

# Package: gtsummary (via r-universe)

October 21, 2024

**Title** Presentation-Ready Data Summary and Analytic Result Tables

**Version** 2.0.3.9003

**Description** Creates presentation-ready tables summarizing data sets, regression models, and more. The code to create the tables is concise and highly customizable. Data frames can be summarized with any function, e.g. `mean()`, `median()`, even user-written functions. Regression models are summarized and include the reference rows for categorical variables. Common regression models, such as logistic regression and Cox proportional hazards regression, are automatically identified and the tables are pre-filled with appropriate column headers.

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**URL** <https://github.com/ddsjoberg/gtsummary>,  
<https://www.danieldsjoberg.com/gtsummary/>

**BugReports** <https://github.com/ddsjoberg/gtsummary/issues>

**Depends** R (>= 4.2)

**Imports** cards (>= 0.3.0), cli (>= 3.6.3), dplyr (>= 1.1.3), glue (>= 1.8.0), gt (>= 0.10.0), lifecycle (>= 1.0.3), rlang (>= 1.1.1), tidyr (>= 1.3.0), vctrs (>= 0.6.4)

**Suggests** aod (>= 1.3.3), broom (>= 1.0.5), broom.helpers (>= 1.17.0), broom.mixed (>= 0.2.9), car (>= 3.0-11), cardx (>= 0.2.1), cmprsk, effectsize (>= 0.6.0), emmeans (>= 1.7.3), flextable (>= 0.8.1), geopack (>= 1.3.10), ggstats (>= 0.2.1), huxtable (>= 5.4.0), insight (>= 0.15.0), kableExtra (>= 1.3.4), knitr (>= 1.37), lme4 (>= 1.1-31), mice (>= 3.10.0), nnet, officer, openxlsx, parameters (>= 0.20.2), parsnip (>= 0.1.7), rmarkdown, smd (>= 0.6.6), spelling, survey (>= 4.2), survival (>= 3.6-4), testthat (>= 3.2.0), withr (>= 2.5.0), workflows (>= 0.2.4)

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---

add_ci	<i>Add CI Column</i>
--------	----------------------

---

## Description

Add a new column with the confidence intervals for proportions, means, etc.

## Usage

```
add_ci(x, ...)

## S3 method for class 'tbl_summary'
add_ci(
  x,
  method = list(all_continuous() ~ "t.test", all_categorical() ~ "wilson"),
  include = everything(),
  statistic = list(all_continuous() ~ "{conf.low}, {conf.high}", all_categorical() ~
    "{conf.low}%, {conf.high}%"),
  conf.level = 0.95,
  style_fun = list(all_continuous() ~ label_style_sigfig(), all_categorical() ~
    label_style_sigfig(scale = 100)),
  pattern = NULL,
  ...
)
```

**Arguments**

x	(tbl_summary) a summary table of class 'tblsummary'
...	These dots are for future extensions and must be empty.
method	( <a href="#">formula-list-selector</a> ) Confidence interval method. Default is <code>list(all_continuous() ~ "t.test", all_categorical() ~ "wilson")</code> . See details below.
include	( <a href="#">tidy-select</a> ) Variables to include in the summary table. Default is <code>everything()</code> .
statistic	( <a href="#">formula-list-selector</a> ) Indicates how the confidence interval will be displayed. Default is <code>list(all_continuous() ~ "{conf.low}, {conf.high}", all_categorical() ~ "{conf.low}%, {conf.high}%")</code>
conf.level	(scalar real) Confidence level. Default is 0.95
style_fun	(function) Function to style upper and lower bound of confidence interval. Default is <code>list(all_continuous() ~ label_style_sigfig(), all_categorical() ~ label_style_sigfig(s = 100))</code> .
pattern	(string) Indicates the pattern to use to merge the CI with the statistics cell. The default is NULL, where no columns are merged. The two columns that will be merged are the statistics column, represented by "{stat}" and the CI column represented by "{ci}", e.g. <code>pattern = "{stat} ({ci})"</code> will merge the two columns with the CI in parentheses. Default is NULL, and no merging is performed.

**Value**

gtsummary table

**method argument**

Must be one of

- "wilson", "wilson.no.correct" calculated via `prop.test(correct = c(TRUE, FALSE))` for **categorical** variables
- "exact" calculated via `stats::binom.test()` for **categorical** variables
- "wald", "wald.no.correct" calculated via `cardx::proportion_ci_wald(correct = c(TRUE, FALSE))` for **categorical** variables
- "agresti.coull" calculated via `cardx::proportion_ci_agresti_coull()` for **categorical** variables
- "jeffreys" calculated via `cardx::proportion_ci_jeffreys()` for **categorical** variables
- "t.test" calculated via `stats::t.test()` for **continuous** variables
- "wilcox.test" calculated via `stats::wilcox.test()` for **continuous** variables

**Examples**

```
# Example 1 -----
trial |>
  tbl_summary(
    missing = "no",
    statistic = all_continuous() ~ "{mean} ({sd})",
    include = c(marker, response, trt)
  ) |>
  add_ci()

# Example 2 -----
trial |>
  select(response, grade) %>%
  tbl_summary(
    statistic = all_categorical() ~ "{p}%",
    missing = "no",
    include = c(response, grade)
  ) |>
  add_ci(pattern = "{stat} ({ci})") |>
  modify_footnote(everything() ~ NA)
```

---

add\_ci.tbl\_svysummary *Add CI Column*

---

**Description**

Add a new column with the confidence intervals for proportions, means, etc.

**Usage**

```
## S3 method for class 'tbl_svysummary'
add_ci(
  x,
  method = list(all_continuous() ~ "svymean", all_categorical() ~ "svyprop.logit"),
  include = everything(),
  statistic = list(all_continuous() ~ "{conf.low}, {conf.high}", all_categorical() ~
    "{conf.low}%, {conf.high}%"),
  conf.level = 0.95,
  style_fun = list(all_continuous() ~ label_style_sigfig(), all_categorical() ~
    label_style_sigfig(scale = 100)),
  pattern = NULL,
  df = survey::degf(x$inputs$data),
  ...
)
```

**Arguments**

x	(tbl_summary) a summary table of class 'tblsummary'
method	( <a href="#">formula-list-selector</a> ) Confidence interval method. Default is <code>list(all_continuous() ~ "svymean", all_categorical() ~ "svyprop.logit")</code> . See details below.
include	( <a href="#">tidy-select</a> ) Variables to include in the summary table. Default is <code>everything()</code> .
statistic	( <a href="#">formula-list-selector</a> ) Indicates how the confidence interval will be displayed. Default is <code>list(all_continuous() ~ "{conf.low}, {conf.high}", all_categorical() ~ "{conf.low}%, {conf.high}%")</code>
conf.level	(scalar real) Confidence level. Default is 0.95
style_fun	(function) Function to style upper and lower bound of confidence interval. Default is <code>list(all_continuous() ~ label_style_sigfig(), all_categorical() ~ label_style_sigfig(s = 100))</code> .
pattern	(string) Indicates the pattern to use to merge the CI with the statistics cell. The default is NULL, where no columns are merged. The two columns that will be merged are the statistics column, represented by "{stat}" and the CI column represented by "{ci}", e.g. <code>pattern = "{stat} ({ci})"</code> will merge the two columns with the CI in parentheses. Default is NULL, and no merging is performed.
df	(numeric) denominator degrees of freedom, passed to <code>survey::svyciprop(df)</code> or <code>confint(df)</code> . Default is <code>survey::degf(x\$inputs\$data)</code> .
...	These dots are for future extensions and must be empty.

**Value**

gtsummary table

**method argument**

Must be one of

- "svyprop.logit", "svyprop.likelihood", "svyprop.asin", "svyprop.beta", "svyprop.mean", "svyprop.xlogit" calculated via `survey::svyciprop()` for **categorical** variables
- "svymean" calculated via `survey::svymean()` for **continuous** variables
- "svymedian.mean", "svymedian.beta", "svymedian.xlogit", "svymedian.asin", "svymedian.score" calculated via `survey::svyquantile(quantiles = 0.5)` for **continuous** variables

**Examples**

```

data(api, package = "survey")
survey::svydesign(id = ~dnum, weights = ~pw, data = apiclus1, fpc = ~fpc) |>
  tbl_svsummary(
    by = "both",
    include = c(api00, stype),
    statistic = all_continuous() ~ "{mean} ({sd})"
  ) |>
  add_stat_label() |>
  add_ci(pattern = "{stat} (95% CI {ci})") |>
  modify_header(all_stat_cols() ~ "**{level}**") |>
  modify_spanning_header(all_stat_cols() ~ "**Survived**")

```

---

```
add_difference.tbl_summary
```

*Add differences between groups*

---

**Description**

Adds difference to tables created by `tbl_summary()`. The difference between two groups (typically mean or rate difference) is added to the table along with the difference's confidence interval and a p-value (when applicable).

**Usage**

```

## S3 method for class 'tbl_summary'
add_difference(
  x,
  test = NULL,
  group = NULL,
  adj.vars = NULL,
  test.args = NULL,
  conf.level = 0.95,
  include = everything(),
  pvalue_fun = label_style_pvalue(digits = 1),
  estimate_fun = list(c(all_continuous(), all_categorical(FALSE)) ~ label_style_sigfig(),
    all_dichotomous() ~ label_style_sigfig(scale = 100, suffix = "%"), all_tests("smd")
    ~ label_style_sigfig()),
  ...
)

```

**Arguments**

x (tbl\_summary)  
table created with `tbl_summary()`



test	( <a href="#">formula-list-selector</a> ) Specifies the tests/methods to perform for each variable, e.g. <code>list(all_continuous() ~ "t.test", all_dichotomous() ~ "prop.test", all_categorical(FALSE) ~ "smd")</code> . See below for details on default tests and <a href="#">?tests</a> for details on available tests and creating custom tests.
group	( <a href="#">tidy-select</a> ) Variable name of an ID or grouping variable. The column can be used to calculate p-values with correlated data. Default is NULL. See <a href="#">tests</a> for methods that utilize the group argument.
adj.vars	( <a href="#">tidy-select</a> ) Variables to include in adjusted calculations (e.g. in ANCOVA models). Default is NULL.
test.args	( <a href="#">formula-list-selector</a> ) Containing additional arguments to pass to tests that accept arguments. For example, add an argument for all t-tests, use <code>test.args = all_tests("t.test") ~ list(var.equal = TRUE)</code> .
conf.level	(numeric) a scalar in the interval (0, 1) indicating the confidence level. Default is 0.95
include	( <a href="#">tidy-select</a> ) Variables to include in output. Default is <code>everything()</code> .
pvalue_fun	(function) Function to round and format p-values. Default is <code>label_style_pvalue()</code> . The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. <code>pvalue_fun = label_style_pvalue(digits = 2)</code> ).
estimate_fun	( <a href="#">formula-list-selector</a> ) List of formulas specifying the functions to round and format differences and confidence limits.
...	These dots are for future extensions and must be empty.

**Value**

a gtsummary table of class "tbl\_summary"

**Examples**

```
# Example 1 -----
trial |>
  select(trt, age, marker, response, death) %>%
  tbl_summary(
    by = trt,
    statistic =
      list(
        all_continuous() ~ "{mean} ({sd})",
        all_dichotomous() ~ "{p}%"
      ),
```

```

      missing = "no"
    ) |>
    add_n() |>
    add_difference()

# Example 2 -----
# ANCOVA adjusted for grade and stage
trial |>
  select(trt, age, marker, grade, stage) %>%
  tbl_summary(
    by = trt,
    statistic = list(all_continuous() ~ "{mean} ({sd})"),
    missing = "no",
    include = c(age, marker, trt)
  ) |>
  add_n() |>
  add_difference(adj.vars = c(grade, stage))

```

---

```
add_difference.tbl_svsummary
```

*Add differences between groups*

---

## Description

Adds difference to tables created by `tbl_summary()`. The difference between two groups (typically mean or rate difference) is added to the table along with the difference's confidence interval and a p-value (when applicable).

## Usage

```

## S3 method for class 'tbl_svsummary'
add_difference(
  x,
  test = NULL,
  group = NULL,
  adj.vars = NULL,
  test.args = NULL,
  conf.level = 0.95,
  include = everything(),
  pvalue_fun = label_style_pvalue(digits = 1),
  estimate_fun = list(c(all_continuous(), all_categorical(FALSE)) ~ label_style_sigfig(),
    all_dichotomous() ~ label_style_sigfig(scale = 100, suffix = "%"), all_tests("smd")
    ~ label_style_sigfig()),
  ...
)

```

**Arguments**

x	(tbl_summary) table created with tbl_summary()
test	(formula-list-selector) Specifies the tests/methods to perform for each variable, e.g. list(all_continuous() ~ "t.test", all_dichotomous() ~ "prop.test", all_categorical(FALSE) ~ "smd").  See below for details on default tests and <a href="#">?tests</a> for details on available tests and creating custom tests.
group	(tidy-select) Variable name of an ID or grouping variable. The column can be used to calculate p-values with correlated data. Default is NULL. See <a href="#">tests</a> for methods that utilize the group argument.
adj.vars	(tidy-select) Variables to include in adjusted calculations (e.g. in ANCOVA models). Default is NULL.
test.args	(formula-list-selector) Containing additional arguments to pass to tests that accept arguments. For example, add an argument for all t-tests, use test.args = all_tests("t.test") ~ list(var.equal = TRUE).
conf.level	(numeric) a scalar in the interval (0, 1) indicating the confidence level. Default is 0.95
include	(tidy-select) Variables to include in output. Default is everything().
pvalue_fun	(function) Function to round and format p-values. Default is label_style_pvalue(). The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. pvalue_fun = label_style_pvalue(digits = 2)).
estimate_fun	(formula-list-selector) List of formulas specifying the functions to round and format differences and confidence limits. Default is list(c(all_continuous(), all_categorical(FALSE)) ~ label_style
...	These dots are for future extensions and must be empty.

**Value**

a gtsummary table of class "tbl\_summary"

**Examples**

---

 add\_glance

*Add model statistics*


---

## Description

Add model statistics returned from `broom::glance()`. Statistics can either be appended to the table (`add_glance_table()`), or added as a table source note (`add_glance_source_note()`).

## Usage

```
add_glance_table(
  x,
  include = everything(),
  label = NULL,
  fmt_fun = list(everything() ~ label_style_sigfig(digits = 3), any_of("p.value") ~
    label_style_pvalue(digits = 1), c(where(is.integer), starts_with("df")) ~
    label_style_number()),
  glance_fun = glance_fun_s3(x$inputs$x)
)

add_glance_source_note(
  x,
  include = everything(),
  label = NULL,
  fmt_fun = list(everything() ~ label_style_sigfig(digits = 3), any_of("p.value") ~
    label_style_pvalue(digits = 1), c(where(is.integer), starts_with("df")) ~
    label_style_number()),
  glance_fun = glance_fun_s3(x$inputs$x),
  text_interpret = c("md", "html"),
  sep1 = " = ",
  sep2 = "; "
)
```

## Arguments

<code>x</code>	( <code>tbl_regression</code> ) a 'tbl_regression' object
<code>include</code>	( <code>tidy-select</code> ) names of statistics to include in output. Must be column names of the tibble returned by <code>broom::glance()</code> or from the <code>glance_fun</code> argument. The <code>include</code> argument can also be used to specify the order the statistics appear in the table.
<code>label</code>	( <code>formula-list-selector</code> ) specifies statistic labels, e.g. <code>list(r.squared = "R2", p.value = "P")</code>
<code>fmt_fun</code>	( <code>formula-list-selector</code> ) Specifies the the functions used to format/round the glance statistics. The default is to round the number of observations and degrees of freedom to the nearest

	integer, p-values are styled with <code>style_pvalue()</code> and the remaining statistics are styled with <code>style_sigfig(x, digits = 3)</code>
<code>glance_fun</code>	(function) function that returns model statistics. Default is <code>glance_fun()</code> (which is <code>broom::glance()</code> for most model objects). Custom functions must return a single row tibble.
<code>text_interpret</code>	(string) String indicates whether source note text will be interpreted with <code>gt::md()</code> or <code>gt::html()</code> . Must be "md" (default) or "html".
<code>sep1</code>	(string) Separator between statistic name and statistic. Default is " = ", e.g. "R2 = 0.456"
<code>sep2</code>	(string) Separator between statistics. Default is "; "

**Value**

gtsummary table

**Tips**

When combining `add_glance_table()` with `tbl_merge()`, the ordering of the model terms and the glance statistics may become jumbled. To re-order the rows with glance statistics on bottom, use the script below:

```
tbl_merge(list(tbl1, tbl2)) %>%
  modify_table_body(~.x %>% arrange(row_type == "glance_statistic"))
```

**Examples**

```
mod <- lm(age ~ marker + grade, trial) |> tbl_regression()

# Example 1 -----
mod |>
  add_glance_table(
    label = list(sigma = "\U03C3"),
    include = c(r.squared, AIC, sigma)
  )

# Example 2 -----
mod |>
  add_glance_source_note(
    label = list(sigma = "\U03C3"),
    include = c(r.squared, AIC, sigma)
  )
```

---

add_global_p	<i>Add the global p-values</i>
--------------	--------------------------------

---

**Description**

This function uses `car::Anova()` (by default) to calculate global p-values for model covariates. Output from `tbl_regression` and `tbl_uvregression` objects supported.

**Usage**

```
add_global_p(x, ...)

## S3 method for class 'tbl_regression'
add_global_p(
  x,
  include = everything(),
  keep = FALSE,
  anova_fun = global_pvalue_fun,
  type = "III",
  quiet,
  ...
)

## S3 method for class 'tbl_uvregression'
add_global_p(
  x,
  include = everything(),
  keep = FALSE,
  anova_fun = global_pvalue_fun,
  type = "III",
  quiet,
  ...
)
```

**Arguments**

x	(tbl_regression, tbl_uvregression) Object with class 'tbl_regression' or 'tbl_uvregression'
...	Additional arguments to be passed to <code>car::Anova</code> , <code>aod::wald.test()</code> or <code>anova_fun</code> (if specified)
include	( <a href="#">tidy-select</a> ) Variables to calculate global p-value for. Default is <code>everything()</code>
keep	(scalar logical) Logical argument indicating whether to also retain the individual p-values in the table output for each level of the categorical variable. Default is <code>FALSE</code> .

anova_fun	(function) Function used to calculate global p-values. Default is generic <code>global_pvalue_fun()</code> , which wraps <code>car::Anova()</code> for most models. The <code>type</code> argument is passed to this function. See help file for details.  To pass a custom function, it must accept as its first argument is a model. Note that anything passed in <code>...</code> will be passed to this function. The function must return an object of class 'cards' (see <code>cardx::ard_car_anova()</code> as an example), or a tibble with columns 'term' and 'p.value' (e.g. <code>\(x, type, ...)</code> <code>car::Anova(x, type, ...)</code>  >
type	Type argument passed to <code>anova_fun</code> . Default is "III"
quiet	<b>[Deprecated]</b>

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
lm(marker ~ age + grade, trial) |>
tbl_regression() |>
add_global_p()

# Example 2 -----
trial[c("response", "age", "trt", "grade")] |>
tbl_uvregression(
  method = glm,
  y = response,
  method.args = list(family = binomial),
  exponentiate = TRUE
) |>
add_global_p()
```

---

add\_n.tbl\_survfit      *Add N*

---

**Description**

For each `survfit()` object summarized with `tbl_survfit()` this function will add the total number of observations in a new column.

**Usage**

```
## S3 method for class 'tbl_survfit'
add_n(x, ...)
```

**Arguments**

x	object of class "tbl_survfit"
...	Not used

**Examples**

```
library(survival)
fit1 <- survfit(Surv(ttdeath, death) ~ 1, trial)
fit2 <- survfit(Surv(ttdeath, death) ~ trt, trial)

# Example 1 -----
list(fit1, fit2) |>
  tbl_survfit(times = c(12, 24)) |>
  add_n()
```

---

add\_nevent.tbl\_survfit

*Add event N*

---

**Description**

For each `survfit()` object summarized with `tbl_survfit()` this function will add the total number of events observed in a new column.

**Usage**

```
## S3 method for class 'tbl_survfit'
add_nevent(x, ...)
```

**Arguments**

x	object of class 'tbl_survfit'
...	Not used

**See Also**

Other `tbl_survfit` tools: [add\\_p.tbl\\_survfit\(\)](#)

**Examples**

```
library(survival)
fit1 <- survfit(Surv(ttdeath, death) ~ 1, trial)
fit2 <- survfit(Surv(ttdeath, death) ~ trt, trial)

# Example 1 -----
list(fit1, fit2) |>
  tbl_survfit(times = c(12, 24)) |>
```



```
add_n() |>
add_nevent()
```

---

```
add_nevent_regression Add event N
```

---

## Description

Add event N

## Usage

```
add_nevent(x, ...)

