvalues <- psych::corr.test(dat)$p

  # use `kableExtra::cell_spec` to color significant cells
  for (i in 1:nrow(correlations)) {
    for (j in 1:ncol(correlations)) {
      if (pvalues[i, j] < 0.05 && i != j) {
        correlations[i, j] <- cell_spec(correlations[i, j], background = "pink")
      }
    }
  }
  return(correlations)
}

# The `escape=FALSE` is important here!
datasummary_correlation(dat, method = cor_fun, escape = FALSE)

## End(Not run)

```

```
datasummary_correlation_format
```

Format the content of a correlation table

Description

Mostly for internal use, but can be useful when users supply a function to the method argument of `datasummary_correlation`.

Usage

```

datasummary_correlation_format(
  x,
  fmt,
  leading_zero = FALSE,
  diagonal = NULL,
  upper_triangle = NULL
)

```

Arguments

x	square numeric matrix
fmt	determines how to format numeric values <ul style="list-style-type: none"> integer: the number of digits to keep after the period <code>format(round(x, fmt), nsmall=fmt)</code> character: passed to the <code>sprintf</code> function (e.g., <code>"\</code> function: returns a formatted character string.
leading_zero	boolean. If <code>FALSE</code> , leading zeros are removed
diagonal	character or <code>NULL</code> . If character, all elements of the diagonal are replaced by the same character (e.g., <code>"1"</code>).
upper_triangle	character or <code>NULL</code> . If character, all elements of the upper triangle are replaced by the same character (e.g., <code>" "</code> or <code> "."</code>).

Examples

```

library(modelsummary)

dat <- mtcars[, c("mpg", "hp", "disp")]

cor_fun <- function(x) {
  out <- cor(x, method = "kendall")
  datasummary_correlation_format(
    out,
    fmt = 2,
    upper_triangle = "x",
    diagonal = ".")
}

datasummary_correlation(dat, method = cor_fun)

```

 datasummary_df

Draw a table from a data.frame

Description

Draw a table from a data.frame

Usage

```

datasummary_df(
  data,
  output = "default",
  fmt = 2,
  align = NULL,
  hrule = NULL,
  title = NULL,
  notes = NULL,
  add_rows = NULL,
  add_columns = NULL,
  ...
)

```

Arguments

data	A data.frame (or tibble)
output	filename or object type (character string) <ul style="list-style-type: none"> • Supported filename extensions: .html, .tex, .md, .txt, .png, .jpg. • Supported object types: "default", "html", "markdown", "latex", "latex_tabular", "data.frame", "modelsummary_list", "gt", "kableExtra", "huxtable", "flextable". • To change the default output format, type <code>options(modelsummary_default = "latex")</code>, where <code>latex</code> can be any of the valid object types listed above. • Warning: the output argument <i>cannot</i> be used when customizing tables with external packages. • See the 'Details' section below for more information.
fmt	determines how to format numeric values <ul style="list-style-type: none"> • integer: the number of digits to keep after the period format (<code>round(x, fmt), nsmall=fmt</code>) • character: passed to the <code>sprintf</code> function (e.g., <code>\</code>) • function: returns a formatted character string.
align	A character string of length equal to the number of columns in the table. "lcr" means that the first column will be left-aligned, the 2nd column center-aligned, and the 3rd column right-aligned.
hrule	position of horizontal rules (integer vector)
title	string
notes	list or vector of notes to append to the bottom of the table.
add_rows	a data.frame (or tibble) with the same number of columns as your main table. By default, rows are appended to the bottom of the table. You can define a "position" attribute of integers to set the row positions. See Examples section below.
add_columns	a data.frame (or tibble) with the same number of rows as your main table.
...	all other arguments are passed through to the extractor and table-making functions. This allows users to pass arguments directly to <code>modelsummary</code> in order to affect the behavior of other functions behind the scenes. Examples include: <ul style="list-style-type: none"> • <code>broom::tidy(exponentiate=TRUE)</code> to exponentiate logistic regression • <code>kableExtra::kbl(escape=FALSE)</code> to avoid escaping math characters in <code>kableExtra</code> tables.

- `performance::model_performance(metrics="RMSE")` to select goodness-of-fit statistics to extract using the performance package (must have set `options(modelsummary_get="easystats")` first).

 datasummary_skim

Quick overview of numeric or categorical variables

Description

This function was inspired by the excellent `skimr` package for R.

Usage

```
datasummary_skim(
  data,
  type = "numeric",
  output = "default",
  fmt = "%.1f",
  histogram = TRUE,
  title = NULL,
  notes = NULL,
  align = NULL,
  ...
)
```

Arguments

<code>data</code>	A <code>data.frame</code> (or tibble)
<code>type</code>	of variables to summarize: "numeric" or "categorical" (character)
<code>output</code>	filename or object type (character string) <ul style="list-style-type: none"> • Supported filename extensions: <code>.html</code>, <code>.tex</code>, <code>.md</code>, <code>.txt</code>, <code>.png</code>, <code>.jpg</code>. • Supported object types: "default", "html", "markdown", "latex", "latex_tabular", "data.frame", "modelsummary_list", "gt", "kableExtra", "huxtable", "flextable". • To change the default output format, type <code>options(modelsummary_default = "latex")</code>, where <code>latex</code> can be any of the valid object types listed above. • Warning: the output argument <i>cannot</i> be used when customizing tables with external packages. • See the 'Details' section below for more information.
<code>fmt</code>	determines how to format numeric values <ul style="list-style-type: none"> • integer: the number of digits to keep after the period format (<code>round(x, fmt)</code>, <code>nsmall=fmt</code>) • character: passed to the <code>sprintf</code> function (e.g., <code>%</code>) • function: returns a formatted character string.
<code>histogram</code>	include a histogram (TRUE/FALSE). Supported for: <ul style="list-style-type: none"> • <code>type = "numeric"</code> • output is "html", "default", "jpg", "png", or "kableExtra" • PDF and HTML documents compiled via Rmarkdown or knitr • See the examples section below for an example of how to use <code>datasummary</code> to include histograms in other formats such as markdown.

title	string
notes	list or vector of notes to append to the bottom of the table.
align	A character string of length equal to the number of columns in the table. "lcr" means that the first column will be left-aligned, the 2nd column center-aligned, and the 3rd column right-aligned.
...	all other arguments are passed through to the extractor and table-making functions. This allows users to pass arguments directly to <code>modelsummary</code> in order to affect the behavior of other functions behind the scenes. Examples include: <ul style="list-style-type: none"> • <code>broom::tidy(exponentiate=TRUE)</code> to exponentiate logistic regression • <code>kableExtra::kbl(escape=FALSE)</code> to avoid escaping math characters in <code>kableExtra</code> tables. • <code>performance::model_performance(metrics="RMSE")</code> to select goodness-of-fit statistics to extract using the <code>performance</code> package (must have set <code>options(modelsummary_get="easystats")</code> first).

Examples