## S3 method for class 'tbl_regression'
add_nevent(x, location = "label", ...)

## S3 method for class 'tbl_uvregression'
add_nevent(x, location = "label", ...)
```

## Arguments

x	(tbl_regression, tbl_uvregression) a tbl_regression or tbl_uvregression table
...	These dots are for future extensions and must be empty.
location	(character) location to place Ns. Select one or more of c('label', 'level'). Default is 'label'. When "label" total Ns are placed on each variable's label row. When "level" level counts are placed on the variable level for categorical variables, and total N on the variable's label row for continuous.

## Examples

```
# Example 1 -----
trial |>
  select(response, trt, grade) |>
  tbl_uvregression(
    y = response,
    exponentiate = TRUE,
    method = glm,
    method.args = list(family = binomial),
  ) |>
  add_nevent()

# Example 2 -----
glm(response ~ age + grade, trial, family = binomial) |>
```

```
tbl_regression(exponentiate = TRUE) |>
add_nevent(location = "level")
```

---

```
add_n_regression      Add N to regression table
```

---

## Description

Add N to regression table

## Usage

```
## S3 method for class 'tbl_regression'
add_n(x, location = "label", ...)

## S3 method for class 'tbl_uvregression'
add_n(x, location = "label", ...)
```

## Arguments

`x` (tbl\_regression, tbl\_uvregression)  
a tbl\_regression or tbl\_uvregression table

`location` (character)  
location to place Ns. Select one or more of c('label', 'level'). Default is 'label'.  
When "label" total Ns are placed on each variable's label row. When "level" level counts are placed on the variable level for categorical variables, and total N on the variable's label row for continuous.

... These dots are for future extensions and must be empty.

## Examples

```
# Example 1 -----
trial |>
  select(response, age, grade) |>
  tbl_uvregression(
    y = response,
    exponentiate = TRUE,
    method = glm,
    method.args = list(family = binomial),
    hide_n = TRUE
  ) |>
  add_n(location = "label")

# Example 2 -----
glm(response ~ age + grade, trial, family = binomial) |>
tbl_regression(exponentiate = TRUE) |>
add_n(location = "level")
```

---

add_n_summary	<i>Add column with N</i>
---------------	--------------------------

---

## Description

For each variable in a `tbl_summary` table, the `add_n` function adds a column with the total number of non-missing (or missing) observations

## Usage

```
## S3 method for class 'tbl_summary'
add_n(
  x,
  statistic = "{N_nonmiss}",
  col_label = "**N**",
  footnote = FALSE,
  last = FALSE,
  ...
)

## S3 method for class 'tbl_svsummary'
add_n(
  x,
  statistic = "{N_nonmiss}",
  col_label = "**N**",
  footnote = FALSE,
  last = FALSE,
  ...
)

## S3 method for class 'tbl_likert'
add_n(
  x,
  statistic = "{N_nonmiss}",
  col_label = "**N**",
  footnote = FALSE,
  last = FALSE,
  ...
)
```

## Arguments

<code>x</code>	( <code>tbl_summary</code> ) Object with class 'tbl_summary' created with <code>tbl_summary()</code> function.
<code>statistic</code>	(string) String indicating the statistic to report. Default is the number of non-missing ob-

ervation for each variable, `statistic = "{N_nonmiss}"`. All statistics available to report include:

- `"{N_obs}"` total number of observations,
- `"{N_nonmiss}"` number of non-missing observations,
- `"{N_miss}"` number of missing observations,
- `"{p_nonmiss}"` percent non-missing data,
- `"{p_miss}"` percent missing data

The argument uses `glue::glue()` syntax and multiple statistics may be reported, e.g. `statistic = "{N_nonmiss} / {N_obs} ({p_nonmiss}%)"`

<code>col_label</code>	(string) String indicating the column label. Default is <code>"**N**"</code>
<code>footnote</code>	(scalar logical) Logical argument indicating whether to print a footnote clarifying the statistics presented. Default is <code>FALSE</code>
<code>last</code>	(scalar logical) Logical indicator to include N column last in table. Default is <code>FALSE</code> , which will display N column first.
<code>...</code>	These dots are for future extensions and must be empty.

### Value

A table of class `c('tbl_summary', 'gtsummary')`

### Author(s)

Daniel D. Sjoberg

### Examples

```
# Example 1 -----
trial |>
  tbl_summary(by = trt, include = c(trt, age, grade, response)) |>
  add_n()

# Example 2 -----
survey::svydesign(~1, data = as.data.frame(Titanic), weights = ~Freq) |>
  tbl_svysummary(by = Survived, percent = "row", include = c(Class, Age)) |>
  add_n()
```

---

add_overall	<i>Add overall column</i>
-------------	---------------------------

---

### Description

Adds a column with overall summary statistics to tables created by `tbl_summary()`, `tbl_svsummary()`, `tbl_continuous()` or `tbl_custom_summary()`.

### Usage

```
add_overall(x, ...)
```

```
## S3 method for class 'tbl_summary'
add_overall(
  x,
  last = FALSE,
  col_label = "**Overall** \nN = {style_number(N)}",
  statistic = NULL,
  digits = NULL,
  ...
)
```

```
## S3 method for class 'tbl_continuous'
add_overall(
  x,
  last = FALSE,
  col_label = "**Overall** \nN = {style_number(N)}",
  statistic = NULL,
  digits = NULL,
  ...
)
```

```
## S3 method for class 'tbl_svsummary'
add_overall(
  x,
  last = FALSE,
  col_label = "**Overall** \nN = {style_number(N)}",
  statistic = NULL,
  digits = NULL,
  ...
)
```

```
## S3 method for class 'tbl_custom_summary'
add_overall(
  x,
  last = FALSE,
  col_label = "**Overall** \nN = {style_number(N)}",
```

```

    statistic = NULL,
    digits = NULL,
    ...
  )

```

### Arguments

x	(tbl_summary, tbl_svsummary, tbl_continuous, tbl_custom_summary) A stratified 'gtsummary' table
...	These dots are for future extensions and must be empty.
last	(scalar logical) Logical indicator to display overall column last in table. Default is FALSE, which will display overall column first.
col_label	(string) String indicating the column label. Default is "**Overall** \nN = {style_number(N)}"
statistic	( <a href="#">formula-list-selector</a> ) Override the statistic argument in initial tbl_* function call. Default is NULL.
digits	( <a href="#">formula-list-selector</a> ) Override the digits argument in initial tbl_* function call. Default is NULL.

### Value

A gtsummary of same class as x

### Author(s)

Daniel D. Sjoberg

### Examples

```

# Example 1 -----
trial |>
  tbl_summary(include = c(age, grade), by = trt) |>
  add_overall()

# Example 2 -----
trial |>
  tbl_summary(
    include = grade,
    by = trt,
    percent = "row",
    statistic = ~"{p}%",
    digits = ~1
  ) |>
  add_overall(
    last = TRUE,
    statistic = ~"{p}% (n={n})",
    digits = ~ c(1, 0)
  )

```

```
# Example 3 -----
trial |>
  tbl_continuous(
    variable = age,
    by = trt,
    include = grade
  ) |>
  add_overall(last = TRUE)
```

---

add_overall_ard	<i>ARD add overall column</i>
-----------------	-------------------------------

---

### Description

Adds a column with overall summary statistics to tables created by `tbl_ard_summary()`.

### Usage

```
## S3 method for class 'tbl_ard_summary'
add_overall(
  x,
  cards,
  last = FALSE,
  col_label = "**Overall**",
  statistic = NULL,
  ...
)
```

### Arguments

<code>x</code>	(tbl_ard_summary) A stratified 'gtsummary' table
<code>cards</code>	(card) An ARD object of class "card" typically created with <code>cards::ard_*</code> () functions.
<code>last</code>	(scalar logical) Logical indicator to display overall column last in table. Default is FALSE, which will display overall column first.
<code>col_label</code>	(string) String indicating the column label. Default is " <b>Overall</b> "
<code>statistic</code>	( <a href="#">formula-list-selector</a> ) Override the statistic argument in initial <code>tbl_*</code> function call. Default is NULL.
<code>...</code>	These dots are for future extensions and must be empty.

### Value

A `gtsummary` of same class as `x`

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
# build primary table
tbl <-
  cards::ard_stack(
    trial,
    .by = trt,
    cards::ard_continuous(variables = age),
    cards::ard_categorical(variables = grade),
    .missing = TRUE,
    .attributes = TRUE,
    .total_n = TRUE
  ) |>
  tbl_ard_summary(by = trt)

# create ARD with overall results
ard_overall <-
  cards::ard_stack(
    trial,
    cards::ard_continuous(variables = age),
    cards::ard_categorical(variables = grade),
    .missing = TRUE,
    .attributes = TRUE,
    .total_n = TRUE
  )

# add an overall column
tbl |>
  add_overall(cards = ard_overall)
```

---

add\_p.tbl\_continuous *Add p-values*

---

**Description**

Add p-values

**Usage**

```
## S3 method for class 'tbl_continuous'
add_p(
  x,
  test = NULL,
  pvalue_fun = label_style_pvalue(digits = 1),
  include = everything(),
```



```

  test.args = NULL,
  group = NULL,
  ...
)

```

## Arguments

x	(tbl_continuous) table created with tbl_continuous()
test	List of formulas specifying statistical tests to perform for each variable. Default is two-way ANOVA when by= is not NULL, and has the same defaults as add_p.tbl_continuous() when by = NULL. See <a href="#">tests</a> for details, more tests, and instruction for implementing a custom test.
pvalue_fun	(function) Function to round and format p-values. Default is label_style_pvalue(). The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. pvalue_fun = label_style_pvalue(digits = 2)).
include	(tidy-select) Variables to include in output. Default is everything().
test.args	(formula-list-selector) Containing additional arguments to pass to tests that accept arguments. For example, add an argument for all t-tests, use test.args = all_tests("t.test") ~ list(var.equal = TRUE).
group	(tidy-select) Variable name of an ID or grouping variable. The column can be used to calculate p-values with correlated data. Default is NULL. See <a href="#">tests</a> for methods that utilize the group argument.
...	These dots are for future extensions and must be empty.

## Value

'tbl\_continuous' object

## Examples

```

trial |>
  tbl_continuous(variable = age, by = trt, include = grade) |>
  add_p()

```

---

add_p.tbl_cross	<i>Add p-value</i>
-----------------	--------------------

---

### Description

Calculate and add a p-value comparing the two variables in the cross table. If missing levels are included in the tables, they are also included in p-value calculation.

### Usage

```
## S3 method for class 'tbl_cross'
add_p(
  x,
  test = NULL,
  pvalue_fun = ifelse(source_note, label_style_pvalue(digits = 1, prepend_p = TRUE),
    label_style_pvalue(digits = 1)),
  source_note = FALSE,
  test.args = NULL,
  ...
)
```

### Arguments

x	(tbl_cross) Object with class <code>tbl_cross</code> created with the <code>tbl_cross()</code> function
test	(string) A string specifying statistical test to perform. Default is "chisq.test" when expected cell counts $\geq 5$ and "fisher.test" when expected cell counts $< 5$ .
pvalue_fun	(function) Function to round and format p-value. Default is <code>label_style_pvalue(digits = 1)</code> , except when <code>source_note = TRUE</code> when the default is <code>label_style_pvalue(digits = 1, prepend_p = TRUE)</code>
source_note	(scalar logical) Logical value indicating whether to show p-value in the {gt} table source notes rather than a column.
test.args	(named list) Named list containing additional arguments to pass to the test (if it accepts additional arguments). For example, add an argument for a chi-squared test with <code>test.args = list(correct = TRUE)</code>
...	These dots are for future extensions and must be empty.

### Author(s)

Karissa Whiting, Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
trial |>
  tbl_cross(row = stage, col = trt) |>
  add_p()

# Example 2 -----
trial |>
  tbl_cross(row = stage, col = trt) |>
  add_p(source_note = TRUE)
```

---

add\_p.tbl\_summary      *Add p-values*

---

**Description**

Adds p-values to tables created by `tbl_summary()` by comparing values across groups.

**Usage**

```
## S3 method for class 'tbl_summary'
add_p(
  x,
  test = NULL,
  pvalue_fun = label_style_pvalue(digits = 1),
  group = NULL,
  include = everything(),
  test.args = NULL,
  adj.vars = NULL,
  ...
)
```

**Arguments**

<code>x</code>	(tbl_summary) table created with <code>tbl_summary()</code>
<code>test</code>	( <a href="#">formula-list-selector</a> ) Specifies the statistical tests to perform for each variable, e.g. <code>list(all_continuous() ~ "t.test", all_categorical() ~ "fisher.test")</code> . See below for details on default tests and <a href="#">?tests</a> for details on available tests and creating custom tests.
<code>pvalue_fun</code>	(function) Function to round and format p-values. Default is <code>label_style_pvalue()</code> . The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. <code>pvalue_fun = label_style_pvalue(digits = 2)</code> ).

group	( <a href="#">tidy-select</a> ) Variable name of an ID or grouping variable. The column can be used to calculate p-values with correlated data. Default is NULL. See <a href="#">tests</a> for methods that utilize the group argument.
include	( <a href="#">tidy-select</a> ) Variables to include in output. Default is everything().
test.args	( <a href="#">formula-list-selector</a> ) Containing additional arguments to pass to tests that accept arguments. For example, add an argument for all t-tests, use <code>test.args = all_tests("t.test") ~ list(var.equal = TRUE)</code> .
adj.vars	( <a href="#">tidy-select</a> ) Variables to include in adjusted calculations (e.g. in ANCOVA models). Default is NULL.
...	These dots are for future extensions and must be empty.

**Value**

a gtsummary table of class "tbl\_summary"

**test argument**

See the [?tests](#) help file for details on available tests and creating custom tests. The [?tests](#) help file also includes pseudo-code for each test to be clear precisely how the calculation is performed.

The default test used in `add_p()` primarily depends on these factors:

- whether the variable is categorical/dichotomous vs continuous
- number of levels in the `tbl_summary(by)` variable
- whether the `add_p(group)` argument is specified
- whether the `add_p(adj.vars)` argument is specified

**Specified neither `add_p(group)` nor `add_p(adj.vars)`:**

- `"wilcox.test"` when by variable has two levels and variable is continuous.
- `"kruskal.test"` when by variable has more than two levels and variable is continuous.
- `"chisq.test.no.correct"` for categorical variables with all expected cell counts  $\geq 5$ , and `"fisher.test"` for categorical variables with any expected cell count  $< 5$ .

**Specified `add_p(group)` and not `add_p(adj.vars)`:**

- `"lme4"` when by variable has two levels for all summary types.

*There is no default for grouped data when by variable has more than two levels. Users must create custom tests for this scenario.*

**Specified `add_p(adj.vars)` and not `add_p(group)`:**

- `"ancova"` when variable is continuous and by variable has two levels.

**Examples**

```
# Example 1 -----
trial |>
  tbl_summary(by = trt, include = c(age, grade)) |>
  add_p()

# Example 2 -----
trial |>
  select(trt, age, marker) |>
  tbl_summary(by = trt, missing = "no") |>
  add_p(
    # perform t-test for all variables
    test = everything() ~ "t.test",
    # assume equal variance in the t-test
    test.args = all_tests("t.test") ~ list(var.equal = TRUE)
  )
```

---

add_p.tbl_survfit	<i>Add p-value</i>
-------------------	--------------------

---

**Description**

Calculate and add a p-value to stratified `tbl_survfit()` tables.

**Usage**

```
## S3 method for class 'tbl_survfit'
add_p(
  x,
  test = "logrank",
  test.args = NULL,
  pvalue_fun = label_style_pvalue(digits = 1),
  include = everything(),
  quiet,
  ...
)
```

**Arguments**

x	(tbl_survfit) Object of class "tbl_survfit"
test	(string) string indicating test to use. Must be one of "logrank", "tarone", "survdiff", "petopeto_gehanwilcoxon", "coxph_lrt", "coxph_wald", "coxph_score". See details below

test.args	(named list) named list of arguments that will be passed to the method specified in the test argument. Default is NULL.
pvalue_fun	(function) Function to round and format p-values. Default is label_style_pvalue(). The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. pvalue_fun = label_style_pvalue(digits = 2)).
include	(tidy-select) Variables to include in output. Default is everything().
quiet	<b>[Deprecated]</b>
...	These dots are for future extensions and must be empty.

### test argument

The most common way to specify `test=` is by using a single string indicating the test name. However, if you need to specify different tests within the same table, the input is flexible using the list notation common throughout the `gtsummary` package. For example, the following code would call the log-rank test, and a second test of the *G-rho* family.

```
... |>
  add_p(test = list(trt ~ "logrank", grade ~ "survdiff"),
        test.args = grade ~ list(rho = 0.5))
```

### Note

To calculate the p-values, the formula is re-constructed from the the call in the original `survfit()` object. When the `survfit()` object is created a for loop, `lapply()`, `purrr::map()` setting the call *may not* reflect the true formula which may result in an error or an incorrect calculation.

To ensure correct results, the call formula in `survfit()` must represent the formula that will be used in `survival::survdiff()`. If you utilize the `tbl_survfit.data.frame()` S3 method, this is handled for you.

### See Also

Other `tbl_survfit` tools: [add\\_nevent.tbl\\_survfit\(\)](#)

### Examples

```
library(survival)

gts_survfit <-
  list(
    survfit(Surv(ttdeath, death) ~ grade, trial),
    survfit(Surv(ttdeath, death) ~ trt, trial)
  ) |>
  tbl_survfit(times = c(12, 24))
```

```

# Example 1 -----
gts_survfit |>
  add_p()

# Example 2 -----
# Pass `rho=` argument to `survdiff()`
gts_survfit |>
  add_p(test = "survdiff", test.args = list(rho = 0.5))

```

---

add\_p.tbl\_svysummary *Add p-values*

---

## Description

Adds p-values to tables created by `tbl_svysummary()` by comparing values across groups.

## Usage

```

## S3 method for class 'tbl_svysummary'
add_p(
  x,
  test = list(all_continuous() ~ "svy.wilcox.test", all_categorical() ~ "svy.chisq.test"),
  pvalue_fun = label_style_pvalue(digits = 1),
  include = everything(),
  test.args = NULL,
  ...
)

```

## Arguments

<code>x</code>	(tbl_svysummary) table created with <code>tbl_svysummary()</code>
<code>test</code>	( <a href="#">formula-list-selector</a> ) List of formulas specifying statistical tests to perform. Default is <code>list(all_continuous() ~ "svy.wilcox.test", all_categorical() ~ "svy.chisq.test")</code> . See below for details on default tests and <a href="#">?tests</a> for details on available tests and creating custom tests.
<code>pvalue_fun</code>	(function) Function to round and format p-values. Default is <code>label_style_pvalue()</code> . The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. <code>pvalue_fun = label_style_pvalue(digits = 2)</code> ).
<code>include</code>	( <a href="#">tidy-select</a> ) Variables to include in output. Default is <code>everything()</code> .

```
test.args      (formula-list-selector)
                Containing additional arguments to pass to tests that accept arguments. For ex-
                ample, add an argument for all t-tests, use test.args = all_tests("t.test")
                ~ list(var.equal = TRUE).
...           These dots are for future extensions and must be empty.
```

## Value

a gsummary table of class "tbl\_svysummary"

## Examples

```
# Example 1 -----
# A simple weighted dataset
survey::svydesign(~1, data = as.data.frame(Titanic), weights = ~Freq) |>
  tbl_svysummary(by = Survived, include = c(Sex, Age)) |>
  add_p()

# A dataset with a complex design
data(api, package = "survey")
d_clust <- survey::svydesign(id = ~dnum, weights = ~pw, data = apiclus1, fpc = ~fpc)

# Example 2 -----
tbl_svysummary(d_clust, by = both, include = c(api00, api99)) |>
  add_p()

# Example 3 -----
# change tests to svy t-test and Wald test
tbl_svysummary(d_clust, by = both, include = c(api00, api99, stype)) |>
  add_p(
    test = list(
      all_continuous() ~ "svy.t.test",
      all_categorical() ~ "svy.wald.test"
    )
  )
```

---

add\_q

*Add multiple comparison adjustment*

---

## Description

Adjustments to p-values are performed with `stats::p.adjust()`.

## Usage

```
add_q(x, method = "fdr", pvalue_fun = NULL, quiet = NULL)
```



**Arguments**

x	(gtsummary) a gtsummary object with a column named "p.value"
method	(string) String indicating method to be used for p-value adjustment. Methods from <code>stats::p.adjust()</code> are accepted. Default is <code>method='fdr'</code> . Must be one of <code>'holm'</code> , <code>'hochberg'</code> , <code>'hommel'</code> , <code>'bonferroni'</code> , <code>'BH'</code> , <code>'BY'</code> , <code>'fdr'</code> , <code>'none'</code>
pvalue_fun	(function) Function to round and format q-values. Default is the function specified to round the existing 'p.value' column.
quiet	<b>[Deprecated]</b>

**Author(s)**

Daniel D. Sjoberg, Esther Drill

**Examples**

```
# Example 1 -----
add_q_ex1 <-
  trial |>
  tbl_summary(by = trt, include = c(trt, age, grade, response)) |>
  add_p() |>
  add_q()

# Example 2 -----
trial |>
  tbl_uvregression(
    y = response,
    include = c("trt", "age", "grade"),
    method = glm,
    method.args = list(family = binomial),
    exponentiate = TRUE
  ) |>
  add_global_p() |>
  add_q()
```

---

add\_significance\_stars

*Add significance stars*

---

**Description**

Add significance stars to estimates with small p-values

**Usage**

```
add_significance_stars(
  x,
  pattern = ifelse(inherits(x, c("tbl_regression", "tbl_uvregression")),
    "{estimate}{stars}", "{p.value}{stars}"),
  thresholds = c(0.001, 0.01, 0.05),
  hide_ci = TRUE,
  hide_p = inherits(x, c("tbl_regression", "tbl_uvregression")),
  hide_se = FALSE
)
```

**Arguments**

x	(gtsummary) A 'gtsummary' object with a 'p.value' column
pattern	(string) glue-syntax string indicating what to display in formatted column. Default is "{estimate}{stars}" for regression summaries and "{p.value}{stars}" otherwise. A footnote is placed on the first column listed in the pattern. Other common patterns are "{estimate}{stars} ({conf.low}, {conf.high})" and "{estimate} ({conf.low} to {conf.high}){stars}"
thresholds	(numeric) Thresholds for significance stars. Default is c(0.001, 0.01, 0.05)
hide_ci	(scalar logical) logical whether to hide confidence interval. Default is TRUE
hide_p	(scalar logical) logical whether to hide p-value. Default is TRUE for regression summaries, and FALSE otherwise.
hide_se	(scalar logical) logical whether to hide standard error. Default is FALSE

**Value**

a 'gtsummary' table

**Examples**

```
tbl <-
  lm(time ~ ph.ecog + sex, survival::lung) |>
  tbl_regression(label = list(ph.ecog = "ECOG Score", sex = "Sex"))

# Example 1 -----
tbl |>
  add_significance_stars(hide_ci = FALSE, hide_p = FALSE)

# Example 2 -----
tbl |>
  add_significance_stars(
```

```

    pattern = "{estimate} ({conf.low}, {conf.high}){stars}",
    hide_ci = TRUE, hide_se = TRUE
  ) |>
  modify_header(estimate = "***Beta (95% CI)**") |>
  modify_footnote(estimate = "CI = Confidence Interval", abbreviation = TRUE)

# Example 3 -----
# Use ' \n' to put a line break between beta and SE
tbl |>
  add_significance_stars(
    hide_se = TRUE,
    pattern = "{estimate}{stars} \n({std.error})"
  ) |>
  modify_header(estimate = "***Beta \n(SE)**") |>
  modify_footnote(estimate = "SE = Standard Error", abbreviation = TRUE) |>
  as_gt() |>
  gt::fmt_markdown(columns = everything()) |>
  gt::tab_style(
    style = "vertical-align:top",
    locations = gt::cells_body(columns = label)
  )

# Example 4 -----
lm(marker ~ stage + grade, data = trial) |>
  tbl_regression() |>
  add_global_p() |>
  add_significance_stars(
    hide_p = FALSE,
    pattern = "{p.value}{stars}"
  )

```

---

 add\_stat

*Add a custom statistic*


---

## Description

The function allows a user to add a new column (or columns) of statistics to an existing `tbl_summary`, `tbl_svsummary`, or `tbl_continuous` object.

## Usage

```
add_stat(x, fns, location = everything() ~ "label")
```

## Arguments

`x` (tbl\_summary/tbl\_svsummary/tbl\_continuous)  
 A gtsummary table of class 'tbl\_summary', 'tbl\_svsummary', or 'tbl\_continuous'.

`fns` ([formula-list-selector](#))  
 Indicates the functions that create the statistic. See details below.

location (formula-list-selector)  
 Indicates the location the new statistics are placed. The values must be one of `c("label", "level", "missing")`. When "label", a single statistic is placed on the variable label row. When "level" the statistics are placed on the variable level rows. The length of the vector of statistics returned from the `fns` function must match the dimension of levels. Default is to place the new statistics on the label row.

### Value

A 'gtsummary' of the same class as the input

### Details

The returns from custom functions passed in `fns=` are required to follow a specified format. Each of these function will execute on a single variable.

1. Each function must return a tibble or a vector. If a vector is returned, it will be converted to a tibble with one column and number of rows equal to the length of the vector.
2. When `location='label'`, the returned statistic from the custom function must be a tibble with one row. When `location='level'` the tibble must have the same number of rows as there are levels in the variable (excluding the row for unknown values).
3. Each function may take the following arguments: `foo(data, variable, by, tbl, ...)`
  - `data=` is the input data frame passed to `tbl_summary()`
  - `variable=` is a string indicating the variable to perform the calculation on. This is the variable in the label column of the table.
  - `by=` is a string indicating the by variable from `tbl_summary=`, if present
  - `tbl=` the original `tbl_summary()/tbl_svysummary()` object is also available to utilize

The user-defined function does not need to utilize each of these inputs. It's encouraged the user-defined function accept `...` as each of the arguments *will* be passed to the function, even if not all inputs are utilized by the user's function, e.g. `foo(data, variable, by, ...)`

- Use `modify_header()` to update the column headers
- Use `modify_fmt_fun()` to update the functions that format the statistics
- Use `modify_footnote()` to add a explanatory footnote

If you return a tibble with column names `p.value` or `q.value`, default p-value formatting will be applied, and you may take advantage of subsequent p-value formatting functions, such as `bold_p()` or `add_q()`.

### Examples

```
# Example 1 -----
# fn returns t-test pvalue
my_ttest <- function(data, variable, by, ...) {
  t.test(data[[variable]] ~ as.factor(data[[by]]))$p.value
}
```

```

trial |>
  tbl_summary(
    by = trt,
    include = c(trt, age, marker),
    missing = "no"
  ) |>
  add_stat(fns = everything() ~ my_ttest) |>
  modify_header(add_stat_1 = "**p-value**", all_stat_cols() ~ "**{level}**")

# Example 2 -----
# fn returns t-test test statistic and pvalue
my_ttest2 <- function(data, variable, by, ...) {
  t.test(data[[variable]] ~ as.factor(data[[by]])) |>
    broom::tidy() %>%
    dplyr::mutate(
      stat = glue::glue("t={style_sigfig(statistic)}, {style_pvalue(p.value, prepend_p = TRUE)}")
    ) %>%
    dplyr::pull(stat)
}

trial |>
  tbl_summary(
    by = trt,
    include = c(trt, age, marker),
    missing = "no"
  ) |>
  add_stat(fns = everything() ~ my_ttest2) |>
  modify_header(add_stat_1 = "**Treatment Comparison**")

# Example 3 -----
# return test statistic and p-value is separate columns
my_ttest3 <- function(data, variable, by, ...) {
  t.test(data[[variable]] ~ as.factor(data[[by]])) %>%
    broom::tidy() %>%
    select(statistic, p.value)
}

trial |>
  tbl_summary(
    by = trt,
    include = c(trt, age, marker),
    missing = "no"
  ) |>
  add_stat(fns = everything() ~ my_ttest3) |>
  modify_header(statistic = "**t-statistic**", p.value = "**p-value**") |>
  modify_fmt_fun(statistic = label_style_sigfig(), p.value = label_style_pvalue(digits = 2))

```

**Description****[Questioning]**

Adds or modifies labels describing the summary statistics presented for each variable in a `tbl_summary()` table.

**Usage**

```
add_stat_label(x, ...)

## S3 method for class 'tbl_summary'
add_stat_label(x, location = c("row", "column"), label = NULL, ...)

## S3 method for class 'tbl_svysummary'
add_stat_label(x, location = c("row", "column"), label = NULL, ...)

## S3 method for class 'tbl_ard_summary'
add_stat_label(x, location = c("row", "column"), label = NULL, ...)
```

**Arguments**

<code>x</code>	(tbl_summary) Object with class 'tbl_summary' or with class 'tbl_svysummary'
<code>...</code>	These dots are for future extensions and must be empty.
<code>location</code>	(string) Location where statistic label will be included. "row" (the default) to add the statistic label to the variable label row, and "column" adds a column with the statistic label.
<code>label</code>	(formula-list-selector) indicates the updates to the statistic label, e.g. <code>label = all_categorical() ~ "No. (%)"</code> . When not specified, the default statistic labels are used.

**Value**

A `tbl_summary` or `tbl_svysummary` object

**Tips**

When using `add_stat_label(location='row')` with subsequent `tbl_merge()`, it's important to have somewhat of an understanding of the underlying structure of the `gtsummary` table. `add_stat_label(location='row')` works by adding a new column called "stat\_label" to `x$table_body`. The "label" and "stat\_label" columns are merged when the `gtsummary` table is printed. The `tbl_merge()` function merges on the "label" column (among others), which is typically the first column you see in a `gtsummary` table. Therefore, when you want to merge a table that has run `add_stat_label(location='row')` you need to match the "label" column values before the "stat\_column" is merged with it.

For example, the following two tables merge properly

```
tbl1 <- trial %>% select(age, grade) |> tbl_summary() |> add_stat_label()
tbl2 <- lm(marker ~ age + grade, trial) |> tbl_regression()
```

```
tbl_merge(list(tbl1, tbl2))
```

The addition of the new "stat\_label" column requires a default labels for categorical variables, which is "No. (%)". This can be changed to either desired text or left blank using NA\_character\_. The blank option is useful in the location="row" case to keep the output for categorical variables identical what was produced without a "add\_stat\_label()" function call.

### Author(s)

Daniel D. Sjoberg

### Examples

```
tbl <- trial |>
  dplyr::select(trt, age, grade, response) |>
  tbl_summary(by = trt)

# Example 1 -----
# Add statistic presented to the variable label row
tbl |>
  add_stat_label(
    # update default statistic label for continuous variables
    label = all_continuous() ~ "med. (iqr)"
  )

# Example 2 -----
tbl |>
  add_stat_label(
    # add a new column with statistic labels
    location = "column"
  )

# Example 3 -----
trial |>
  select(age, grade, trt) |>
  tbl_summary(
    by = trt,
    type = all_continuous() ~ "continuous2",
    statistic = all_continuous() ~ c("{median} ({p25}, {p75})", "{min} - {max}"),
  ) |>
  add_stat_label(label = age ~ c("IQR", "Range"))
```

---

add\_vif

*Add Variance Inflation Factor*

---

### Description

Add the variance inflation factor (VIF) or generalized VIF (GVIF) to the regression table. Function uses `car::vif()` to calculate the VIF.

**Usage**

```
add_vif(x, statistic = NULL, estimate_fun = label_style_sigfig(digits = 2))
```

**Arguments**

`x` 'tbl\_regression' object

`statistic` "VIF" (variance inflation factors, for models with no categorical terms) or one of/combination of "GVIF" (generalized variance inflation factors), "aGVIF" 'adjusted GVIF, i.e.  $GVIF^{[1/(2*df)]}$  and/or "df" (degrees of freedom). See `car::vif()` for details.

`estimate_fun` Default is `label_style_sigfig(digits = 2)`.

**See Also**

Review [list](#), [formula](#), and [selector syntax](#) used throughout `gtsummary`

**Examples**

```
# Example 1 -----
lm(age ~ grade + marker, trial) |>
  tbl_regression() |>
  add_vif()

# Example 2 -----
lm(age ~ grade + marker, trial) |>
  tbl_regression() |>
  add_vif(c("aGVIF", "df"))
```

---

assign\_summary\_digits *Assign Default Digits*

---

**Description**

Used to assign the default formatting for variables summarized with `tbl_summary()`.

**Usage**

```
assign_summary_digits(data, statistic, type, digits = NULL)
```

**Arguments**

`data` (data.frame)  
a data frame

`statistic` (named list)  
a named list; notably, *not* a [formula-list-selector](#)



type (named list)  
a named list; notably, *not* a [formula-list-selector](#)

digits (named list)  
a named list; notably, *not* a [formula-list-selector](#). Default is NULL

**Value**

a named list

**Examples**

```
assign_summary_digits(
  mtcars,
  statistic = list(mpg = "{mean}"),
  type = list(mpg = "continuous")
)
```

---

assign\_summary\_type *Assign Default Summary Type*

---

**Description**

Function inspects data and assigns a summary type when not specified in the type argument.

**Usage**

```
assign_summary_type(data, variables, value, type = NULL, cat_threshold = 10L)
```

**Arguments**

data (data.frame)  
a data frame

variables (character)  
character vector of column names in data

value (named list)  
named list of values to show for dichotomous variables, where the names are the variables

type (named list)  
named list of summary types, where names are the variables

cat\_threshold (integer)  
for base R numeric classes with fewer levels than this threshold will default to a categorical summary. Default is 10L

**Value**

named list

**Examples**

```
assign_summary_type(
  data = trial,
  variables = c("age", "grade", "response"),
  value = NULL
)
```

---

 assign\_tests

*Assign Test*


---

**Description**

This function is used to assign default tests for `add_p()` and `add_difference()`.

**Usage**

```
assign_tests(x, ...)
```

```
## S3 method for class 'tbl_summary'
assign_tests(
  x,
  include,
  by = x$inputs$by,
  test = NULL,
  group = NULL,
  adj.vars = NULL,
  summary_type = x$inputs$type,
  calling_fun = c("add_p", "add_difference"),
  ...
)
```

```
## S3 method for class 'tbl_svysummary'
assign_tests(
  x,
  include,
  by = x$inputs$by,
  test = NULL,
  group = NULL,
  adj.vars = NULL,
  summary_type = x$inputs$type,
  calling_fun = c("add_p", "add_difference"),
  ...
)
```

```
## S3 method for class 'tbl_continuous'
assign_tests(x, include, by, cont_variable, test = NULL, group = NULL, ...)
```

```
## S3 method for class 'tbl_survfit'
assign_tests(x, include, test = NULL, ...)
```

### Arguments

x	(gtsummary) a table of class 'gtsummary'
...	Passed to <code>rlang::abort()</code> , <code>rlang::warn()</code> or <code>rlang::inform()</code> .
include	(character) Character vector of column names to assign a default tests.
by	(string) a single stratifying column name
test	(named list) a named list of tests.
group	(string) a variable name indicating the grouping column for correlated data. Default is NULL.
adj.vars	(character) Variables to include in adjusted calculations (e.g. in ANCOVA models).
summary_type	(named list) named list of summary types
calling_fun	(string) Must be one of 'add_p' and 'add_difference'. Depending on the context, different defaults are set.
cont_variable	(string) a column name of the continuous summary variable in <code>tbl_continuous()</code>

### Value

A table of class 'gtsummary'

### Examples

```
trial |>
  tbl_summary(
    by = trt,
    include = c(age, stage)
  ) |>
  assign_tests(include = c("age", "stage"), calling_fun = "add_p")
```

---

as_flex_table	<i>Convert gtsummary object to a flextable object</i>
---------------	---

---

### Description

Function converts a gtsummary object to a flextable object. A user can use this function if they wish to add customized formatting available via the flextable functions. The flextable output is particularly useful when combined with R markdown with Word output, since the gt package does not support Word.

### Usage

```
as_flex_table(x, include = everything(), return_calls = FALSE, ...)
```

### Arguments

x	(gtsummary) An object of class "gtsummary"
include	Commands to include in output. Input may be a vector of quoted or unquoted names. tidyselect and gtsummary select helper functions are also accepted. Default is everything().
return_calls	Logical. Default is FALSE. If TRUE, the calls are returned as a list of expressions.
...	Not used

### Details

The as\_flex\_table() function supports bold and italic markdown syntax in column headers and spanning headers ('\*\*' and '\_' only). Text wrapped in double stars ('\*\*bold\*\*') will be made bold, and text between single underscores ('\_italic\_') will be made italic. No other markdown syntax is supported and the double-star and underscore cannot be combined. To further style your table, you may convert the table to flextable with as\_flex\_table(), then utilize any of the flextable functions.

### Value

A 'flextable' object

### Author(s)

Daniel D. Sjoberg

**Examples**

```

trial |>
  select(trt, age, grade) |>
  tbl_summary(by = trt) |>
  add_p() |>
  as_flex_table()

```

as\_gt

*Convert gtsummary object to gt***Description**

Function converts a gtsummary object to a "gt\_tbl" object, that is, a table created with `gt::gt()`. Function is used in the background when the results are printed or knit. A user can use this function if they wish to add customized formatting available via the [gt package](#).

**Usage**

```
as_gt(x, include = everything(), return_calls = FALSE, ...)
```

**Arguments**

x	(gtsummary) An object of class "gtsummary"
include	Commands to include in output. Input may be a vector of quoted or unquoted names. tidyselect and gtsummary select helper functions are also accepted. Default is everything().
return_calls	Logical. Default is FALSE. If TRUE, the calls are returned as a list of expressions.
...	Arguments passed on to <code>gt::gt(...)</code>

**Value**

A gt\_tbl object

**Note**

As of 2024-08-15, line breaks (e.g. '\n') do not render properly for PDF output. For now, these line breaks are stripped when rendering to PDF with Quarto and R markdown.

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
trial |>
  tbl_summary(by = trt, include = c(age, grade, response)) |>
  as_gt()
```

---

as_hux_table	<i>Convert gtsummary object to a huxtable object</i>
--------------	--

---

**Description**

Function converts a gtsummary object to a huxtable object. A user can use this function if they wish to add customized formatting available via the huxtable functions. The huxtable package supports output to PDF via LaTeX, as well as HTML and Word.