```
## Not run:
dat <- mtcars
dat$vs <- as.logical(dat$vs)
dat$cyl <- as.factor(dat$vs)
datasummary_skim(dat)
datasummary_skim(dat, "categorical")

# You can use `datasummary` to produce a similar table in different formats.
# Note that the `Histogram` function relies on unicode characters. These
# characters will only display correctly in some operating systems, under some
# locales, using some fonts. Displaying such histograms on Windows computers
# is notoriously tricky. The `modelsummary` authors cannot provide support to
# display these unicode histograms.

f <- All(mtcars) ~ Mean + SD + Min + Median + Max + Histogram
datasummary(f, mtcars, output="markdown")

## End(Not run)
```

dvnames

Title models with their dependent variables

Description

A convenience function for use with a regression model or list of regression models. Returns a named list of models, where the names are the models' respective dependent variables. Pass your list of models to `dvnames` before sending to `modelsummary` to automatically get dependent variable-titled columns.

Usage

```
dvnames(models, number = FALSE, fill = "Model")
```

Arguments

models	A regression model or list of regression models
number	Should the models be numbered (1), (2), etc., in addition to their dependent variable names?
fill	If <code>insight::find_response()</code> cannot find a response, the column title to use in its place. Set to ' ' to leave blank.

Examples

```
m1 <- lm(mpg ~ hp, data = mtcars)
m2 <- lm(mpg ~ hp + wt, data = mtcars)

# Without dvnames, column names are Model 1 and Model 2
modelsummary(list(m1, m2))

# With dvnames, they are "mpg" and "mpg"
modelsummary(dvnames(list(m1, m2)))
```

get_estimates	<i>Extract model estimates. A mostly internal function with some potential uses outside.</i>
---------------	--

Description

Extract model estimates. A mostly internal function with some potential uses outside.

Usage

```
get_estimates(model, conf_level = 0.95, vcov = NULL, ...)
```

Arguments

model	a single model object
conf_level	confidence level to use for confidence intervals
vcov	robust standard errors and other manual statistics. The <code>vcov</code> argument accepts five types of input (see the 'Details' and 'Examples' sections below): <ul style="list-style-type: none"> • string, vector, or list of strings: "robust", "HC", "HC0", "HC1", "HC2", "HC3", "HC4", "HC4m", "HC5", "stata", or "classical" (alias "constant" or "iid"). • formula or list of formulas with the cluster variable(s) on the right-hand side (e.g., <code>~clusterid</code>). • function or list of functions which return variance-covariance matrices with row and column names equal to the names of your coefficient estimates (e.g., <code>stats::vcov</code>, <code>sandwich::vcovHC</code>). • list of <code>length(models)</code> variance-covariance matrices with row and column names equal to the names of your coefficient estimates.

- a list of `length(models)` vectors with names equal to the names of your coefficient estimates. See 'Examples' section below. Warning: since this list of vectors can include arbitrary strings or numbers, `modelsummary` cannot automatically calculate p values. The `stars` argument may thus use incorrect significance thresholds when `vcov` is a list of vectors.
- ...
- all other arguments are passed through to the extractor and table-making functions. This allows users to pass arguments directly to `modelsummary` in order to affect the behavior of other functions behind the scenes. Examples include:
- `broom::tidy(exponentiate=TRUE)` to exponentiate logistic regression
 - `kableExtra::kbl(escape=FALSE)` to avoid escaping math characters in `kableExtra` tables.
 - `performance::model_performance(metrics="RMSE")` to select goodness-of-fit statistics to extract using the `performance` package (must have set `options(modelsummary_get="easystats")` first).

get_gof

Extract model gof A mostly internal function with some potential uses outside.

Description

Extract model gof A mostly internal function with some potential uses outside.

Usage

```
get_gof(model, vcov_type = NULL, ...)
```

Arguments

`model` a single model object

`vcov_type` string `vcov` type to add at the bottom of the table

- ...
- all other arguments are passed through to the extractor and table-making functions. This allows users to pass arguments directly to `modelsummary` in order to affect the behavior of other functions behind the scenes. Examples include:
- `broom::tidy(exponentiate=TRUE)` to exponentiate logistic regression
 - `kableExtra::kbl(escape=FALSE)` to avoid escaping math characters in `kableExtra` tables.
 - `performance::model_performance(metrics="RMSE")` to select goodness-of-fit statistics to extract using the `performance` package (must have set `options(modelsummary_get="easystats")` first).

glance.modelsummary_list

Extract goodness-of-fit statistics from a modelsummary_list object.

Description

A modelsummary_list is a simple list which holds two data.frames names "glance" and "tidy", and to which we have attached the class "modelsummary_list". This allows us to manually create objects that will be printed nicely by modelsummary (see Examples section).

Usage