**Usage**

```
as_hux_table(
  x,
  include = everything(),
  return_calls = FALSE,
  strip_md_bold = FALSE
)

as_hux_xlsx(x, file, include = everything(), bold_header_rows = TRUE)
```

**Arguments**

x	(gtsummary) An object of class "gtsummary"
include	Commands to include in output. Input may be a vector of quoted or unquoted names. tidyselect and gtsummary select helper functions are also accepted. Default is everything().
return_calls	Logical. Default is FALSE. If TRUE, the calls are returned as a list of expressions.
strip_md_bold	<b>[Deprecated]</b>
file	File path for the output.
bold_header_rows	(scalar logical) logical indicating whether to bold header rows. Default is TRUE

**Value**

A {huxtable} object

## Excel Output

Use the `as_hux_xlsx()` function to save a copy of the table in an excel file. The file is saved using `huxtable::quick_xlsx()`.

## Author(s)

David Hugh-Jones, Daniel D. Sjöberg

## Examples

```
trial |>
  tbl_summary(by = trt, include = c(age, grade)) |>
  add_p() |>
  as_hux_table()
```

---

 as\_kable

---

*Convert gtsummary object to a kable object*


---

## Description

Output from `knitr::kable()` is less full featured compared to summary tables produced with `gt`. For example, kable summary tables do not include indentation, footnotes, or spanning header rows.

Line breaks (`\n`) are removed from column headers and table cells.

## Usage

```
as_kable(x, ..., include = everything(), return_calls = FALSE)
```

## Arguments

<code>x</code>	(gtsummary) Object created by a function from the gtsummary package (e.g. <code>tbl_summary</code> or <code>tbl_regression</code> )
<code>...</code>	Additional arguments passed to <code>knitr::kable()</code>
<code>include</code>	Commands to include in output. Input may be a vector of quoted or unquoted names. <code>tidyselect</code> and <code>gtsummary</code> select helper functions are also accepted. Default is <code>everything()</code> .
<code>return_calls</code>	Logical. Default is FALSE. If TRUE, the calls are returned as a list of expressions.

## Details

Tip: To better distinguish variable labels and level labels when indenting is not supported, try `bold_labels()` or `italicize_levels()`.

**Value**

A knitr\_kable object

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
trial |>
  tbl_summary(by = trt) |>
  bold_labels() |>
  as_kable()
```

---

as\_kable\_extra

*Convert gtsummary object to a kableExtra object*

---

**Description**

Function converts a gtsummary object to a knitr\_kable + kableExtra object. This allows the customized formatting available via knitr::kable() and {kableExtra}; as\_kable\_extra() supports arguments in knitr::kable(). as\_kable\_extra() output via gtsummary supports bold and italic cells for table bodies. Users are encouraged to leverage as\_kable\_extra() for enhanced pdf printing; for html output options there is better support via as\_gt().

**Usage**

```
as_kable_extra(
  x,
  escape = FALSE,
  format = NULL,
  ...,
  include = everything(),
  addtl_fmt = TRUE,
  return_calls = FALSE
)
```

**Arguments**

x (gtsummary)  
Object created by a function from the gtsummary package (e.g. [tbl\\_summary](#) or [tbl\\_regression](#))

format, escape, ...  
arguments passed to knitr::kable(). Default is escape = FALSE, and the format is auto-detected.



include	Commands to include in output. Input may be a vector of quoted or unquoted names. tidyselect and gtsummary select helper functions are also accepted. Default is everything().
addtl_fmt	logical indicating whether to include additional formatting. Default is TRUE. This is primarily used to escape special characters, convert markdown to LaTeX, and remove line breaks from the footnote.
return_calls	Logical. Default is FALSE. If TRUE, the calls are returned as a list of expressions.

**Value**

A {kableExtra} table

**PDF/LaTeX**

This section shows options intended for use with `output: pdf_document` in yaml of `.Rmd`.

When the default values of `as_kable_extra(escape = FALSE, addtl_fmt = TRUE)` are utilized, the following formatting occurs.

- Markdown bold, italic, and underline syntax in the headers, spanning headers, caption, and footnote will be converted to escaped LaTeX code
- Special characters in the table body, headers, spanning headers, caption, and footnote will be escaped with `.escape_latex()` or `.escape_latex2()`
- The `"\n"` symbol will be recognized as a line break in the table headers, spanning headers, caption, and the table body
- The `"\n"` symbol is removed from the footnotes

To suppress *these* additional formats, set `as_kable_extra(addtl_fmt = FALSE)`

Additional styling is available with `kableExtra::kable_styling()` as shown in Example 2, which implements row striping and repeated column headers in the presence of page breaks.

**HTML**

This section discusses options intended for use with `output: html_document` in yaml of `.Rmd`.

When the default values of `as_kable_extra(escape = FALSE, addtl_fmt = TRUE)` are utilized, the following formatting occurs.

- The default markdown syntax in the headers and spanning headers is removed
- Special characters in the table body, headers, spanning headers, caption, and footnote will be escaped with `.escape_html()`
- The `"\n"` symbol is removed from the footnotes

To suppress the additional formatting, set `as_kable_extra(addtl_fmt = FALSE)`

**Author(s)**

Daniel D. Sjoberg

**Examples**

```

# basic gtsummary tbl to build upon
as_kable_extra_base <-
  trial |>
  tbl_summary(by = trt, include = c(age, stage)) |>
  bold_labels()

# Example 1 (PDF via LaTeX) -----
# add linebreak in table header with '\n'
as_kable_extra_ex1_pdf <-
  as_kable_extra_base |>
  modify_header(all_stat_cols() ~ "**{level}** \n*N = {n}*") |>
  as_kable_extra()

# Example 2 (PDF via LaTeX) -----
# additional styling in `knitr::kable()` and with
# call to `kableExtra::kable_styling()`
as_kable_extra_ex2_pdf <-
  as_kable_extra_base |>
  as_kable_extra(
    booktabs = TRUE,
    longtable = TRUE,
    linesep = ""
  ) |>
  kableExtra::kable_styling(
    position = "left",
    latex_options = c("striped", "repeat_header"),
    stripe_color = "gray!15"
  )

```

---

as\_tibble.gtsummary    *Convert gtsummary object to a tibble*

---

**Description**

Function converts a gtsummary object to a tibble.

**Usage**

```

## S3 method for class 'gtsummary'
as_tibble(
  x,
  include = everything(),
  col_labels = TRUE,
  return_calls = FALSE,
  fmt_missing = FALSE,
  ...
)

```

```
## S3 method for class 'gtsummary'
as.data.frame(...)
```

### Arguments

x	(gtsummary) An object of class "gtsummary"
include	Commands to include in output. Input may be a vector of quoted or unquoted names. tidyselect and gtsummary select helper functions are also accepted. Default is everything().
col_labels	(scalar logical) Logical argument adding column labels to output tibble. Default is TRUE.
return_calls	Logical. Default is FALSE. If TRUE, the calls are returned as a list of expressions.
fmt_missing	(scalar logical) Logical argument adding the missing value formats.
...	Arguments passed on to gt::gt(...)

### Value

a **tibble**

### Author(s)

Daniel D. Sjoberg

### Examples

```
tbl <-
  trial |>
  tbl_summary(by = trt, include = c(age, grade, response))

as_tibble(tbl)

# without column labels
as_tibble(tbl, col_labels = FALSE)
```

---

bold\_italicize\_labels\_levels

*Bold or Italicize*

---

### Description

Bold or italicize labels or levels in gtsummary tables

**Usage**

```
bold_labels(x)

italicize_labels(x)

bold_levels(x)

italicize_levels(x)

## S3 method for class 'gtsummary'
bold_labels(x)

## S3 method for class 'gtsummary'
bold_levels(x)

## S3 method for class 'gtsummary'
italicize_labels(x)

## S3 method for class 'gtsummary'
italicize_levels(x)

## S3 method for class 'tbl_cross'
bold_labels(x)

## S3 method for class 'tbl_cross'
bold_levels(x)

## S3 method for class 'tbl_cross'
italicize_labels(x)

## S3 method for class 'tbl_cross'
italicize_levels(x)
```

**Arguments**

x (gtsummary) An object of class 'gtsummary'

**Value**

Functions return the same class of gtsummary object supplied

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
tbl_summary(trial, include = c("trt", "age", "response")) |>
```

```
bold_labels() |>
bold_levels() |>
italicize_labels() |>
italicize_levels()
```

---

bold_p	<i>Bold significant p-values</i>
--------	----------------------------------

---

### Description

Bold values below a chosen threshold (e.g.  $<0.05$ ) in a gtsummary tables.

### Usage

```
bold_p(x, t = 0.05, q = FALSE)
```

### Arguments

x	(gtsummary) Object created using gtsummary functions
t	(scalar numeric) Threshold below which values will be bold. Default is 0.05.
q	(scalar logical) When TRUE will bold the q-value column rather than the p-value. Default is FALSE.

### Author(s)

Daniel D. Sjoberg, Esther Drill

### Examples

```
# Example 1 -----
trial |>
tbl_summary(by = trt, include = c(response, marker, trt), missing = "no") |>
add_p() |>
bold_p(t = 0.1)

# Example 2 -----
glm(response ~ trt + grade, trial, family = binomial(link = "logit")) |>
tbl_regression(exponentiate = TRUE) |>
bold_p(t = 0.65)
```

---

brdg\_continuous      *Continuous Summary Table Bridges*


---

### Description

Bridge function for converting `tbl_continuous()` cards to basic `gtsummary` objects. This bridge function converts the 'cards' object to a format suitable to pass to `brdg_summary()`: no `pier_*()` functions required.

### Usage

```
brdg_continuous(cards, by = NULL, statistic, include, variable, type)
```

### Arguments

cards	(card) An ARD object of class "card" typically created with <code>cards::ard_*()</code> functions.
by	(string) string indicating the stratifying column
statistic	(named list) named list of summary statistic names
include	( <a href="#">tidy-select</a> ) Variables to include in the summary table. Default is <code>everything()</code> .
variable	( <a href="#">tidy-select</a> ) A single column from data. Variable name of the continuous column to be summarized.
type	(named list) named list of summary types

### Value

a `gtsummary` object

### Examples

```
library(cards)

bind_ard(
  # the primary ARD with the results
  ard_continuous(trial, by = grade, variables = age),
  # add missing and attributes ARD
  ard_missing(trial, by = grade, variables = age),
  ard_attributes(trial, variables = c(grade, age))
) |>
  # adding the column name
  dplyr::mutate(
```

```

      gts_column =
        ifelse(!context %in% "attributes", "stat_0", NA_character_)
    ) |>
  brdg_continuous(
    variable = "age",
    include = "grade",
    statistic = list(grade = "{median} ({p25}, {p75})"),
    type = list(grade = "categorical")
  ) |>
  as_tibble()

```

---

brdg\_hierarchical      *Hierarchy table bridge*

---

## Description

Bridge function for converting `tbl_hierarchical()` (and similar) cards to basic `gtsummary` objects. All bridge functions begin with prefix `brdg_*()`.

This file also contains helper functions for constructing the bridge, referred to as the piers (supports for a bridge) and begin with `pier_*()`.

- `brdg_hierarchical()`: The bridge function ingests an ARD data frame and returns a `gtsummary` table that includes `.$table_body` and a basic `.$table_styling`. The `.$table_styling$header` data frame includes the header statistics. Based on context, this function adds a column to the ARD data frame named `"gts_column"`. This column is used during the reshaping in the `pier_*()` functions defining column names.
- `pier_*()`: these functions accept a cards tibble and returns a tibble that is a piece of the `.$table_body`. Typically these will be stacked to construct the final table body data frame. The ARD object passed here will have two primary parts: the calculated summary statistics and the attributes ARD. The attributes ARD is used for labeling. The ARD data frame passed to this function must include a `"gts_column"` column, which is added in `brdg_hierarchical()`.

## Usage

```

brdg_hierarchical(
  cards,
  variables,
  by,
  include,
  statistic,
  type,
  overall_row,
  count,
  is_ordered,
  label
)

pier_summary_hierarchical(cards, variables, include, statistic)

```

**Arguments**

cards	(card) an ARD object of class "card" created with <code>cards::ard_hierarchical_stack()</code> .
variables	(character) character list of hierarchy variables.
by	(string) string indicating the stratifying column.
include	(character) character list of hierarchy variables to include summary statistics for.
statistic	(named list) named list of summary statistic names.
type	(named list) named list of summary types.
overall_row	(scalar logical) whether an overall summary row should be included at the top of the table. The default is FALSE.
count	(scalar logical) whether <code>tbl_hierarchical_count()</code> (TRUE) or <code>tbl_hierarchical()</code> (FALSE) is being applied.
is_ordered	(scalar logical) whether the last variable in <code>variables</code> is ordered.
label	(named list) named list of hierarchy variable labels.

**Value**

a `gtsummary` object

**See Also**

Review [list](#), [formula](#), and [selector syntax](#) used throughout `gtsummary`

---

brdg\_summary

*Summary table bridge*


---

**Description**

Bridge function for converting `tbl_summary()` (and similar) cards to basic `gtsummary` objects. All bridge functions begin with prefix `brdg_*()`.

This file also contains helper functions for constructing the bridge, referred to as the piers (supports for a bridge) and begin with `pier_*()`.



- `brdg_summary()`: The bridge function ingests an ARD data frame and returns a `gtsummary` table that includes `.$table_body` and a basic `.$table_styling`. The `.$table_styling$header` data frame includes the header statistics. Based on context, this function adds a column to the ARD data frame named `"gts_column"`. This column is used during the reshaping in the `pier_*`() functions defining column names.
- `pier_*`(): these functions accept a `cards` tibble and returns a tibble that is a piece of the `.$table_body`. Typically these will be stacked to construct the final table body data frame. The ARD object passed here will have two primary parts: the calculated summary statistics and the attributes ARD. The attributes ARD is used for labeling. The ARD data frame passed to this function must include a `"gts_column"` column, which is added in `brdg_summary()`.

### Usage

```
brdg_summary(
  cards,
  variables,
  type,
  statistic,
  by = NULL,
  missing = "no",
  missing_stat = "{N_miss}",
  missing_text = "Unknown"
)

pier_summary_dichotomous(cards, variables, statistic)

pier_summary_categorical(cards, variables, statistic)

pier_summary_continuous2(cards, variables, statistic)

pier_summary_continuous(cards, variables, statistic)

pier_summary_missing_row(
  cards,
  variables,
  missing = "no",
  missing_stat = "{N_miss}",
  missing_text = "Unknown"
)
```

### Arguments

<code>cards</code>	(card) An ARD object of class "card" typically created with <code>cards::ard_*</code> () functions.
<code>variables</code>	(character) character list of variables
<code>type</code>	(named list) named list of summary types

**statistic** (named list)  
 named list of summary statistic names

**by** (string)  
 string indicating the stratifying column

**missing, missing\_text, missing\_stat**  
 Arguments dictating how and if missing values are presented:
 

- **missing**: must be one of `c("ifany", "no", "always")`
- **missing\_text**: string indicating text shown on missing row. Default is "Unknown"
- **missing\_stat**: statistic to show on missing row. Default is "{N\_miss}". Possible values are `N_miss, N_obs, N_nonmiss, p_miss, p_nonmiss`.

### Value

a gtsummary object

### Examples

```

library(cards)

# first build ARD data frame
cards <-
  ard_stack(
    mtcars,
    ard_continuous(variables = c("mpg", "hp")),
    ard_categorical(variables = "cyl"),
    ard_dichotomous(variables = "am"),
    .missing = TRUE,
    .attributes = TRUE
  ) |>
  # this column is used by the `pier_*()` functions
  dplyr::mutate(gts_column = ifelse(context == "attributes", NA, "stat_0"))

brdg_summary(
  cards = cards,
  variables = c("cyl", "am", "mpg", "hp"),
  type =
    list(
      cyl = "categorical",
      am = "dichotomous",
      mpg = "continuous",
      hp = "continuous2"
    ),
  statistic =
    list(
      cyl = "{n} / {N}",
      am = "{n} / {N}",
      mpg = "{mean} ({sd})",
      hp = c("{median} ({p25}, {p75})", "{mean} ({sd})")
    )
) |>

```

```

  as_tibble()

  pier_summary_dichotomous(
    cards = cards,
    variables = "am",
    statistic = list(am = "{n} ({p})")
  )

  pier_summary_categorical(
    cards = cards,
    variables = "cyl",
    statistic = list(cyl = "{n} ({p})")
  )

  pier_summary_continuous2(
    cards = cards,
    variables = "hp",
    statistic = list(hp = c("{median}", "{mean}"))
  )

  pier_summary_continuous(
    cards = cards,
    variables = "mpg",
    statistic = list(mpg = "{median}")
  )

```

---

brdg\_wide\_summary      *Wide summary table bridge*


---

## Description

Bridge function for converting `tbl_wide_summary()` (and similar) cards to basic `gtsummary` objects. All bridge functions begin with prefix `brdg_*()`.

## Usage

```
brdg_wide_summary(cards, variables, statistic, type)
```

## Arguments

cards	(card) An ARD object of class "card" typically created with <code>cards::ard_*()</code> functions.
variables	(character) character list of variables
statistic	(named list) named list of summary statistic names
type	(named list) named list of summary types

**Value**

a gtsummary object

**Examples**

```
library(cards)

bind_ard(
  ard_continuous(trial, variables = c(age, marker)),
  ard_attributes(trial, variables = c(age, marker))
) |>
  brdg_wide_summary(
    variables = c("age", "marker"),
    statistic = list(age = c("{mean}", "{sd}"), marker = c("{mean}", "{sd}")),
    type = list(age = "continuous", marker = "continuous")
  )
```

---

 combine\_terms

*Combine terms*


---

**Description**

The function combines terms from a regression model, and replaces the terms with a single row in the output table. The p-value is calculated using `stats::anova()`.

**Usage**

```
combine_terms(x, formula_update, label = NULL, quiet, ...)
```

**Arguments**

x	(tbl_regression) A tbl_regression object
formula_update	(formula) formula update passed to the <code>stats::update()</code> . This updated formula is used to construct a reduced model, and is subsequently passed to <code>stats::anova()</code> to calculate the p-value for the group of removed terms. See the <code>stats::update()</code> function's <code>formula. =</code> argument for proper syntax.
label	(string) Optional string argument labeling the combined rows
quiet	<b>[Deprecated]</b>
...	Additional arguments passed to <code>stats::anova</code>

**Value**

tbl\_regression object

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
# Logistic Regression Example, LRT p-value
glm(response ~ marker + I(marker^2) + grade,
     trial[c("response", "marker", "grade")] |> na.omit(), # keep complete cases only!
     family = binomial) |>
tbl_regression(label = grade ~ "Grade", exponentiate = TRUE) |>
# collapse non-linear terms to a single row in output using anova
combine_terms(
  formula_update = . ~ . - marker - I(marker^2),
  label = "Marker (non-linear terms)",
  test = "LRT"
)
```

---

 custom\_tidiers

*Custom tidiers*


---

**Description**

**[Maturing]** Collection of tidiers that can be utilized in gtsummary. See details below.

**Usage**

```
tidy_standardize(
  x,
  exponentiate = FALSE,
  conf.level = 0.95,
  conf.int = TRUE,
  ...,
  quiet = FALSE
)
```

```
tidy_bootstrap(
  x,
  exponentiate = FALSE,
  conf.level = 0.95,
  conf.int = TRUE,
  ...,
  quiet = FALSE
)
```

```
tidy_robust(
  x,
```

```

  exponentiate = FALSE,
  conf.level = 0.95,
  conf.int = TRUE,
  vcov = NULL,
  vcov_args = NULL,
  ...,
  quiet = FALSE
)

pool_and_tidy_mice(x, pool.args = NULL, ..., quiet = FALSE)

tidy_gam(x, conf.int = FALSE, exponentiate = FALSE, conf.level = 0.95, ...)

tidy_wald_test(x, tidy_fun = NULL, ...)

```

### Arguments

x	(model) Regression model object
exponentiate	(scalar logical) Logical indicating whether to exponentiate the coefficient estimates. Default is FALSE.
conf.level	(scalar real) Confidence level for confidence interval/credible interval. Defaults to 0.95.
conf.int	(scalar logical) Logical indicating whether or not to include a confidence interval in the output. Default is TRUE.
...	Arguments passed to method; <ul style="list-style-type: none"> <li>• pool_and_tidy_mice(): mice::tidy(x, ...)</li> <li>• tidy_standardize(): parameters::standardize_parameters(x, ...)</li> <li>• tidy_bootstrap(): parameters::bootstrap_parameters(x, ...)</li> <li>• tidy_robust(): parameters::model_parameters(x, ...)</li> </ul>
quiet	<b>[Deprecated]</b>
vcov, vcov_args	Arguments passed to parameters::model_parameters(). At least one of these arguments <b>must</b> be specified.
pool.args	(named list) Named list of arguments passed to mice::pool() in pool_and_tidy_mice(). Default is NULL
tidy_fun	(function) Tidier function for the model. Default is to use broom::tidy(). If an error occurs, the tidying of the model is attempted with parameters::model_parameters(), if installed.

### Regression Model Tidiers

These tidiers are passed to tbl\_regression() and tbl\_uvregression() to obtain modified results.

- `tidy_standardize()` tidier to report standardized coefficients. The `parameters` package includes a wonderful function to estimate standardized coefficients. The tidier uses the output from `parameters::standardize_parameters()`, and merely takes the result and puts it in `broom::tidy()` format.
- `tidy_bootstrap()` tidier to report bootstrapped coefficients. The `parameters` package includes a wonderful function to estimate bootstrapped coefficients. The tidier uses the output from `parameters::bootstrap_parameters(test = "p")`, and merely takes the result and puts it in `broom::tidy()` format.
- `tidy_robust()` tidier to report robust standard errors, confidence intervals, and p-values. The `parameters` package includes a wonderful function to calculate robust standard errors, confidence intervals, and p-values. The tidier uses the output from `parameters::model_parameters()`, and merely takes the result and puts it in `broom::tidy()` format. To use this function with `tbl_regression()`, pass a function with the arguments for `tidy_robust()` populated.
- `pool_and_tidy_mice()` tidier to report models resulting from multiply imputed data using the `mice` package. Pass the `mice` model object *before* the model results have been pooled. See example.

### Other Tidiers

- `tidy_wald_test()` tidier to report Wald p-values, wrapping the `aod::wald.test()` function. Use this tidier with `add_global_p(anova_fun = tidy_wald_test)`

### Examples

```
# Example 1 -----
mod <- lm(age ~ marker + grade, trial)

tbl_std <- tbl_regression(mod, tidy_fun = tidy_standardize)
tbl <- tbl_regression(mod)

tidy_standardize_ex1 <-
  tbl_merge(
    list(tbl_std, tbl),
    tab_spanner = c("**Standardized Model**", "**Original Model**")
  )

# Example 2 -----
# use "posthoc" method for coef calculation
tbl_regression(mod, tidy_fun = \(x, ...) tidy_standardize(x, method = "posthoc", ...))

# Example 3 -----
# Multiple Imputation using the mice package
set.seed(1123)
pool_and_tidy_mice_ex3 <-
  suppressWarnings(mice::mice(trial, m = 2)) |>
  with(lm(age ~ marker + grade)) |>
  tbl_regression()
```

---

gather\_ard

*Extract ARDs*


---

**Description**

Extract the ARDs from a gtsummary table. If needed, results may be combined with cards: `:bind_ard()`.

**Usage**

```
gather_ard(x)
```

**Arguments**

x (gtsummary)  
a gtsummary table.

**Value**

list

**Examples**

```
tbl_summary(trial, by = trt, include = age) |>
  add_overall() |>
  add_p() |>
  gather_ard()

glm(response ~ trt, data = trial, family = binomial()) |>
  tbl_regression() |>
  gather_ard()
```

---

inline\_text.gtsummary *Report statistics from summary tables inline*


---

**Description**

Report statistics from summary tables inline

**Usage**

```
## S3 method for class 'gtsummary'
inline_text(x, variable, level = NULL, column = NULL, pattern = NULL, ...)
```



**Arguments**

x	(gtsummary) gtsummary object
variable	( <a href="#">tidy-select</a> ) A single variable name of statistic to present
level	(string) Level of the variable to display for categorical variables. Default is NULL
column	( <a href="#">tidy-select</a> ) Column name to return from x\$table_body.
pattern	(string) String indicating the statistics to return. Uses <a href="#">glue::glue()</a> formatting. Default is NULL
...	These dots are for future extensions and must be empty.

**Value**

A string

**column + pattern**

Some gtsummary tables report multiple statistics in a single cell, e.g. "{mean} ({sd})" in `tbl_summary()` or `tbl_svsummary()`. We often need to report just the mean or the SD, and that can be accomplished by using both the `column=` and `pattern=` arguments. When both of these arguments are specified, the `column` argument selects the column to report statistics from, and the `pattern` argument specifies which statistics to report, e.g. `inline_text(x, column = "stat_1", pattern = "{mean}")` reports just the mean from a `tbl_summary()`. *This is not supported for all tables.*

---

```
inline_text.tbl_continuous
```

*Report statistics from summary tables inline*

---

**Description**

Extracts and returns statistics from a `tbl_continuous()` object for inline reporting in an R markdown document. Detailed examples in the [inline\\_text vignette](#)

**Usage**

```
## S3 method for class 'tbl_continuous'
inline_text(
  x,
  variable,
  column = NULL,
  level = NULL,
  pattern = NULL,
```

```

  pvalue_fun = label_style_pvalue(prepend_p = TRUE),
  ...
)

```

### Arguments

x	(tbl_continuous) Object created from tbl_continuous()
variable	(tidy-select) A single variable name of statistic to present
column	(tidy-select) Column name to return from x\$table_body. Can also pass the level of a by variable.
level	(string) Level of the variable to display for categorical variables. Default is NULL
pattern	(string) String indicating the statistics to return. Uses glue::glue() formatting. Default is NULL
pvalue_fun	(function) Function to round and format p-values. Default is label_style_pvalue(). The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. pvalue_fun = label_style_pvalue(digits = 2)).
...	These dots are for future extensions and must be empty.

### Value

A string reporting results from a gtsummary table

### Author(s)

Daniel D. Sjoberg

### Examples

```

t1 <- trial |>
  tbl_summary(by = trt, include = grade) |>
  add_p()

inline_text(t1, variable = grade, level = "I", column = "Drug A", pattern = "{n}/{N} ({p}%)")
inline_text(t1, variable = grade, column = "p.value")

```

---

inline\_text.tbl\_cross *Report statistics from cross table inline*

---

### Description

**[Maturing]** Extracts and returns statistics from a `tbl_cross` object for inline reporting in an R markdown document. Detailed examples in the [inline\\_text vignette](#)

### Usage

```
## S3 method for class 'tbl_cross'
inline_text(
  x,
  col_level,
  row_level = NULL,
  pvalue_fun = label_style_pvalue(prepend_p = TRUE),
  ...
)
```

### Arguments

<code>x</code>	(tbl_cross) A <code>tbl_cross</code> object
<code>col_level</code>	(string) Level of the column variable to display. Can also specify "p.value" for the p-value and "stat_0" for Total column.
<code>row_level</code>	(string) Level of the row variable to display.
<code>pvalue_fun</code>	(function) Function to round and format p-values. Default is <code>label_style_pvalue()</code> . The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. <code>pvalue_fun = label_style_pvalue(digits = 2)</code> ).
<code>...</code>	These dots are for future extensions and must be empty.

### Value

A string reporting results from a `gtsummary` table

### Examples

```
tbl_cross <-
  tbl_cross(trial, row = trt, col = response) %>%
  add_p()

inline_text(tbl_cross, row_level = "Drug A", col_level = "1")
inline_text(tbl_cross, row_level = "Total", col_level = "1")
inline_text(tbl_cross, col_level = "p.value")
```

---

 inline\_text.tbl\_regression

*Report statistics from regression summary tables inline*


---

## Description

Takes an object with class `tbl_regression`, and the location of the statistic to report and returns statistics for reporting inline in an R markdown document. Detailed examples in the [inline\\_text vignette](#)

## Usage

```
## S3 method for class 'tbl_regression'
inline_text(
  x,
  variable,
  level = NULL,
  pattern = "{estimate} ({conf.level*100}% CI {conf.low}, {conf.high}); {p.value})",
  estimate_fun = x$inputs$estimate_fun,
  pvalue_fun = label_style_pvalue(prepend_p = TRUE),
  ...
)
```

## Arguments

<code>x</code>	(tbl_regression) Object created by <code>tbl_regression()</code>
<code>variable</code>	(tidy-select) A single variable name of statistic to present
<code>level</code>	(string) Level of the variable to display for categorical variables. Default is NULL
<code>pattern</code>	(string) String indicating the statistics to return. Uses <code>glue::glue()</code> formatting. Default is "{estimate} ({conf.level }%\ CI {conf.low}, {conf.high}); {p.value})". All columns from <code>x\$table_body</code> are available to print as well as the confidence level ( <code>conf.level</code> ). See below for details.
<code>estimate_fun</code>	(function) Function to style model coefficient estimates. Columns 'estimate', 'conf.low', and 'conf.high' are formatted. Default is <code>x\$inputs\$estimate_fun</code>
<code>pvalue_fun</code>	function to style p-values and/or q-values. Default is <code>label_style_pvalue(prepend_p = TRUE)</code>
<code>...</code>	These dots are for future extensions and must be empty.

## Value

A string reporting results from a gtsummary table

**pattern argument**

The following items (and more) are available to print. Use `print(x$table_body)` to print the table the estimates are extracted from.

- `{estimate}` coefficient estimate formatted with `'estimate_fun'`
- `{conf.low}` lower limit of confidence interval formatted with `'estimate_fun'`
- `{conf.high}` upper limit of confidence interval formatted with `'estimate_fun'`
- `{p.value}` p-value formatted with `'pvalue_fun'`
- `{N}` number of observations in model
- `{label}` variable/variable level label

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
inline_text_ex1 <-
  glm(response ~ age + grade, trial, family = binomial(link = "logit")) %>%
  tbl_regression(exponentiate = TRUE)

inline_text(inline_text_ex1, variable = age)
inline_text(inline_text_ex1, variable = grade, level = "III")
```

---

```
inline_text.tbl_summary
```

*Report statistics from summary tables inline*

---

**Description**

Extracts and returns statistics from a `tbl_summary()` object for inline reporting in an R markdown document. Detailed examples in the [inline\\_text vignette](#)

**Usage**

```
## S3 method for class 'tbl_summary'
inline_text(
  x,
  variable,
  column = NULL,
  level = NULL,
  pattern = NULL,
  pvalue_fun = label_style_pvalue(prepend_p = TRUE),
  ...
)
```

```
## S3 method for class 'tbl_svsummary'
inline_text(
  x,
  variable,
  column = NULL,
  level = NULL,
  pattern = NULL,
  pvalue_fun = label_style_pvalue(prepend_p = TRUE),
  ...
)
```

### Arguments

x	(tbl_summary) Object created from <code>tbl_summary()</code> or <code>tbl_svsummary()</code>
variable	( <a href="#">tidy-select</a> ) A single variable name of statistic to present
column	( <a href="#">tidy-select</a> ) Column name to return from <code>x\$table_body</code> . Can also pass the level of a by variable.
level	(string) Level of the variable to display for categorical variables. Default is NULL
pattern	(string) String indicating the statistics to return. Uses <code>glue::glue()</code> formatting. Default is NULL
pvalue_fun	(function) Function to round and format p-values. Default is <code>label_style_pvalue()</code> . The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. <code>pvalue_fun = label_style_pvalue(digits = 2)</code> ).
...	These dots are for future extensions and must be empty.

### Value

A string reporting results from a `gtsummary` table

### Author(s)

Daniel D. Sjoberg

### Examples

```
t1 <- trial |>
  tbl_summary(by = trt, include = grade) |>
  add_p()

inline_text(t1, variable = grade, level = "I", column = "Drug A", pattern = "{n}/{N} ({p}%)")
inline_text(t1, variable = grade, column = "p.value")
```

---

 inline\_text.tbl\_survfit

*Report statistics from survfit tables inline*


---

## Description

### [Maturing]

Extracts and returns statistics from a `tbl_survfit` object for inline reporting in an R markdown document. Detailed examples in the [inline\\_text vignette](#)

## Usage

```
## S3 method for class 'tbl_survfit'
inline_text(
  x,
  variable = NULL,
  level = NULL,
  pattern = NULL,
  time = NULL,
  prob = NULL,
  column = NULL,
  estimate_fun = x$inputs$estimate_fun,
  pvalue_fun = label_style_pvalue(prepend_p = TRUE),
  ...
)
```

## Arguments

<code>x</code>	( <code>tbl_survfit</code> ) Object created from <code>tbl_survfit()</code>
<code>variable</code>	( <code>tidy-select</code> ) Variable name of statistic to present.
<code>level</code>	(string) Level of the variable to display for categorical variables. Can also specify the 'Unknown' row. Default is NULL
<code>pattern</code>	(string) String indicating the statistics to return.
<code>time, prob</code>	(numeric scalar) time or probability for which to return result
<code>column</code>	( <code>tidy-select</code> ) column to print from <code>x\$table_body</code> . Columns may be selected with <code>time</code> or <code>prob</code> arguments as well.
<code>estimate_fun</code>	(function) Function to round and format estimate and confidence limits. Default is the same function used in <code>tbl_survfit()</code>

pvalue_fun	(function) Function to round and format p-values. Default is label_style_pvalue(). The function must have a numeric vector input, and return a string that is the rounded/formatted p-value (e.g. pvalue_fun = label_style_pvalue(digits = 2)).
...	These dots are for future extensions and must be empty.

**Value**

A string reporting results from a gtsummary table

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
library(survival)

# fit survfit
fit1 <- survfit(Surv(ttdeath, death) ~ trt, trial)
fit2 <- survfit(Surv(ttdeath, death) ~ 1, trial)

# summarize survfit objects
tbl1 <-
  tbl_survfit(
    fit1,
    times = c(12, 24),
    label = ~"Treatment",
    label_header = "**{time} Month**"
  ) %>%
  add_p()

tbl2 <-
  tbl_survfit(
    fit2,
    probs = 0.5,
    label_header = "**Median Survival**"
  )

# report results inline
inline_text(tbl1, time = 24, level = "Drug B")
inline_text(tbl1, time = 24, level = "Drug B",
  pattern = "{estimate} [95% CI {conf.low}, {conf.high}]")
inline_text(tbl1, column = p.value)
inline_text(tbl2, prob = 0.5)
```



---

```
inline_text.tbl_uvregression
```

*Report statistics from regression summary tables inline*

---

## Description

Extracts and returns statistics from a table created by the `tbl_uvregression` function for inline reporting in an R markdown document. Detailed examples in the [inline\\_text vignette](#)

## Usage

```
## S3 method for class 'tbl_uvregression'
inline_text(
  x,
  variable,
  level = NULL,
  pattern = "{estimate} ({conf.level*100}% CI {conf.low}, {conf.high}); {p.value}",
  estimate_fun = x$inputs$estimate_fun,
  pvalue_fun = label_style_pvalue(prepend_p = TRUE),
  ...
)
```

## Arguments

<code>x</code>	(tbl_uvregression) Object created by <code>tbl_uvregression()</code>
<code>variable</code>	(tidy-select) A single variable name of statistic to present
<code>level</code>	(string) Level of the variable to display for categorical variables. Default is NULL
<code>pattern</code>	(string) String indicating the statistics to return. Uses <code>glue::glue()</code> formatting. Default is NULL
<code>estimate_fun</code>	(function) Function to style model coefficient estimates. Columns 'estimate', 'conf.low', and 'conf.high' are formatted. Default is <code>x\$inputs\$estimate_fun</code>
<code>pvalue_fun</code>	function to style p-values and/or q-values. Default is <code>label_style_pvalue(prepend_p = TRUE)</code>
<code>...</code>	These dots are for future extensions and must be empty.

## Value

A string reporting results from a gtsummary table

**pattern argument**

The following items (and more) are available to print. Use `print(x$table_body)` to print the table the estimates are extracted from.

- `{estimate}` coefficient estimate formatted with `'estimate_fun'`
- `{conf.low}` lower limit of confidence interval formatted with `'estimate_fun'`
- `{conf.high}` upper limit of confidence interval formatted with `'estimate_fun'`
- `{p.value}` p-value formatted with `'pvalue_fun'`
- `{N}` number of observations in model
- `{label}` variable/variable level label

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
inline_text_ex1 <-
  trial[c("response", "age", "grade")] %>%
  tbl_uvregression(
    method = glm,
    method.args = list(family = binomial),
    y = response,
    exponentiate = TRUE
  )

inline_text(inline_text_ex1, variable = age)
inline_text(inline_text_ex1, variable = grade, level = "III")
```

---

label\_style

*Style Functions*

---

**Description**

Similar to the `style_*()` family of functions, but these functions return a `style_*()` **function** rather than performing the styling.