```
## S3 method for class 'modelsummary_list'
glance(x, ...)
```

Arguments

x a list of class modelsummary_list (see Examples section)
 ... other parameters are accepted by ignored

Examples

```
## Not run:
mod <- list(
  tidy = data.frame(term = c("A", "B"),
                    estimate = 1:2,
                    std.error = 3:4),
  glance = data.frame(nobs = 10))
class(mod) <- c("modelsummary_list", class(mod))

modelsummary(mod)

## End(Not run)
```

glance_custom

Extract custom information from a model object and turn it into a tidy data.frame or tibble with a single row.

Description

To customize the output of a model of class lm, you can define a new method called glance_custom.lm which returns a one-row data.frame.

Usage

```
glance_custom(x)
```

Arguments

x model or other R object to convert to single-row data frame

Methods

No methods found in currently loaded packages.

gof_map	<i>Data.frame used to clean up and format goodness-of-fit statistics</i>
---------	--

Description

By default, this data frame is passed to the 'gof_map' argument of the 'modelsummary' function. Users can modify this data frame to customize the list of statistics to display and their format. See example below.

Usage

```
gof_map
```

Format

data.frame with 4 columns of character data: raw, clean, fmt, omit

Examples

```
## Not run:

library(modelsummary)
mod <- lm(wt ~ drat, data = mtcars)
gm <- modelsummary::gof_map
gm$omit[gm$raw == 'deviance'] <- FALSE
gm$fmt[gm$raw == 'r.squared'] <- "%.5f"
modelsummary(mod, gof_map = gm)

## End(Not run)
```

Histogram	<i>datasummary statistic shortcut</i>
-----------	---------------------------------------

Description

This function uses Unicode characters to create a histogram. This can sometimes be useful, but is generally discouraged. Unicode characters can only display a limited number of heights for bars, and the accuracy of output is highly dependent on the platform (typeface, output type, windows vs. mac, etc.). We recommend you use the `kableExtra::spec_hist` function instead.

Usage

```
Histogram(x, bins = 10)
```

Arguments

x	variable to summarize
bins	number of histogram bars

Max	<i>datasummary statistic shortcut</i>
-----	---------------------------------------

Description

datasummary statistic shortcut

Usage

```
Max(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary:::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
             Min + Max + SD + Var,
             data = mtcars)

## End(Not run)
```

Mean	<i>datasummary statistic shortcut</i>
------	---------------------------------------

Description

datasummary statistic shortcut

Usage

```
Mean(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary:::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + P0 + P25 + P50 + P75 + P100 +
            Min + Max + SD + Var,
            data = mtcars)

## End(Not run)
```

Median	<i>datasummary statistic shortcut</i>
--------	---------------------------------------

Description

datasummary statistic shortcut

Usage

```
Median(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
            Min + Max + SD + Var,
            data = mtcars)

## End(Not run)
```

Min	<i>datasummary statistic shortcut</i>
-----	---------------------------------------

Description

datasummary statistic shortcut

Usage

```
Min(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
             Min + Max + SD + Var,
             data = mtcars)

## End(Not run)
```

modelplot

Model Summary Plots with Estimates and Confidence Intervals

Description

Model Summary Plots with Estimates and Confidence Intervals

Usage

```
modelplot(
  models,
  conf_level = 0.95,
  coef_map = NULL,
  coef_omit = NULL,
  coef_rename = NULL,
  vcov = NULL,
  add_rows = NULL,
  facet = FALSE,
  draw = TRUE,
  background = NULL,
  ...
)
```

Arguments

models	a model or (optionally named) list of models
conf_level	confidence level to use for confidence intervals
coef_map	character vector. Subset, rename, and reorder coefficients. Coefficients omitted from this vector are omitted from the table. The order of the vector determines the order of the table. <code>coef_map</code> can be a named or an unnamed character vector (see the Examples section below). If <code>coef_map</code> is a named vector, its values define the labels that must appear in the table, and its names identify the original term names stored in the model object: <code>c("hp:mpg"="HPxM/G")</code> .

coef_omit	string regular expression. Omits all matching coefficients from the table using <code>grep1(per1=TRUE)</code> .
coef_rename	named character vector or function which returns a named vector. Values of the vector refer to the variable names that will appear in the table. Names refer to the original term names stored in the model object, e.g. <code>c("hp:mpg"="hp X mpg")</code> for an interaction term.
vcov	robust standard errors and other manual statistics. The <code>vcov</code> argument accepts five types of input (see the 'Details' and 'Examples' sections below): <ul style="list-style-type: none"> • string, vector, or list of strings: "robust", "HC", "HC0", "HC1", "HC2", "HC3", "HC4", "HC4m", "HC5", "stata", or "classical" (alias "constant" or "iid"). • formula or list of formulas with the cluster variable(s) on the right-hand side (e.g., <code>~clusterid</code>). • function or list of functions which return variance-covariance matrices with row and column names equal to the names of your coefficient estimates (e.g., <code>stats::vcov</code>, <code>sandwich::vcovHC</code>). • list of <code>length(models)</code> variance-covariance matrices with row and column names equal to the names of your coefficient estimates. • a list of <code>length(models)</code> vectors with names equal to the names of your coefficient estimates. See 'Examples' section below. Warning: since this list of vectors can include arbitrary strings or numbers, <code>modelsummary</code> cannot automatically calculate p values. The <code>stars</code> argument may thus use incorrect significance thresholds when <code>vcov</code> is a list of vectors.
add_rows	a <code>data.frame</code> (or <code>tibble</code>) with the same number of columns as your main table. By default, rows are appended to the bottom of the table. You can define a "position" attribute of integers to set the row positions. See Examples section below.
facet	TRUE or FALSE. When the 'models' argument includes several model objects, TRUE draws terms in separate facets, and FALSE draws terms side-by-side (dodged).
draw	TRUE returns a 'ggplot2' object, FALSE returns the <code>data.frame</code> used to draw the plot.
background	A list of 'ggplot2' geoms to add to the background of the plot. This is especially useful to display annotations "behind" the 'geom_pointrange' that 'modelplot' draws.
...	all other arguments are passed through to the extractor and table-making functions. This allows users to pass arguments directly to <code>modelsummary</code> in order to affect the behavior of other functions behind the scenes. Examples include: <ul style="list-style-type: none"> • <code>broom::tidy(exponentiate=TRUE)</code> to exponentiate logistic regression • <code>kableExtra::kbl(escape=FALSE)</code> to avoid escaping math characters in <code>kableExtra</code> tables. • <code>performance::model_performance(metrics="RMSE")</code> to select goodness-of-fit statistics to extract using the <code>performance</code> package (must have set <code>options(modelsummary_get="easystats")</code> first).

Examples