**Usage**

```
label_style_number(
  digits = 0,
  big.mark = ifelse(decimal.mark == ",", " ", ", "),
  decimal.mark = getOption("OutDec"),
  scale = 1,
  prefix = "",
  suffix = "",
```

```
    ...
)

label_style_sigfig(
  digits = 2,
  scale = 1,
  big.mark = ifelse(decimal.mark == ",", " ", ","),
  decimal.mark = getOption("OutDec"),
  prefix = "",
  suffix = "",
  ...
)

label_style_pvalue(
  digits = 1,
  prepend_p = FALSE,
  big.mark = ifelse(decimal.mark == ",", " ", ","),
  decimal.mark = getOption("OutDec"),
  ...
)

label_style_ratio(
  digits = 2,
  big.mark = ifelse(decimal.mark == ",", " ", ","),
  decimal.mark = getOption("OutDec"),
  prefix = "",
  suffix = "",
  ...
)

label_style_percent(
  prefix = "",
  suffix = "",
  digits = 0,
  big.mark = ifelse(decimal.mark == ",", " ", ","),
  decimal.mark = getOption("OutDec"),
  ...
)
```

### Arguments

digits, big.mark, decimal.mark, scale, prepend\_p, prefix, suffix, ...  
arguments passed to the style\_\*( ) functions

### Value

a function

**See Also**

Other style tools: `style_sigfig()`

**Examples**

```
my_style <- label_style_number(digits = 1)
my_style(3.14)
```

---

modify

*Modify column headers, footnotes, and spanning headers*

---

**Description**

These functions assist with modifying the aesthetics/style of a table.

- `modify_header()` update column headers
- `modify_footnote()` update/add table footnotes
- `modify_spanning_header()` update/add spanning headers

The functions often require users to know the underlying column names. Run `show_header_names()` to print the column names to the console.

**Usage**

```
modify_header(x, ..., text_interpret = c("md", "html"), quiet, update)
```

```
modify_footnote(
  x,
  ...,
  abbreviation = FALSE,
  text_interpret = c("md", "html"),
  update,
  quiet
)
```

```
modify_spanning_header(x, ..., text_interpret = c("md", "html"), quiet, update)
```

```
show_header_names(x, include_example, quiet)
```

**Arguments**

x (gtsummary)  
A gtsummary object

... [dynamic-dots](#)  
 Used to assign updates to headers, spanning headers, and footnotes.  
 Use `modify_*(colname='new header/footer')` to update a single column.  
 Using a formula will invoke `tidyselect`, e.g. `modify_*(all_stat_cols() ~ "**{level}**")`. The dynamic dots allow syntax like `modify_header(x, !!!list(label = "Variable"))`. See examples below.  
 Use the `show_header_names()` to see the column names that can be modified.

`text_interpret` (string)  
 String indicates whether text will be interpreted with `gt::md()` or `gt::html()`.  
 Must be "md" (default) or "html".

`update, quiet` **[Deprecated]**

`abbreviation` (scalar logical)  
 Logical indicating if an abbreviation is being updated.

`include_example`  
**[Deprecated]**

**Value**

Updated `gtsummary` object

`tbl_summary()`, `tbl_svsummary()`, and `tbl_cross()`

When assigning column headers, footnotes, and spanning headers, you may use `{N}` to insert the number of observations. `tbl_svsummary` objects additionally have `{N_unweighted}` available.

When there is a stratifying `by=` argument present, the following fields are additionally available to stratifying columns: `{level}`, `{n}`, and `{p}` (`{n_unweighted}` and `{p_unweighted}` for `tbl_svsummary` objects)

Syntax follows `glue::glue()`, e.g. `all_stat_cols() ~ "**{level}**", N = {n}"`.

**tbl\_regression()**

When assigning column headers for `tbl_regression` tables, you may use `{N}` to insert the number of observations, and `{N_event}` for the number of events (when applicable).

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# create summary table
tbl <- trial |>
  tbl_summary(by = trt, missing = "no", include = c("age", "grade", "trt")) |>
  add_p()

# print the column names that can be modified
show_header_names(tbl)
```

```

# Example 1 -----
# updating column headers and footnote
tbl |>
  modify_header(label = "**Variable**", p.value = "**P**") |>
  modify_footnote(all_stat_cols() ~ "median (IQR) for Age; n (%) for Grade")

# Example 2 -----
# updating headers, remove all footnotes, add spanning header
tbl |>
  modify_header(all_stat_cols() ~ "**{level}**", N = {n} ({{style_percent(p)}}%)") |>
  modify_footnote(everything() ~ NA) |>
  modify_spanning_header(all_stat_cols() ~ "**Treatment Received**")

# Example 3 -----
# updating an abbreviation in table footnote
glm(response ~ age + grade, trial, family = binomial) |>
  tbl_regression(exponentiate = TRUE) |>
  modify_footnote(conf.low = "CI = Credible Interval", abbreviation = TRUE)

```

---

modify\_caption

*Modify table caption*

---

## Description

Captions are assigned based on output type.

- `gt::gt(caption=)`
- `flextable::set_caption(caption=)`
- `huxtable::set_caption(value=)`
- `knitr::kable(caption=)`

## Usage

```
modify_caption(x, caption, text_interpret = c("md", "html"))
```

## Arguments

<code>x</code>	(gtsummary) A gtsummary object
<code>caption</code>	(string) A string for the table caption/title
<code>text_interpret</code>	(string) String indicates whether text will be interpreted with <code>gt::md()</code> or <code>gt::html()</code> . Must be "md" (default) or "html".

## Value

Updated gtsummary object

**Examples**

```
trial |>
  tbl_summary(by = trt, include = c(marker, stage)) |>
  modify_caption(caption = "**Baseline Characteristics** N = {N}")
```

---

```
modify_column_alignment      Modify column alignment
```

---

**Description**

Update column alignment/justification in a gtsummary table.

**Usage**

```
modify_column_alignment(x, columns, align = c("left", "right", "center"))
```

**Arguments**

x	(gtsummary) gtsummary object
columns	( <a href="#">tidy-select</a> ) Selector of columns in x\$table_body
align	(string) String indicating alignment of column, must be one of c("left", "right", "center")

**Examples**

```
# Example 1 -----
lm(age ~ marker + grade, trial) %>%
  tbl_regression() %>%
  modify_column_alignment(columns = everything(), align = "left")
```

---

```
modify_column_hide      Modify hidden columns
```

---

**Description**

Use these functions to hide or unhide columns in a gtsummary table. Use `show_header_names(show_hidden=TRUE)` to print available columns to update.

**Usage**

```
modify_column_hide(x, columns)

modify_column_unhide(x, columns)
```

**Arguments**

x	(gtsummary) gtsummary object
columns	( <a href="#">tidy-select</a> ) Selector of columns in x\$table_body

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
# hide 95% CI, and replace with standard error
lm(age ~ marker + grade, trial) |>
  tbl_regression() |>
  modify_column_hide(conf.low) |>
  modify_column_unhide(columns = std.error)
```

---

modify\_column\_indent *Modify column indentation*

---

**Description**

Add, increase, or reduce indentation for columns.

**Usage**

```
modify_column_indent(x, columns, rows = NULL, indent = 4L, double_indent, undo)
```

**Arguments**

x	(gtsummary) gtsummary object
columns	( <a href="#">tidy-select</a> ) Selector of columns in x\$table_body
rows	(predicate expression) Predicate expression to select rows in x\$table_body. Can be used to style footnote, formatting functions, missing symbols, and text formatting. Default is NULL. See details below.
indent	(integer) An integer indicating how many space to indent text
double_indent, undo	<b>[Deprecated]</b>



**Value**

a gtsummary table

**See Also**

Other Advanced modifiers: [modify\\_column\\_merge\(\)](#), [modify\\_table\\_styling\(\)](#)

**Examples**

```
# remove indentation from `tbl_summary()`
trial |>
  tbl_summary(include = grade) |>
  modify_column_indent(columns = label, indent = 0L)

# increase indentation in `tbl_summary`
trial |>
  tbl_summary(include = grade) |>
  modify_column_indent(columns = label, rows = !row_type %in% 'label', indent = 8L)
```

---

modify\_column\_merge     *Modify Column Merging*

---

**Description**

Merge two or more columns in a gtsummary table. Use `show_header_names()` to print underlying column names.

**Usage**

```
modify_column_merge(x, pattern, rows = NULL)
```

**Arguments**

x	(gtsummary) gtsummary object
pattern	glue syntax string indicating how to merge columns in <code>x\$table_body</code> . For example, to construct a confidence interval use " <code>{conf.low}</code> , <code>{conf.high}</code> ".
rows	(predicate expression) Predicate expression to select rows in <code>x\$table_body</code> . Can be used to style footnote, formatting functions, missing symbols, and text formatting. Default is NULL. See details below.

**Value**

gtsummary table

## Details

1. Calling this function merely records the instructions to merge columns. The actual merging occurs when the gtsummary table is printed or converted with a function like `as_gt()`.
2. Because the column merging is delayed, it is recommended to perform major modifications to the table, such as those with `tbl_merge()` and `tbl_stack()`, before assigning merging instructions. Otherwise, unexpected formatting may occur in the final table.
3. If this functionality is used in conjunction with `tbl_stack()` (which includes `tbl_uvregression()`), there may be potential issues with printing. When columns are stack AND when the column-merging is defined with a quosure, you may run into issues due to the loss of the environment when 2 or more quosures are combined. If the expression version of the quosure is the same as the quosure (i.e. no evaluated objects), there should be no issues.

This function is used internally with care, and **it is *not* recommended for users.**

## Future Updates

There are planned updates to the implementation of this function with respect to the `pattern=` argument. Currently, this function replaces a numeric column with a formatted character column following `pattern=`. Once `gt::cols_merge()` gains the `rows=` argument the implementation will be updated to use it, which will keep numeric columns numeric. For the *vast majority* of users, *the planned change will be go unnoticed.*

## See Also

Other Advanced modifiers: [modify\\_column\\_indent\(\)](#), [modify\\_table\\_styling\(\)](#)

## Examples

```
# Example 1 -----
trial |>
  tbl_summary(by = trt, missing = "no", include = c(age, marker, trt)) |>
  add_p(all_continuous() ~ "t.test", pvalue_fun = label_style_pvalue(prepend_p = TRUE)) |>
  modify_fmt_fun(statistic ~ label_style_sigfig()) |>
  modify_column_merge(pattern = "t = {statistic}; {p.value}") |>
  modify_header(statistic = "**t-test**")

# Example 2 -----
lm(marker ~ age + grade, trial) |>
  tbl_regression() |>
  modify_column_merge(
    pattern = "{estimate} ({conf.low}, {conf.high})",
    rows = !is.na(estimate)
  )
```

---

modify_fmt_fun	<i>Modify formatting functions</i>
----------------	------------------------------------

---

## Description

Use this function to update the way numeric columns and rows of `.$table_body` are formatted

## Usage

```
modify_fmt_fun(x, ..., rows = NULL, update, quiet)
```

## Arguments

x	(gtsummary) A gtsummary object
...	<a href="#">dynamic-dots</a> Used to assign updates to formatting functions. Use <code>modify_fmt_fun(colname = &lt;fmt fn&gt;)</code> to update a single column. Using a formula will invoke <code>tidyselect</code> , e.g. <code>modify_fmt_fun(c(estimate, conf.low, conf.high) ~ &lt;fmt_fn&gt;)</code> Use the <code>show_header_names()</code> to see the column names that can be modified.
rows	(predicate expression) Predicate expression to select rows in <code>x\$table_body</code> . Can be used to style footnote, formatting functions, missing symbols, and text formatting. Default is <code>NULL</code> . See details below.
update, quiet	<b>[Deprecated]</b>

## rows argument

The `rows` argument accepts a predicate expression that is used to specify rows to apply formatting. The expression must evaluate to a logical when evaluated in `x$table_body`. For example, to apply formatting to the age rows pass `rows = variable == "age"`. A vector of row numbers is NOT acceptable.

A couple of things to note when using the `rows` argument.

1. You can use saved objects to create the predicate argument, e.g. `rows = variable == letters[1]`.
2. The saved object cannot share a name with a column in `x$table_body`. The reason for this is that in `tbl_merge()` the columns are renamed, and the renaming process cannot disambiguate the `variable` column from an external object named `variable` in the following expression `rows = .data$variable = .env$variable`.

## Examples

```
# Example 1 -----
# show 'grade' p-values to 3 decimal places and estimates to 4 sig figs
lm(age ~ marker + grade, trial) |>
  tbl_regression() %>%
```

```

modify_fmt_fun(
  p.value = label_style_pvalue(digits = 3),
  c(estimate, conf.low, conf.high) ~ label_style_sigfig(digits = 4),
  rows = variable == "grade"
)

```

---

modify\_table\_body      *Modify Table Body*

---

## Description

Function is for advanced manipulation of gtsummary tables. It allow users to modify the `.$table_body` data frame included in each gtsummary object.

If a new column is added to the table, default printing instructions will then be added to `.$table_styling`. By default, columns are hidden. To show a column, add a column header with `modify_header()` or call `modify_column_unhide()`.

## Usage

```
modify_table_body(x, fun, ...)
```

## Arguments

x	(gtsummary) A 'gtsummary' object
fun	(function) A function or formula. If a <i>function</i> , it is used as is. If a <i>formula</i> , e.g. <code>fun = ~.x  &gt; arrange(variable)</code> , it is converted to a function. The argument passed to fun is <code>x\$table_body</code> .
...	Additional arguments passed on to the function

## Value

A 'gtsummary' object

## Examples

```

# Example 1 -----
# Add number of cases and controls to regression table
trial |>
tbl_uvregression(
  y = response,
  include = c(age, marker),
  method = glm,
  method.args = list(family = binomial),
  exponentiate = TRUE,
  hide_n = TRUE
) |>

```

```

# adding number of non-events to table
modify_table_body(
  ~ .x %>%
    dplyr::mutate(N_nonevent = N_obs - N_event) |>
    dplyr::relocate(c(N_event, N_nonevent), .before = estimate)
) |>
# assigning header labels
modify_header(N_nonevent = "**Control N**", N_event = "**Case N**") |>
modify_fmt_fun(c(N_event, N_nonevent) ~ style_number)

```

---

modify\_table\_styling *Modify Table Styling*

---

### Description

This is a function meant for advanced users to gain more control over the characteristics of the resulting gtsummary table by directly modifying `.$table_styling`. *This function is primarily used in the development of other gtsummary functions, and very little checking of the passed arguments is performed.*

### Usage

```

modify_table_styling(
  x,
  columns,
  rows = NULL,
  label = NULL,
  spanning_header = NULL,
  hide = NULL,
  footnote = NULL,
  footnote_abbrev = NULL,
  align = NULL,
  missing_symbol = NULL,
  fmt_fun = NULL,
  text_format = NULL,
  undo_text_format = NULL,
  indent = NULL,
  text_interpret = c("md", "html"),
  cols_merge_pattern = NULL
)

```

### Arguments

x	(gtsummary) gtsummary object
columns	(tidy-select) Selector of columns in x\$table_body

rows	(predicate expression) Predicate expression to select rows in <code>x\$table_body</code> . Can be used to style footnote, formatting functions, missing symbols, and text formatting. Default is NULL. See details below.
label	(character) Character vector of column label(s). Must be the same length as columns.
spanning_header	(string) string with text for spanning header
hide	(scalar logical) Logical indicating whether to hide column from output
footnote	(string) string with text for footnote
footnote_abbrev	(string) string with abbreviation definition, e.g. "CI = Confidence Interval"
align	(string) String indicating alignment of column, must be one of <code>c("left", "right", "center")</code>
missing_symbol	(string) string indicating how missing values are formatted.
fmt_fun	(function) function that formats the statistics in the columns/rows in columns and rows
text_format, undo_text_format	(string) String indicated which type of text formatting to apply/remove to the rows and columns. Must be one of <code>c("bold", "italic")</code> .
indent	(integer) An integer indicating how many space to indent text
text_interpret	(string) Must be one of "md" or "html" and indicates the processing function as <code>gt::md()</code> or <code>gt::html()</code> . Use this in conjunction with arguments for header and footnotes.
cols_merge_pattern	(string) <b>[Experimental]</b> glue-syntax string indicating how to merge columns in <code>x\$table_body</code> . For example, to construct a confidence interval use <code>"{conf.low}, {conf.high}"</code> . The first column listed in the pattern string must match the single column name passed in <code>columns=</code> .

## Details

Review the [gtsummary definition](#) vignette for information on `.stable_styling` objects.

### rows argument

The `rows` argument accepts a predicate expression that is used to specify rows to apply formatting. The expression must evaluate to a logical when evaluated in `x$table_body`. For example, to apply

formatting to the age rows pass `rows = variable == "age"`. A vector of row numbers is NOT acceptable.

A couple of things to note when using the `rows` argument.

1. You can use saved objects to create the predicate argument, e.g. `rows = variable == letters[1]`.
2. The saved object cannot share a name with a column in `x$table_body`. The reason for this is that in `tbl_merge()` the columns are renamed, and the renaming process cannot disambiguate the `variable` column from an external object named `variable` in the following expression `rows = .data$variable = .env$variable`.

### **cols\_merge\_pattern argument**

There are planned updates to the implementation of column merging. Currently, this function replaces the numeric column with a formatted character column following `cols_merge_pattern=`. Once `gt::cols_merge()` gains the `rows=` argument the implementation will be updated to use it, which will keep numeric columns numeric. For the *vast majority* of users, *the planned change will be go unnoticed*.

If this functionality is used in conjunction with `tbl_stack()` (which includes `tbl_uvregression()`), there is potential issue with printing. When columns are stack AND when the column-merging is defined with a quosure, you may run into issues due to the loss of the environment when 2 or more quosures are combined. If the expression version of the quosure is the same as the quosure (i.e. no evaluated objects), there should be no issues. Regardless, this argument is used internally with care, and it is *not* recommended for users.

### **See Also**

See [gtsummary internals vignette](#)

Other Advanced modifiers: [modify\\_column\\_indent\(\)](#), [modify\\_column\\_merge\(\)](#)

---

plot

*Plot Regression Coefficients*

---

### **Description**

The `plot()` function extracts `x$table_body` and passes the it to `ggstats::ggcoef_plot()` along with formatting options.

### **Usage**

```
## S3 method for class 'tbl_regression'
plot(x, remove_header_rows = TRUE, remove_reference_rows = FALSE, ...)

## S3 method for class 'tbl_uvregression'
plot(x, remove_header_rows = TRUE, remove_reference_rows = FALSE, ...)
```

**Arguments**

`x` (tbl\_regression, tbl\_uvregression)  
 A 'tbl\_regression' or 'tbl\_uvregression' object

`remove_header_rows`  
 (scalar logical)  
 logical indicating whether to remove header rows for categorical variables. Default is TRUE

`remove_reference_rows`  
 (scalar logical)  
 logical indicating whether to remove reference rows for categorical variables. Default is FALSE.

`...` arguments passed to `ggstats::ggcoef_plot(...)`

**Details****[Experimental]****Value**

a ggplot

**Examples**

```
glm(response ~ marker + grade, trial, family = binomial) |>
  tbl_regression(
    add_estimate_to_reference_rows = TRUE,
    exponentiate = TRUE
  ) |>
  plot()
```

---

 proportion\_summary      *Summarize a proportion*


---

**Description**

**[Experimental]** This helper, to be used with `tbl_custom_summary()`, creates a function computing a proportion and its confidence interval.