```
## Not run:
```

```

library(modelsummary)

# single model
mod <- lm(hp ~ vs + drat, mtcars)
modelplot(mod)

# omit terms with string matches or regexes
modelplot(mod, coef_omit = 'Interc')

# rename, reorder and subset with 'coef_map'
cm <- c('vs' = 'V-shape engine',
        'drat' = 'Rear axle ratio')
modelplot(mod, coef_map = cm)

# several models
models <- list()
models[['Small model']] <- lm(hp ~ vs, mtcars)
models[['Medium model']] <- lm(hp ~ vs + factor(cyl), mtcars)
models[['Large model']] <- lm(hp ~ vs + drat + factor(cyl), mtcars)
modelplot(models)

# add_rows: add an empty reference category
mod <- lm(hp ~ factor(cyl), mtcars)

add_rows = data.frame(
  term = "factory(cyl)4",
  model = "Model 1",
  estimate = NA)
attr(add_rows, "position") = 3
modelplot(mod, add_rows = add_rows)

# customize your plots with 'ggplot2' functions
library(ggplot2)

modelplot(models) +
  scale_color_brewer(type = 'qual') +
  theme_classic()

# pass arguments to 'geom_pointrange' through the ... ellipsis
modelplot(mod, color = 'red', size = 1, fatten = .5)

# add geoms to the background, behind geom_pointrange
b <- list(geom_vline(xintercept = 0, color = 'orange'),
  annotate("rect", alpha = .1,
    xmin = -.5, xmax = .5,
    ymin = -Inf, ymax = Inf),
  geom_point(aes(y = term, x = estimate), alpha = .3,
    size = 10, color = 'red', shape = 'square'))
modelplot(mod, background = b)

## End(Not run)

```

Description

The content of the tables can be altered with the function's arguments, or by calling options, as described in the *Details* section below. The look of the tables can be customized by specifying the output argument, and by using functions from one of the supported table customization packages: `kableExtra`, `gt`, `flextable`, `huxtable`.

Usage