**Usage**

```
proportion_summary(
  variable,
  value,
  weights = NULL,
  na.rm = TRUE,
  conf.level = 0.95,
```



```

method = c("wilson", "wilson.no.correct", "wald", "wald.no.correct", "exact",
"agresti.coull", "jeffreys")
)

```

### Arguments

variable	(string) String indicating the name of the variable from which the proportion will be computed.
value	(scalar) Value (or list of values) of variable to be taken into account in the numerator.
weights	(string) Optional string indicating the name of a frequency weighting variable. If NULL, all observations will be assumed to have a weight equal to 1.
na.rm	(scalar logical) Should missing values be removed before computing the proportion? (default is TRUE)
conf.level	(scalar numeric) Confidence level for the returned confidence interval. Must be strictly greater than 0 and less than 1. Default to 0.95, which corresponds to a 95 percent confidence interval.
method	(string) Confidence interval method. Must be one of <code>c("wilson", "wilson.no.correct", "wald", "wald.no.correct", "exact", "agresti.coull", "jeffreys")</code> . See <code>add_ci()</code> for details.

### Details

Computed statistics:

- {n} numerator, number of observations equal to values
- {N} denominator, number of observations
- {prop} proportion, i.e.  $n/N$
- {conf.low} lower confidence interval
- {conf.high} upper confidence interval

Methods `c("wilson", "wilson.no.correct")` are calculated with `stats::prop.test()` (with `correct = c(TRUE, FALSE)`). The default method, "wilson", includes the Yates continuity correction. Methods `c("exact", "asymptotic")` are calculated with `Hmisc::binconf()` and the corresponding method.

### Author(s)

Joseph Larmarange

## Examples

```
# Example 1 -----
Titanic |>
  as.data.frame() |>
  tbl_custom_summary(
    include = c("Age", "Class"),
    by = "Sex",
    stat_fns = ~ proportion_summary("Survived", "Yes", weights = "Freq"),
    statistic = ~ "{prop}% ({n}/{N}) [{conf.low}-{conf.high}]",
    digits = ~ list(
      prop = label_style_percent(digits = 1),
      n = 0,
      N = 0,
      conf.low = label_style_percent(),
      conf.high = label_style_percent()
    ),
    overall_row = TRUE,
    overall_row_last = TRUE
  ) |>
  bold_labels() |>
  modify_footnote(all_stat_cols() ~ "Proportion (%) of survivors (n/N) [95% CI]")
```

---

ratio\_summary

*Summarize the ratio of two variables*

---

## Description

**[Experimental]** This helper, to be used with `tbl_custom_summary()`, creates a function computing the ratio of two continuous variables and its confidence interval.

## Usage

```
ratio_summary(numerator, denominator, na.rm = TRUE, conf.level = 0.95)
```

## Arguments

numerator	(string) String indicating the name of the variable to be summed for computing the numerator.
denominator	(string) String indicating the name of the variable to be summed for computing the denominator.
na.rm	(scalar logical) Should missing values be removed before summing the numerator and the denominator? (default is TRUE)
conf.level	(scalar numeric) Confidence level for the returned confidence interval. Must be strictly greater than 0 and less than 1. Default to 0.95, which corresponds to a 95 percent confidence interval.

**Details**

Computed statistics:

- {num} sum of the variable defined by numerator
- {denom} sum of the variable defined by denominator
- {ratio} ratio of num by denom
- {conf.low} lower confidence interval
- {conf.high} upper confidence interval

Confidence interval is computed with `stats::poisson.test()`, if and only if num is an integer.

**Author(s)**

Joseph Larmarange

**Examples**

```
# Example 1 -----
trial |>
  tbl_custom_summary(
    include = c("stage", "grade"),
    by = "trt",
    stat_fns = ~ ratio_summary("response", "ttdeath"),
    statistic = ~ "{ratio} [{conf.low}; {conf.high}] ({num}/{denom})",
    digits = ~ c(ratio = 3, conf.low = 2, conf.high = 2),
    overall_row = TRUE,
    overall_row_label = "All stages & grades"
  ) |>
  bold_labels() |>
  modify_footnote(all_stat_cols() ~ "Ratio [95% CI] (n/N)")
```

---

remove_row_type	<i>Remove rows</i>
-----------------	--------------------

---

**Description**

Removes either the header, reference, or missing rows from a gtsummary table.

**Usage**

```
remove_row_type(
  x,
  variables = everything(),
  type = c("header", "reference", "missing", "level", "all"),
  level_value = NULL
)
```

**Arguments**

x	(gtsummary) A gtsummary object
variables	(tidy-select) Variables to to remove rows from. Default is everything()
type	(string) Type of row to remove. Must be one of c("header", "reference", "missing", "level", "all")
level_value	(string) When type='level' you can specify the <i>character</i> value of the level to remove. When NULL all levels are removed.

**Value**

Modified gtsummary table

**Examples**

```
# Example 1 -----
trial |>
  dplyr::mutate(
    age60 = ifelse(age < 60, "<60", "60+")
  ) |>
tbl_summary(by = trt, missing = "no", include = c(trt, age, age60)) |>
  remove_row_type(age60, type = "header")
```

---

select\_helpers

*Select helper functions*

---

**Description**

Set of functions to supplement the {tidyselect} set of functions for selecting columns of data frames (and other items as well).

- all\_continuous() selects continuous variables
- all\_continuous2() selects only type "continuous2"
- all\_categorical() selects categorical (including "dichotomous") variables
- all\_dichotomous() selects only type "dichotomous"
- all\_tests() selects variables by the name of the test performed
- all\_stat\_cols() selects columns from tbl\_summary/tbl\_svsummary object with summary statistics (i.e. "stat\_0", "stat\_1", "stat\_2", etc.)
- all\_interaction() selects interaction terms from a regression model
- all\_intercepts() selects intercept terms from a regression model
- all\_contrasts() selects variables in regression model based on their type of contrast

**Usage**

```
all_continuous(continuous2 = TRUE)

all_continuous2()

all_categorical(dichotomous = TRUE)

all_dichotomous()

all_tests(tests)

all_intercepts()

all_interaction()

all_contrasts(
  contrasts_type = c("treatment", "sum", "poly", "helmert", "sdif", "other")
)

all_stat_cols(stat_0 = TRUE)
```

**Arguments**

continuous2	(scalar logical) Logical indicating whether to include continuous2 variables. Default is TRUE
dichotomous	(scalar logical) Logical indicating whether to include dichotomous variables. Default is TRUE
tests	(character) character vector indicating the test type of the variables to select, e.g. select all variables being compared with "t.test".
contrasts_type	(character) type of contrast to select. Select among contrast types c("treatment", "sum", "poly", "helmert", "sdif", "other"). Default is all contrast types.
stat_0	(scalar logical) When FALSE, will not select the "stat_0" column. Default is TRUE

**Value**

A character vector of column names selected

**See Also**

Review [list](#), [formula](#), and [selector syntax](#) used throughout gtsummary

**Examples**

```
select_ex1 <-
  trial |>
```

```
select(age, response, grade) |>
tbl_summary(
  statistic = all_continuous() ~ "{mean} ({sd})",
  type = all_dichotomous() ~ "categorical"
)
```

---

separate\_p\_footnotes *Create footnotes for individual p-values*

---

## Description

### [Questioning]

The usual presentation of footnotes for p-values on a gtsummary table is to have a single footnote that lists all statistical tests that were used to compute p-values on a given table. The `separate_p_footnotes()` function separates aggregated p-value footnotes to individual footnotes that denote the specific test used for each of the p-values.

## Usage

```
separate_p_footnotes(x)
```

## Arguments

x (tbl\_summary, tbl\_svsummary)  
Object with class "tbl\_summary" or "tbl\_svsummary"

## Examples

```
# Example 1 -----
trial |>
tbl_summary(by = trt, include = c(age, grade)) |>
add_p() |>
separate_p_footnotes()
```

---

set\_gtsummary\_theme *Set gtsummary theme*

---

## Description

Functions to **set**, **reset**, **get**, and evaluate **with** gtsummary themes.

- `set_gtsummary_theme()` set a theme
- `reset_gtsummary_theme()` reset themes
- `get_gtsummary_theme()` get a named list with all active theme elements
- `with_gtsummary_theme()` evaluate an expression with a theme temporarily set
- `check_gtsummary_theme()` checks if passed theme is valid

**Usage**

```

set_gtsummary_theme(x, quiet)

reset_gtsummary_theme()

get_gtsummary_theme()

with_gtsummary_theme(
  x,
  expr,
  env = rlang::caller_env(),
  msg_ignored_elements = NULL
)

check_gtsummary_theme(x)

```

**Arguments**

x	(named list) A named list defining a gtsummary theme.
quiet	<b>[Deprecated]</b>
expr	(expression) Expression to be evaluated with the theme specified in x= loaded
env	(environment) The environment in which to evaluate expr=
msg_ignored_elements	(string) Default is NULL with no message printed. Pass a string that will be printed with cli::cli_alert_info(). The "{elements}" object contains vector of theme elements that will be overwritten and ignored.

**Details**

The default formatting and styling throughout the gtsummary package are taken from the published reporting guidelines of the top four urology journals: European Urology, The Journal of Urology, Urology and the British Journal of Urology International. Use this function to change the default reporting style to match another journal, or your own personal style.

**See Also**

[Themes vignette](#)

Available [gtsummary themes](#)

**Examples**

```

# Setting JAMA theme for gtsummary
set_gtsummary_theme(theme_gtsummary_journal("jama"))
# Themes can be combined by including more than one

```

```

set_gtsummary_theme(theme_gtsummary_compact())

set_gtsummary_theme_ex1 <-
  trial |>
  tbl_summary(by = trt, include = c(age, grade, trt)) |>
  add_stat_label() |>
  as_gt()

# reset gtsummary theme
reset_gtsummary_theme()

```

---

 sort\_filter\_p

*Sort/filter by p-values*


---

## Description

Sort/filter by p-values

## Usage

```

sort_p(x, q = FALSE)

filter_p(x, q = FALSE, t = 0.05)

```

## Arguments

x	(gtsummary) An object created using gtsummary functions
q	(scalar logical) When TRUE will check the q-value column rather than the p-value. Default is FALSE.
t	(scalar numeric) Threshold below which values will be retained. Default is 0.05.

## Author(s)

Karissa Whiting, Daniel D. Sjoberg

## Examples

```

# Example 1 -----
trial %>%
  select(age, grade, response, trt) %>%
  tbl_summary(by = trt) %>%
  add_p() %>%
  filter_p(t = 0.8) %>%
  sort_p()

```



```
# Example 2 -----
glm(response ~ trt + grade, trial, family = binomial(link = "logit")) %>%
  tbl_regression(exponentiate = TRUE) %>%
  sort_p()
```

---

style\_number

*Style numbers*


---

## Description

Style numbers

## Usage

```
style_number(
  x,
  digits = 0,
  big.mark = ifelse(decimal.mark == ",", " ", ", "),
  decimal.mark = getOption("OutDec"),
  scale = 1,
  prefix = "",
  suffix = "",
  ...
)
```

## Arguments

x	(numeric) Numeric vector
digits	(non-negative integer) Integer or vector of integers specifying the number of decimals to round x. When vector is passed, each integer is mapped 1:1 to the numeric values in x
big.mark	(string) Character used between every 3 digits to separate hundreds/thousands/millions/etc. Default is ",", except when decimal.mark = " " when the default is a space.
decimal.mark	(string) The character to be used to indicate the numeric decimal point. Default is "." or getOption("OutDec")
scale	(scalar numeric) A scaling factor: x will be multiplied by scale before formatting.
prefix	(string) Additional text to display before the number.
suffix	(string) Additional text to display after the number.
...	Arguments passed on to base::format()

**Value**

formatted character vector

**Examples**

```
c(0.111, 12.3) |> style_number(digits = 1)
c(0.111, 12.3) |> style_number(digits = c(1, 0))
```

---

style_percent	<i>Style percentages</i>
---------------	--------------------------

---

**Description**

Style percentages

**Usage**

```
style_percent(
  x,
  digits = 0,
  big.mark = ifelse(decimal.mark == ",", " ", " "),
  decimal.mark = getOption("OutDec"),
  prefix = "",
  suffix = "",
  symbol,
  ...
)
```

**Arguments**

x	numeric vector of percentages
digits	number of digits to round large percentages (i.e. greater than 10%). Smaller percentages are rounded to digits + 1 places. Default is 0
big.mark	(string) Character used between every 3 digits to separate hundreds/thousands/millions/etc. Default is ",", except when decimal.mark = "." when the default is a space.
decimal.mark	(string) The character to be used to indicate the numeric decimal point. Default is "." or getOption("OutDec")
prefix	(string) Additional text to display before the number.
suffix	(string) Additional text to display after the number.
symbol	Logical indicator to include percent symbol in output. Default is FALSE.
...	Arguments passed on to base::format()

**Value**

A character vector of styled percentages

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
percent_vals <- c(-1, 0, 0.0001, 0.005, 0.01, 0.10, 0.45356, 0.99, 1.45)
style_percent(percent_vals)
style_percent(percent_vals, suffix = "%", digits = 1)
```

---

style_pvalue	<i>Style p-values</i>
--------------	-----------------------

---

**Description**

Style p-values

**Usage**

```
style_pvalue(
  x,
  digits = 1,
  prepend_p = FALSE,
  big.mark = ifelse(decimal.mark == ",", " ", ", "),
  decimal.mark = getOption("OutDec"),
  ...
)
```

**Arguments**

x	(numeric) Numeric vector of p-values.
digits	(integer) Number of digits large p-values are rounded. Must be 1, 2, or 3. Default is 1.
prepend_p	(scalar logical) Logical. Should 'p=' be prepended to formatted p-value. Default is FALSE
big.mark	(string) Character used between every 3 digits to separate hundreds/thousands/millions/etc. Default is " ", except when decimal.mark = ",", when the default is a space.
decimal.mark	(string) The character to be used to indicate the numeric decimal point. Default is "." or getOption("OutDec")
...	Arguments passed on to base::format()

**Value**

A character vector of styled p-values

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
pvals <- c(
  1.5, 1, 0.999, 0.5, 0.25, 0.2, 0.197, 0.12, 0.10, 0.0999, 0.06,
  0.03, 0.002, 0.001, 0.00099, 0.0002, 0.00002, -1
)
style_pvalue(pvals)
style_pvalue(pvals, digits = 2, prepend_p = TRUE)
```

---

style\_ratio

*Style ratios*

---

**Description**

When reporting ratios, such as relative risk or an odds ratio, we'll often want the rounding to be similar on each side of the number 1. For example, if we report an odds ratio of 0.95 with a confidence interval of 0.70 to 1.24, we would want to round to two decimal places for all values. In other words, 2 significant figures for numbers less than 1 and 3 significant figures 1 and larger. `style_ratio()` performs significant figure-like rounding in this manner.

**Usage**

```
style_ratio(
  x,
  digits = 2,
  big.mark = ifelse(decimal.mark == ",", " ", " "),
  decimal.mark = getOption("OutDec"),
  prefix = "",
  suffix = "",
  ...
)
```

**Arguments**

<code>x</code>	(numeric) Numeric vector
<code>digits</code>	(integer) Integer specifying the number of significant digits to display for numbers below 1. Numbers larger than 1 will be be <code>digits + 1</code> . Default is <code>digits = 2</code> .
<code>big.mark</code>	(string) Character used between every 3 digits to separate hundreds/thousands/millions/etc. Default is <code>" "</code> , except when <code>decimal.mark = ","</code> when the default is a space.

decimal.mark	(string) The character to be used to indicate the numeric decimal point. Default is "." or getOption("OutDec")
prefix	(string) Additional text to display before the number.
suffix	(string) Additional text to display after the number.
...	Arguments passed on to base::format()

**Value**

A character vector of styled ratios

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
c(0.123, 0.9, 1.1234, 12.345, 101.234, -0.123, -0.9, -1.1234, -12.345, -101.234) |>
  style_ratio()
```

---

style\_sigfig

*Style significant figure-like rounding*

---

**Description**

Converts a numeric argument into a string that has been rounded to a significant figure-like number. Scientific notation output is avoided, however, and additional significant figures may be displayed for large numbers. For example, if the number of significant digits requested is 2, 123 will be displayed (rather than 120 or  $1.2 \times 10^2$ ).

**Usage**

```
style_sigfig(
  x,
  digits = 2,
  scale = 1,
  big.mark = ifelse(decimal.mark == ",", " ", " ", " "),
  decimal.mark = getOption("OutDec"),
  prefix = "",
  suffix = "",
  ...
)
```

**Arguments**

<code>x</code>	Numeric vector
<code>digits</code>	Integer specifying the minimum number of significant digits to display
<code>scale</code>	(scalar numeric) A scaling factor: <code>x</code> will be multiplied by <code>scale</code> before formatting.
<code>big.mark</code>	(string) Character used between every 3 digits to separate hundreds/thousands/millions/etc. Default is <code>,</code> , except when <code>decimal.mark = "</code> , <code>"</code> when the default is a space.
<code>decimal.mark</code>	(string) The character to be used to indicate the numeric decimal point. Default is <code>.</code> or <code>getOption("OutDec")</code>
<code>prefix</code>	(string) Additional text to display before the number.
<code>suffix</code>	(string) Additional text to display after the number.
<code>...</code>	Arguments passed on to <code>base::format()</code>

**Value**

A character vector of styled numbers

**Details**

- Scientific notation output is avoided.
- If 2 significant figures are requested, the number is rounded to no more than 2 decimal places. For example, a number will be rounded to 2 decimals places when  $\text{abs}(x) < 1$ , 1 decimal place when  $\text{abs}(x) \geq 1$  &  $\text{abs}(x) < 10$ , and to the nearest integer when  $\text{abs}(x) \geq 10$ .
- Additional significant figures may be displayed for large numbers. For example, if the number of significant digits requested is 2, 123 will be displayed (rather than 120 or  $1.2 \times 10^2$ ).

**Author(s)**

Daniel D. Sjoberg

**See Also**

Other style tools: [label\\_style](#)

**Examples**

```
c(0.123, 0.9, 1.1234, 12.345, -0.123, -0.9, -1.1234, -132.345, NA, -0.001) %>%
  style_sigfig()
```

---

tbl\_ard\_continuous      *Summarize continuous variable*


---

**Description****[Experimental]**

Summarize a continuous variable by one or more categorical variables

**Usage**

```
tbl_ard_continuous(
  cards,
  variable,
  include,
  by = NULL,
  label = NULL,
  statistic = everything() ~ "{median} ({p25}, {p75})",
  value = NULL
)
```

**Arguments**

cards	(card) An ARD object of class "card" typically created with <code>cards::ard_*</code> () functions.
variable	(string) A single variable name of the continuous variable being summarized.
include	(character) Character vector of the categorical variables to
by	(string) A single variable name of the stratifying variable.
label	( <a href="#">formula-list-selector</a> ) Used to override default labels in summary table, e.g. <code>list(age = "Age, years")</code> . The default for each variable is the column label attribute, <code>attr(., 'label')</code> . If no label has been set, the column name is used.
statistic	( <a href="#">formula-list-selector</a> ) Specifies summary statistics to display for each variable. The default is <code>everything() ~ "{median} ({p25}, {p75})"</code> .
value	( <a href="#">formula-list-selector</a> ) Supply a value to display a variable on a single row, printing the results for the variable associated with the value (similar to a 'dichotomous' display in <code>tbl_summary()</code> ).

**Value**

a gsummary table of class "tbl\_ard\_summary"

**Examples**