```
modelssummary(
  models,
  output = "default",
  fmt = 3,
  estimate = "estimate",
  statistic = "std.error",
  vcov = NULL,
  conf_level = 0.95,
  stars = FALSE,
  coef_map = NULL,
  coef_omit = NULL,
  coef_rename = NULL,
  gof_map = NULL,
  gof_omit = NULL,
  group = term ~ model,
  group_map = NULL,
  add_rows = NULL,
  align = NULL,
  notes = NULL,
  title = NULL,
  ...
)
```

Arguments

<code>models</code>	a model or (optionally named) list of models
<code>output</code>	filename or object type (character string) <ul style="list-style-type: none"> Supported filename extensions: <code>.html</code>, <code>.tex</code>, <code>.md</code>, <code>.txt</code>, <code>.png</code>, <code>.jpg</code>. Supported object types: <code>"default"</code>, <code>"html"</code>, <code>"markdown"</code>, <code>"latex"</code>, <code>"latex_tabular"</code>, <code>"data.frame"</code>, <code>"modelssummary_list"</code>, <code>"gt"</code>, <code>"kableExtra"</code>, <code>"huxtable"</code>, <code>"flextable"</code>. To change the default output format, type <code>options(modelssummary_default = "latex")</code>, where <code>latex</code> can be any of the valid object types listed above. Warning: the output argument <i>cannot</i> be used when customizing tables with external packages. See the 'Details' section below for more information.
<code>fmt</code>	determines how to format numeric values <ul style="list-style-type: none"> integer: the number of digits to keep after the period <code>format(round(x, fmt), nsmall=fmt)</code>

	<ul style="list-style-type: none"> • character: passed to the <code>sprintf</code> function (e.g., <code>`\</code> • function: returns a formatted character string.
estimate	<p>string or glue string of the estimate to display (or a vector with one string per model). Valid entries include any column name of the data.frame produced by <code>get_estimates(model)</code>. Examples:</p> <ul style="list-style-type: none"> • <code>"estimate"</code> • <code>"{estimate} ({std.error}){stars}"</code> • <code>"{estimate} [{conf.low}, {conf.high}]"</code>
statistic	<p>vector of strings or glue strings which select uncertainty statistics to report vertically below the estimate. NULL omits all uncertainty statistics.</p> <ul style="list-style-type: none"> • <code>"conf.int", "std.error", "statistic", "p.value", "conf.low", "conf.high"</code>, or any column name produced by: <code>get_estimates(model)</code> • glue package strings with braces, such as: <ul style="list-style-type: none"> - <code>"{p.value} [{conf.low}, {conf.high}]"</code> - <code>"Std.Error: {std.error}"</code> • Note: Parentheses are added automatically unless the string includes glue curly braces <code>{}</code>. • Note: To report uncertainty statistics <i>next</i> to coefficients, you can <code>#'</code> supply a glue string to the <code>estimate</code> argument.
vcov	<p>robust standard errors and other manual statistics. The <code>vcov</code> argument accepts five types of input (see the 'Details' and 'Examples' sections below):</p> <ul style="list-style-type: none"> • string, vector, or list of strings: <code>"robust", "HC", "HC0", "HC1", "HC2", "HC3", "HC4", "HC4m", "HC5", "stata"</code>, or <code>"classical"</code> (alias <code>"constant"</code> or <code>"iid"</code>). • formula or list of formulas with the cluster variable(s) on the right-hand side (e.g., <code>~clusterid</code>). • function or list of functions which return variance-covariance matrices with row and column names equal to the names of your coefficient estimates (e.g., <code>stats::vcov</code>, <code>sandwich::vcovHC</code>). • list of <code>length(models)</code> variance-covariance matrices with row and column names equal to the names of your coefficient estimates. • a list of <code>length(models)</code> vectors with names equal to the names of your coefficient estimates. See 'Examples' section below. Warning: since this list of vectors can include arbitrary strings or numbers, <code>modelsummary</code> cannot automatically calculate p values. The <code>stars</code> argument may thus use incorrect significance thresholds when <code>vcov</code> is a list of vectors.
conf_level	confidence level to use for confidence intervals
stars	<p>to indicate statistical significance</p> <ul style="list-style-type: none"> • FALSE (default): no significance stars. • TRUE: <code>*=.1, **=.05, ***=.01</code> • Named numeric vector for custom stars such as <code>c('*' = .1, '+' = .05)</code> • Note: a legend will not be inserted at the bottom of the table when the <code>estimate</code> or <code>statistic</code> arguments use "glue strings" with <code>{stars}</code>.
coef_map	<p>character vector. Subset, rename, and reorder coefficients. Coefficients omitted from this vector are omitted from the table. The order of the vector determines the order of the table. <code>coef_map</code> can be a named or an unnamed character vector (see the Examples section below). If <code>coef_map</code> is a named vector, its values define the labels that must appear in the table, and its names identify the original term names stored in the model object: <code>c("hp:mpg"="HPxM/G")</code>.</p>

<code>coef_omit</code>	string regular expression. Omits all matching coefficients from the table using <code>grep1(per1=TRUE)</code> .
<code>coef_rename</code>	named character vector or function which returns a named vector. Values of the vector refer to the variable names that will appear in the table. Names refer to the original term names stored in the model object, e.g. <code>c("hp:mpg"="hp X mpg")</code> for an interaction term.
<code>gof_map</code>	<ul style="list-style-type: none"> • <code>NULL</code> (default): the <code>modelsummary::gof_map</code> dictionary is used for formatting, and all unknown statistic are included. • <code>data.frame</code> with 3 columns named "raw", "clean", "fmt". Unknown statistics are omitted. See the 'Examples' section below. • list of lists, each of which includes 3 elements named "raw", "clean", "fmt". Unknown statistics are omitted. See the 'Examples section below'.
<code>gof_omit</code>	string regular expression. Omits all matching gof statistics from the table (using <code>grep1(per1=TRUE)</code>).
<code>group</code>	<p>a two-sided formula with two or three components which describes how groups of parameters should be displayed. The formula must include both a "term" and a "model" component. In addition, a component can be used to identify groups of parameters (e.g., outcome levels of a multinomial logit model). This group identifier must be the name of a column in the <code>data.frame</code> produced by <code>get_estimates(model)</code>.</p> <ul style="list-style-type: none"> • <code>term ~ model</code> displays coefficients as rows and models as columns • <code>model ~ term</code> displays models as rows and coefficients as columns • <code>response + term ~ model</code> displays response levels and coefficients as rows and models as columns.
<code>group_map</code>	named or unnamed character vector. Subset, rename, and reorder coefficient groups specified in the <code>group</code> argument. See <code>coef_map</code> .
<code>add_rows</code>	a <code>data.frame</code> (or <code>tibble</code>) with the same number of columns as your main table. By default, rows are appended to the bottom of the table. You can define a "position" attribute of integers to set the row positions. See Examples section below.
<code>align</code>	A character string of length equal to the number of columns in the table. "lcr" means that the first column will be left-aligned, the 2nd column center-aligned, and the 3rd column right-aligned.
<code>notes</code>	list or vector of notes to append to the bottom of the table.
<code>title</code>	string
<code>...</code>	<p>all other arguments are passed through to the extractor and table-making functions. This allows users to pass arguments directly to <code>modelsummary</code> in order to affect the behavior of other functions behind the scenes. Examples include:</p> <ul style="list-style-type: none"> • <code>broom::tidy(exponentiate=TRUE)</code> to exponentiate logistic regression • <code>kableExtra::kbl(escape=FALSE)</code> to avoid escaping math characters in <code>kableExtra</code> tables. • <code>performance::model_performance(metrics="RMSE")</code> to select goodness-of-fit statistics to extract using the <code>performance</code> package (must have set <code>options(modelsummary_get="easystats")</code> first).

Details

`options`

modelsummary supports 4 table-making packages: kableExtra, gt, flextable, and huxtable. Some of these packages have overlapping functionalities. For example, 3 of those packages can export to LaTeX. To change the default backend used for a specific file format, you can use the options function:

```
options(modelsummary_html = 'kableExtra') options(modelsummary_latex = 'gt') options(modelsummary_w
= 'huxtable') options(modelsummary_png = 'gt')
```

modelsummary can use two sets of packages to extract information from statistical models: broom and the easystats family (performance and parameters). By default, it uses broom first and easystats as a fallback if broom fails. You can change the order of priorities or include goodness-of-fit extracted by *both* packages by setting:

```
options(modelsummary_get = "broom") options(modelsummary_get = "easystats") options(modelsummary_ge
= "all")
```

output argument:

The modelsummary_list output type is a lightweight representation of the model results. The modelsummary function can export to this format by setting the output argument, and it can accept objects of this format as input models to create a table. This can be useful to save raw results, in order to print a table later, without having to save and extract from the entire model object. Note that the confidence intervals are only stored in a modelsummary_list if explicitly requested:

```
backup <- modelsummary(models, output = "modelsummary_list"          statistic = "conf.int")
modelsummary(backup)
```

When a file name with a valid extension is supplied to the output argument, the table is written immediately to file. If you want to customize your table by post-processing it with an external package, you need to choose a different output format and saving mechanism. Unfortunately, the approach differs from package to package:

- `gt`: set `output="gt"`, post-process your table, and use the `gt::gtsave` function.
- `kableExtra`: set output to your destination format (e.g., "latex", "html", "markdown"), post-process your table, and use `kableExtra::save_kable` function.

vcov argument:

To use a string such as "robust" or "HCO", your model must be supported by the sandwich package. This includes objects such as: lm, glm, survreg, coxph, mlogit, polr, hurdle, zeroinfl, and more.

"classical", "iid", and "constant" are aliases which do not modify uncertainty estimates and simply report the default standard errors stored in the model object.

One-sided formulas such as `~clusterid` are passed to the `sandwich::vcovCL` function.

Matrices and functions producing variance-covariance matrices are first passed to `lmtest`. If this does not work, `modelsummary` attempts to take the square root of the diagonal to adjust "std.error", but the other uncertainty estimates are not be adjusted.

Numeric vectors are formatted according to `fmt` and placed in brackets. Character vectors printed as given, without parentheses.

If your model type is supported by the `lmtest` package, the `vcov` argument will try to use that package to adjust all the uncertainty estimates, including "std.error", "statistic", "p.value", and "conf.int". If your model is not supported by `lmtest`, only the "std.error" will be adjusted by, for example, taking the square root of the matrix's diagonal.

Value

a regression table in a format determined by the output argument.

Examples

```

## Not run:

# The `modelssummary` website includes \emph{many} examples and tutorials:
# https://vincentarelbundock.github.io/modelssummary

library(modelssummary)

# load data and estimate models
data(trees)
models <- list()
models[['Bivariate']] <- lm(Girth ~ Height, data = trees)
models[['Multivariate']] <- lm(Girth ~ Height + Volume, data = trees)

# simple table
modelssummary(models)

# statistic
modelssummary(models, statistic = NULL)
modelssummary(models, statistic = 'p.value')
modelssummary(models, statistic = 'statistic')
modelssummary(models, statistic = 'conf.int', conf_level = 0.99)
modelssummary(models, statistic = c("t = {statistic}",
                                     "se = {std.error}",
                                     "conf.int"))

# estimate
modelssummary(models,
  statistic = NULL,
  estimate = "{estimate} [{conf.low}, {conf.high}]")
modelssummary(models,
  estimate = c("{estimate}{stars}",
               "{estimate} ({std.error})"))

# vcov
modelssummary(models, vcov = "robust")
modelssummary(models, vcov = list("classical", "stata"))
modelssummary(models, vcov = sandwich::vcovHC)
modelssummary(models,
  vcov = list(stats::vcov, sandwich::vcovHC))
modelssummary(models,
  vcov = list(c("(Intercept)"="!", "Height"="!"),
              c("(Intercept)"="!", "Height"="!", "Volume"="!!")))

# coef_rename
modelssummary(models, coef_map = c('Volume' = 'Large', 'Height' = 'Tall'))

# coef_map
modelssummary(models, coef_map = c('Volume' = 'Large', 'Height' = 'Tall'))
modelssummary(models, coef_map = c('Volume', 'Height'))

# title
modelssummary(models, title = 'This is the title')

# add_rows
rows <- tibble::tribble(~term, ~Bivariate, ~Multivariate,

```

```

      'Empty row', '-', '-',
      'Another empty row', '?', '?')
attr(rows, 'position') <- c(1, 3)
modelsummary(models, add_rows = rows)

# notes
modelsummary(models, notes = list('A first note', 'A second note'))

# gof_map: data.frame
gm <- modelsummary::gof_map
gof_custom$omit[gof_custom$raw == 'deviance'] <- FALSE
gof_custom$fmt[gof_custom$raw == 'r.squared'] <- "%.5f"
modelsummary(models, gof_map = gof_custom)

# gof_map: list of lists
f1 <- function(x) format(round(x, 3), big.mark=",")
f2 <- function(x) format(round(x, 0), big.mark=",")
gm <- list(
  list("raw" = "nobs", "clean" = "N", "fmt" = f2),
  list("raw" = "AIC", "clean" = "aic", "fmt" = f1))
modelsummary(models,
  fmt = f1,
  gof_map = gm)

## End(Not run)

```

modelsummary_wide

Summary Tables for Models with Grouped Coefficients

Description

modelsummary_wide summarizes models with grouped coefficients. For example, these groups could correspond to levels of a multinomial logit outcome variable, or to parameters of a GAMLSS model. This function's arguments are the same as in modelsummary, except for the coef_group and the stacking arguments.

Usage

```

modelsummary_wide(
  models,
  output = "default",
  fmt = 3,
  estimate = "estimate",
  statistic = "std.error",
  vcov = NULL,
  conf_level = 0.95,
  stars = FALSE,
  coef_group = NULL,
  coef_map = NULL,
  coef_omit = NULL,
  coef_rename = NULL,

```

```

gof_map = NULL,
gof_omit = NULL,
add_rows = NULL,
align = NULL,
notes = NULL,
title = NULL,
stacking = "horizontal",
...
)

```

Arguments

models	a model or (optionally named) list of models
output	filename or object type (character string) <ul style="list-style-type: none"> Supported filename extensions: .html, .tex, .md, .txt, .png, .jpg. Supported object types: "default", "html", "markdown", "latex", "latex_tabular", "data.frame", "modelssummary_list", "gt", "kableExtra", "huxtable", "flextable". To change the default output format, type <code>options(modelssummary_default = "latex")</code>, where <code>latex</code> can be any of the valid object types listed above. Warning: the output argument <i>cannot</i> be used when customizing tables with external packages. See the 'Details' section below for more information.
fmt	determines how to format numeric values <ul style="list-style-type: none"> integer: the number of digits to keep after the period <code>format(round(x, fmt), nsmall=fmt)</code> character: passed to the <code>sprintf</code> function (e.g., <code>\</code>) function: returns a formatted character string.
estimate	string or glue string of the estimate to display (or a vector with one string per model). Valid entries include any column name of the data.frame produced by <code>get_estimates(model)</code> . Examples: <ul style="list-style-type: none"> "estimate" "{estimate} ({std.error}){stars}" "{estimate} [{conf.low}, {conf.high}]"
statistic	vector of strings or glue strings which select uncertainty statistics to report vertically below the estimate. NULL omits all uncertainty statistics. <ul style="list-style-type: none"> "conf.int", "std.error", "statistic", "p.value", "conf.low", "conf.high", or any column name produced by: <code>get_estimates(model)</code> glue package strings with braces, such as: <ul style="list-style-type: none"> "{p.value} [{conf.low}, {conf.high}]" "Std.Error: {std.error}" Note: Parentheses are added automatically unless the string includes glue curly braces <code>{}</code>. Note: To report uncertainty statistics <i>next</i> to coefficients, you can <code>#</code> supply a glue string to the estimate argument.
vcov	robust standard errors and other manual statistics. The <code>vcov</code> argument accepts five types of input (see the 'Details' and 'Examples' sections below): <ul style="list-style-type: none"> string, vector, or list of strings: "robust", "HC", "HC0", "HC1", "HC2", "HC3", "HC4", "HC4m", "HC5", "stata", or "classical" (alias "constant" or "iid").

- formula or list of formulas with the cluster variable(s) on the right-hand side (e.g., ~clusterid).
- function or list of functions which return variance-covariance matrices with row and column names equal to the names of your coefficient estimates (e.g., stats::vcov, sandwich::vcovHC).
- list of length(models) variance-covariance matrices with row and column names equal to the names of your coefficient estimates.
- a list of length(models) vectors with names equal to the names of your coefficient estimates. See 'Examples' section below. Warning: since this list of vectors can include arbitrary strings or numbers, modelsummary cannot automatically calculate p values. The stars argument may thus use incorrect significance thresholds when vcov is a list of vectors.

conf_level confidence level to use for confidence intervals

stars to indicate statistical significance

- FALSE (default): no significance stars.
- TRUE: *=.1, **=.05, ***=.01
- Named numeric vector for custom stars such as c('*' = .1, '+' = .05)
- Note: a legend will not be inserted at the bottom of the table when the estimate or statistic arguments use "glue strings" with {stars}.

coef_group the name of the coefficient groups to use as columns (NULL or character). If coef_group is NULL, modelsummary tries to guess the correct coefficient group identifier. To be valid, this identifier must be a column in the data.frame produced by get_estimates(model).

coef_map character vector. Subset, rename, and reorder coefficients. Coefficients omitted from this vector are omitted from the table. The order of the vector determines the order of the table. coef_map can be a named or an unnamed character vector (see the Examples section below). If coef_map is a named vector, its values define the labels that must appear in the table, and its names identify the original term names stored in the model object: c("hp:mpg"="HPxM/G").

coef_omit string regular expression. Omits all matching coefficients from the table using grepl(perl=TRUE).

coef_rename named character vector or function which returns a named vector. Values of the vector refer to the variable names that will appear in the table. Names refer to the original term names stored in the model object, e.g. c("hp:mpg"="hp X mpg") for an interaction term.

gof_map

- NULL (default): the modelsummary::gof_map dictionary is used for formatting, and all unknown statistic are included.
- data.frame with 3 columns named "raw", "clean", "fmt". Unknown statistics are omitted. See the 'Examples' section below.
- list of lists, each of which includes 3 elements named "raw", "clean", "fmt". Unknown statistics are omitted. See the 'Examples section below'.

gof_omit string regular expression. Omits all matching gof statistics from the table (using grepl(perl=TRUE)).

add_rows a data.frame (or tibble) with the same number of columns as your main table. By default, rows are appended to the bottom of the table. You can define a "position" attribute of integers to set the row positions. See Examples section below.

<code>align</code>	A character string of length equal to the number of columns in the table. "lcr" means that the first column will be left-aligned, the 2nd column center-aligned, and the 3rd column right-aligned.
<code>notes</code>	list or vector of notes to append to the bottom of the table.
<code>title</code>	string
<code>stacking</code>	direction in which models are stacked: "horizontal" or "vertical"
<code>...</code>	all other arguments are passed through to the extractor and table-making functions. This allows users to pass arguments directly to <code>modelsummary</code> in order to affect the behavior of other functions behind the scenes. Examples include: <ul style="list-style-type: none"> • <code>broom::tidy(exponentiate=TRUE)</code> to exponentiate logistic regression • <code>kableExtra::kbl(escape=FALSE)</code> to avoid escaping math characters in <code>kableExtra</code> tables. • <code>performance::model_performance(metrics="RMSE")</code> to select goodness-of-fit statistics to extract using the performance package (must have set <code>options(modelsummary_get="easystats")</code> first).

Value

a regression table in a format determined by the output argument.

N	<i>datasummary statistic shortcut</i>
---	---------------------------------------

Description

`datasummary statistic shortcut`

Usage

`N(x)`

Arguments

`x` variable to summarize

Examples

```
## Not run:
datasummary(Factor(cyl) ~ N, data = mtcars)

## End(Not run)
```

Ncol	<i>datasummary statistic shortcut</i>
------	---------------------------------------

Description

datasummary statistic shortcut

Usage

Ncol(x, ...)

Arguments

x	variable to summarize
...	unused

NPercent	<i>datasummary statistic shortcut</i>
----------	---------------------------------------

Description

datasummary statistic shortcut

Usage

NPercent(x, y)

Arguments

x	variable to summarize
y	denominator variable

NUnique	<i>datasummary statistic shortcut</i>
---------	---------------------------------------

Description

datasummary statistic shortcut

Usage

NUnique(x, ...)

Arguments

x	variable to summarize
...	unused

Examples

```
## Not run:
datasummary(cyl + hp ~ NUnique, data = mtcars)

## End(Not run)
```

P0	<i>datasummary statistic shortcut</i>
----	---------------------------------------

Description

datasummary statistic shortcut

Usage

```
P0(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
            Min + Max + SD + Var,
            data = mtcars)

## End(Not run)
```

P100	<i>datasummary statistic shortcut</i>
------	---------------------------------------

Description

datasummary statistic shortcut

Usage

```
P100(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
            Min + Max + SD + Var,
            data = mtcars)

## End(Not run)
```

P25

datasummary statistic shortcut

Description

datasummary statistic shortcut

Usage

```
P25(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
            Min + Max + SD + Var,
            data = mtcars)

## End(Not run)
```

P50 *datasummary statistic shortcut*

Description

datasummary statistic shortcut

Usage

```
P50(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary:::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
            Min + Max + SD + Var,
            data = mtcars)

## End(Not run)
```

P75 *datasummary statistic shortcut*

Description

datasummary statistic shortcut

Usage

```
P75(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary:::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
            Min + Max + SD + Var,
            data = mtcars)

## End(Not run)
```

PercentMissing	<i>datasummary statistic shortcut</i>
----------------	---------------------------------------

Description

datasummary statistic shortcut

Usage

```
PercentMissing(x)
```

Arguments

x	variable to summarize
---	-----------------------

SD	<i>datasummary statistic shortcut</i>
----	---------------------------------------

Description

datasummary statistic shortcut

Usage

```
SD(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

x	variable to summarize
fmt	passed to the <code>modelsummary::rounding</code> function
na.rm	a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
...	unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
            Min + Max + SD + Var,
            data = mtcars)

## End(Not run)
```

supported_models	<i>List of model objects from which modelsummary can extract estimates and statistics</i>
------------------	---

Description

List of model objects from which modelsummary can extract estimates and statistics

Usage

```
supported_models()
```

tidy.modelsummary_list	<i>Extract parameter estimates from a modelsummary_list object.</i>
------------------------	---

Description

A modelsummary_list is a simple list which holds two data.frames names "glance" and "tidy", and to which we have attached the class "modelsummary_list". This allows us to manually create objects that will be printed nicely by modelsummary (see Examples section).

Usage

```
## S3 method for class 'modelsummary_list'
tidy(x, ...)
```

Arguments

x	a list of class modelsummary_list (see Examples section)
...	other parameters are accepted by ignored

Examples

```
## Not run:
mod <- list(
  tidy = data.frame(term = c("A", "B"),
                    estimate = 1:2,
                    std.error = 3:4),
  glance = data.frame(nobs = 10))
class(mod) <- c("modelsummary_list", class(mod))

modelsummary(mod)

## End(Not run)
```

tidy_custom	<i>Extract custom information from a model object and turn it into a tidy data.frame or tibble</i>
-------------	--

Description

To customize the output of a model of class `lm`, you can define a method called `tidy_custom.lm` which returns a `data.frame` with a column called "term", and the other columns you want to use as "estimate" or "statistic" in your `modelsummary()` call. The output of this method must be similar to the result of `tidy(model)`.

Usage

```
tidy_custom(x)
```

Arguments

`x` An object to be converted into a tidy `data.frame` or `tibble`.

Value

A `data.frame` or `tibble` with information about model components.

Var	<i>datasummary statistic shortcut</i>
-----	---------------------------------------

Description

`datasummary statistic shortcut`

Usage

```
Var(x, fmt = NULL, na.rm = TRUE, ...)
```

Arguments

`x` variable to summarize
`fmt` passed to the `modelsummary::rounding` function
`na.rm` a logical value indicating whether 'NA' values should be stripped before the computation proceeds.
`...` unused

Examples

```
## Not run:
datasummary(mpg + hp ~ Mean + Median + P0 + P25 + P50 + P75 + P100 +
  Min + Max + SD + Var,
  data = mtcars)

## End(Not run)
```

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