```

library(cards)

# Example 1 -----
# the primary ARD with the results
ard_continuous(
  # the order variables are passed is important for the `by` variable.
  # 'trt' is the column stratifying variable and needs to be listed first.
  trial, by = c(trt, grade), variables = age
) |>
# adding OPTIONAL information about the summary variables
bind_ard(
  # add univariate trt tabulation
  ard_categorical(trial, variables = trt),
  # add missing and attributes ARD
  ard_missing(trial, by = c(trt, grade), variables = age),
  ard_attributes(trial, variables = c(trt, grade, age))
) |>
tbl_ard_continuous(by = "trt", variable = "age", include = "grade")

# Example 2 -----
# the primary ARD with the results
ard_continuous(trial, by = grade, variables = age) |>
# adding OPTIONAL information about the summary variables
bind_ard(
  # add missing and attributes ARD
  ard_missing(trial, by = grade, variables = age),
  ard_attributes(trial, variables = c(grade, age))
) |>
tbl_ard_continuous(variable = "age", include = "grade")

```

---

tbl\_ard\_hierarchical *ARD Hierarchical Table*


---

**Description****[Experimental]**

*This is an preview of this function. There will be changes in the coming releases, and changes will not undergo a formal deprecation cycle.*

Constructs tables from nested or hierarchical data structures (e.g. adverse events).

**Usage**

```
tbl_ard_hierarchical(
  cards,
  variables,
  by = NULL,
  include = everything(),
  statistic = "{n} ({p})",

```



```
  label = NULL
)
```

### Arguments

cards	(card) An ARD object of class "card" typically created with <code>cards::ard_*</code> () functions.
variables	( <a href="#">tidy-select</a> ) character vector or tidy-selector of columns in data used to create a hierarchy. Hierarchy will be built with variables in the order given.
by	( <a href="#">tidy-select</a> ) a single column from data. Summary statistics will be stratified by this variable. Default is NULL.
include	( <a href="#">tidy-select</a> ) variables from hierarchy for which summary statistics should be returned (on the variable label rows) Including the last element of hierarchy has no effect since each level has its own row for this variable. The default is <code>everything()</code> .
statistic	(string) used to specify the summary statistics to display for all variables in <code>tbl_hierarchical()</code> . The default is "{n} ({p})".
label	( <a href="#">formula-list-selector</a> ) used to override default labels in hierarchical table, e.g. <code>list(AESOC = "System Organ Class")</code> . The default for each variable is the column label attribute, <code>attr(, 'label')</code> . If no label has been set, the column name is used.

### Value

a gtssummary table of class "tbl\_ard\_hierarchical"

### Examples

```
ADAE_subset <- cards::ADAE |>
  dplyr::filter(
    AESOC %in% unique(cards::ADAE$AESOC)[1:5],
    AETERM %in% unique(cards::ADAE$AETERM)[1:5]
  )

# Example 1: Event Rates -----
# First, build the ARD
ard <-
  cards::ard_stack_hierarchical(
    data = ADAE_subset,
    variables = c(AESOC, AETERM),
    by = TRTA,
    denominator = cards::ADSL |> mutate(TRTA = ARM),
    id = USBJID
  )
```

```

# Second, build table from the ARD
tbl_ard_hierarchical(
  cards = ard,
  variables = c(AESOC, AETERM),
  by = TRTA
)

# Example 2: Event Counts -----
ard <-
  cards::ard_stack_hierarchical_count(
    data = ADAE_subset,
    variables = c(AESOC, AETERM),
    by = TRTA,
    denominator = cards::ADSL |> mutate(TRTA = ARM)
  )

tbl_ard_hierarchical(
  cards = ard,
  variables = c(AESOC, AETERM),
  by = TRTA,
  statistic = "{n}"
)

```

---

tbl_ard_summary	<i>ARD summary table</i>
-----------------	--------------------------

---

## Description

### [Experimental]

The `tbl_ard_summary()` function tables descriptive statistics for continuous, categorical, and dichotomous variables. The functions accepts an ARD object.

## Usage

```

tbl_ard_summary(
  cards,
  by = NULL,
  statistic = list(all_continuous() ~ "{median} ({p25}, {p75})", all_categorical() ~
    "{n} ({p}%)"),
  type = NULL,
  label = NULL,
  missing = c("no", "ifany", "always"),
  missing_text = "Unknown",
  missing_stat = "{N_miss}",
  include = everything(),
  overall = FALSE
)

```

**Arguments**

cards	(card) An ARD object of class "card" typically created with <code>cards::ard_*()</code> functions.
by	(tidy-select) A single column from data. Summary statistics will be stratified by this variable. Default is NULL
statistic	(formula-list-selector) Used to specify the summary statistics for each variable. Each of the statistics must be present in card as no new statistics are calculated in this function. The default is <code>list(all_continuous() ~ "{median} ({p25}, {p75})", all_categorical() ~ "{n} ({p}%)")</code> .
type	(formula-list-selector) Specifies the summary type. Accepted values are <code>c("continuous", "continuous2", "categorical", "dichotomous")</code> . Continuous summaries may be assigned <code>c("continuous", "continuous2")</code> , while categorical and dichotomous cannot be modified.
label	(formula-list-selector) Used to override default labels in summary table, e.g. <code>list(age = "Age, years")</code> . The default for each variable is the column label attribute, <code>attr(, 'label')</code> . If no label has been set, the column name is used.
missing, missing_text, missing_stat	Arguments dictating how and if missing values are presented: <ul style="list-style-type: none"> <li>• <code>missing</code>: must be one of <code>c("no", "ifany", "always")</code></li> <li>• <code>missing_text</code>: string indicating text shown on missing row. Default is "Unknown"</li> <li>• <code>missing_stat</code>: statistic to show on missing row. Default is "{N_miss}". Possible values are <code>N_miss, N_obs, N_nonmiss, p_miss, p_nonmiss</code></li> </ul>
include	(tidy-select) Variables to include in the summary table. Default is <code>everything()</code>
overall	(scalar logical) When TRUE, the cards input is parsed into two parts to run <code>tbl_ard_summary(cards_by)  &gt; add_overall(cards_overall)</code> . Can only be used when <code>by</code> argument is specified. Default is FALSE.

**Value**

a gtsummary table of class "tbl\_ard\_summary"

**Examples**

```
library(cards)

ard_stack(
  data = ADSL,
  ard_categorical(variables = "AGEGR1"),
```

```

ard_continuous(variables = "AGE"),
. attributes = TRUE,
. missing = TRUE,
. total_n = TRUE
) |>
tbl_ard_summary()

ard_stack(
  data = ADSL,
  . by = ARM,
  ard_categorical(variables = "AGEGR1"),
  ard_continuous(variables = "AGE"),
  . attributes = TRUE,
  . missing = TRUE,
  . total_n = TRUE
) |>
tbl_ard_summary(by = ARM)

ard_stack(
  data = ADSL,
  . by = ARM,
  ard_categorical(variables = "AGEGR1"),
  ard_continuous(variables = "AGE"),
  . attributes = TRUE,
  . missing = TRUE,
  . total_n = TRUE,
  . overall = TRUE
) |>
tbl_ard_summary(by = ARM, overall = TRUE)

```

---

tbl\_ard\_wide\_summary *Wide ARD summary table*

---

## Description

### [Experimental]

This function is similar to `tbl_ard_summary()`, but places summary statistics wide, in separate columns. All included variables must be of the same summary type, e.g. all continuous summaries or all categorical summaries (which encompasses dichotomous variables).

## Usage

```

tbl_ard_wide_summary(
  cards,
  statistic = switch(type[[1]], continuous = c("{median}", "{p25}", "{p75}"), c("{n}",
    "{p}%")),
  type = NULL,
  label = NULL,
  value = NULL,
  include = everything()
)

```

**Arguments**

cards	(card) An ARD object of class "card" typically created with <code>cards::ard_*</code> () functions.
statistic	(character) character vector of the statistics to present. Each element of the vector will result in a column in the summary table. Default is <code>c("{median}", "{p25}", "{p75}")</code> for continuous summaries, and <code>c("{n}", "{p}%")</code> for categorical/dichotomous summaries
type	( <a href="#">formula-list-selector</a> ) Specifies the summary type. Accepted values are <code>c("continuous", "continuous2", "categorical", "dichotomous")</code> . If not specified, default type is assigned via <code>assign_summary_type()</code> . See below for details.
label	( <a href="#">formula-list-selector</a> ) Used to override default labels in summary table, e.g. <code>list(age = "Age, years")</code> . The default for each variable is the column label attribute, <code>attr(., 'label')</code> . If no label has been set, the column name is used.
value	( <a href="#">formula-list-selector</a> ) Specifies the level of a variable to display on a single row. The <code>gtsummary</code> type selectors, e.g. <code>all_dichotomous()</code> , cannot be used with this argument. Default is <code>NULL</code> . See below for details.
include	( <a href="#">tidy-select</a> ) Variables to include in the summary table. Default is <code>everything()</code> .

**Value**

a `gtsummary` table of class `'tbl_wide_summary'`

**Examples**

```
library(cards)

ard_stack(
  trial,
  ard_continuous(variables = age),
  .missing = TRUE,
  .attributes = TRUE,
  .total_n = TRUE
) |>
tbl_ard_wide_summary()

ard_stack(
  trial,
  ard_dichotomous(variables = response),
  ard_categorical(variables = grade),
  .missing = TRUE,
  .attributes = TRUE,
  .total_n = TRUE
)
```

```
) |>
tbl_ard_wide_summary()
```

---

tbl\_butcher

*Butcher table*


---

### Description

Some gtsummary objects can become large and the size becomes cumbersome when working with the object. The function removes all elements from a gtsummary object, except those required to print the table. This may result in gtsummary functions that add information or modify the table, such as `add_global_p()`, will no longer execute after the excess elements have been removed (aka butchered). Of note, the majority of `inline_text()` calls will continue to execute properly.

### Usage

```
tbl_butcher(x, include = c("table_body", "table_styling"))
```

### Arguments

x	(gtsummary) a gtsummary object
include	(character) names of additional elements to retain in the gtsummary object. <code>c("table_body", "table_styling")</code> will always be retained.

### Value

a gtsummary object

### Examples

```
tbl_large <-
  trial |>
  tbl_uvregression(
    y = age,
    method = lm
  )

tbl_butchered <-
  tbl_large |>
  tbl_butcher()

# size comparison
object.size(tbl_large) |> format(units = "Mb")
object.size(tbl_butchered) |> format(units = "Mb")
```

---

tbl\_continuous      *Summarize continuous variable*


---

## Description

Summarize a continuous variable by one or more categorical variables

## Usage

```
tbl_continuous(
  data,
  variable,
  include = everything(),
  digits = NULL,
  by = NULL,
  statistic = everything() ~ "{median} ({p25}, {p75})",
  label = NULL,
  value = NULL
)
```

## Arguments

data	(data.frame) A data frame.
variable	( <a href="#">tidy-select</a> ) A single column from data. Variable name of the continuous column to be summarized.
include	( <a href="#">tidy-select</a> ) Variables to include in the summary table. Default is everything().
digits	( <a href="#">formula-list-selector</a> ) Specifies how summary statistics are rounded. Values may be either integer(s) or function(s). If not specified, default formatting is assigned via <code>assign_summary_digits()</code> . See below for details.
by	( <a href="#">tidy-select</a> ) A single column from data. Summary statistics will be stratified by this variable. Default is NULL.
statistic	( <a href="#">formula-list-selector</a> ) Specifies summary statistics to display for each variable. The default is <code>everything() ~ "{median} ({p25}, {p75})"</code> .
label	( <a href="#">formula-list-selector</a> ) Used to override default labels in summary table, e.g. <code>list(age = "Age, years")</code> . The default for each variable is the column label attribute, <code>attr(., 'label')</code> . If no label has been set, the column name is used.

value (formula-list-selector)  
 Supply a value to display a variable on a single row, printing the results for the variable associated with the value (similar to a 'dichotomous' display in `tbl_summary()`).

### Value

a gtsummary table

### Examples

```
# Example 1 -----
tbl_continuous(
  data = trial,
  variable = age,
  by = trt,
  include = grade
)

# Example 2 -----
trial |>
  dplyr::mutate(all_subjects = 1) |>
  tbl_continuous(
    variable = age,
    statistic = ~"{mean} ({sd})",
    by = trt,
    include = c(all_subjects, stage, grade),
    value = all_subjects ~ 1,
    label = list(all_subjects = "All Subjects")
  )
```

---

tbl\_cross

*Cross table*

---

### Description

The function creates a cross table of categorical variables.

### Usage

```
tbl_cross(
  data,
  row = 1L,
  col = 2L,
  label = NULL,
  statistic = ifelse(percent == "none", "{n}", "{n} ({p}%"),
  digits = NULL,
  percent = c("none", "column", "row", "cell"),
  margin = c("column", "row"),
```



```

missing = c("ifany", "always", "no"),
missing_text = "Unknown",
margin_text = "Total"
)

```

## Arguments

data	(data.frame) A data frame.
row	(tidy-select) Column name in data to be used for the rows of cross table. Default is the first column in data.
col	(tidy-select) Column name in data to be used for the columns of cross table. Default is the second column in data.
label	(formula-list-selector) Used to override default labels in summary table, e.g. list(age = "Age, years"). The default for each variable is the column label attribute, attr(., 'label'). If no label has been set, the column name is used.
statistic	(string) A string with the statistic name in curly brackets to be replaced with the numeric statistic (see glue::glue). The default is {n}. If percent argument is "column", "row", or "cell", default is "{n} ({p}%)".
digits	(numeric/list/function) Specifies the number of decimal places to round the summary statistics. This argument is passed to tbl_summary(digits = ~digits). By default integers are shown to the zero decimal places, and percentages are formatted with style_percent(). If you would like to modify either of these, pass a vector of integers indicating the number of decimal places to round the statistics. For example, if the statistic being calculated is "{n} ({p}%)", and you want the percent rounded to 2 decimal places use digits = c(0, 2). User may also pass a styling function: digits = style_sigfig
percent	(string) Indicates the type of percentage to return. Must be one of "none", "column", "row", or "cell". Default is "cell" when {N} or {p} is used in statistic.
margin	(character) Indicates which margins to add to the table. Default is c("row", "column"). Use margin = NULL to suppress both row and column margins.
missing	(string) Must be one of c("ifany", "no", "always").
missing_text	(string) String indicating text shown on missing row. Default is "Unknown"
margin_text	(string) Text to display for margin totals. Default is "Total"

**Value**

A `tbl_cross` object

**Author(s)**

Karissa Whiting, Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
trial |>
  tbl_cross(row = trt, col = response) |>
  bold_labels()

# Example 2 -----
trial |>
  tbl_cross(row = stage, col = trt, percent = "cell") |>
  add_p() |>
  bold_labels()
```

---

`tbl_custom_summary`      *Create a table of summary statistics using a custom summary function*

---

**Description****[Experimental]**

The `tbl_custom_summary()` function calculates descriptive statistics for continuous, categorical, and dichotomous variables. This function is similar to `tbl_summary()` but allows you to provide a custom function in charge of computing the statistics (see Details).

**Usage**

```
tbl_custom_summary(
  data,
  by = NULL,
  label = NULL,
  stat_fns,
  statistic,
  digits = NULL,
  type = NULL,
  value = NULL,
  missing = c("ifany", "no", "always"),
  missing_text = "Unknown",
  missing_stat = "{N_miss}",
  include = everything(),
  overall_row = FALSE,
  overall_row_last = FALSE,
  overall_row_label = "Overall"
)
```

**Arguments**

data	(data.frame) A data frame.
by	(tidy-select) A single column from data. Summary statistics will be stratified by this variable. Default is NULL.
label	(formula-list-selector) Used to override default labels in summary table, e.g. list(age = "Age, years"). The default for each variable is the column label attribute, attr(., 'label'). If no label has been set, the column name is used.
stat_fns	(formula-list-selector) Specifies the function to be used to compute the statistics (see below for details and examples). You can also use dedicated helpers such as <a href="#">ratio_summary()</a> or <a href="#">proportion_summary()</a> .
statistic	(formula-list-selector) Specifies summary statistics to display for each variable. The default is list(all_continuous() ~ "{median} ({p25}, {p75})", all_categorical() ~ "{n} ({p}%)"). See below for details.
digits	(formula-list-selector) Specifies how summary statistics are rounded. Values may be either integer(s) or function(s). If not specified, default formatting is assigned via <a href="#">assign_summary_digits()</a> . See below for details.
type	(formula-list-selector) Specifies the summary type. Accepted values are c("continuous", "continuous2", "categorical", "dichotomous"). If not specified, default type is assigned via <a href="#">assign_summary_type()</a> . See below for details.
value	(formula-list-selector) Specifies the level of a variable to display on a single row. The gtsummary type selectors, e.g. <a href="#">all_dichotomous()</a> , cannot be used with this argument. Default is NULL. See below for details.
missing, missing_text, missing_stat	Arguments dictating how and if missing values are presented: <ul style="list-style-type: none"> <li>• missing: must be one of c("ifany", "no", "always")</li> <li>• missing_text: string indicating text shown on missing row. Default is "Unknown"</li> <li>• missing_stat: statistic to show on missing row. Default is "{N_miss}". Possible values are N_miss, N_obs, N_nonmiss, p_miss, p_nonmiss.</li> </ul>
include	(tidy-select) Variables to include in the summary table. Default is everything().
overall_row	(scalar logical) Logical indicator to display an overall row. Default is FALSE. Use <a href="#">add_overall()</a> to add an overall column.
overall_row_last	(scalar logical) Logical indicator to display overall row last in table. Default is FALSE, which will display overall row first.

overall\_row\_label  
 (string)  
 String indicating the overall row label. Default is "Overall".

### Value

A `tbl_custom_summary` object

### Similarities with `tbl_summary()`

Please refer to the help file of `tbl_summary()` regarding the use of select helpers, and arguments include, by, type, value, digits, missing and missing\_text.

### stat\_fns argument

The `stat_fns` argument specify the custom function(s) to be used for computing the summary statistics. For example, `stat_fns = everything() ~ foo`.

Each function may take the following arguments: `foo(data, full_data, variable, by, type, ...)`

- `data=` is the input data frame passed to `tbl_custom_summary()`, subset according to the level of `by` or `variable` if any, excluding NA values of the current variable
- `full_data=` is the full input data frame passed to `tbl_custom_summary()`
- `variable=` is a string indicating the variable to perform the calculation on
- `by=` is a string indicating the by variable from `tbl_custom_summary=`, if present
- `type=` is a string indicating the type of variable (continuous, categorical, ...)
- `stat_display=` a string indicating the statistic to display (for the `statistic` argument, for that variable)

The user-defined does not need to utilize each of these inputs. It's encouraged the user-defined function accept `...` as each of the arguments *will* be passed to the function, even if not all inputs are utilized by the user's function, e.g. `foo(data, ...)` (see examples).

The user-defined function should return a one row `dplyr::tibble()` with one column per summary statistics (see examples).

### statistic argument

The `statistic` argument specifies the statistics presented in the table. The input is a list of formulas that specify the statistics to report. For example, `statistic = list(age ~ "{mean} ({sd})")`. A statistic name that appears between curly brackets will be replaced with the numeric statistic (see `glue::glue()`). All the statistics indicated in the `statistic` argument should be returned by the functions defined in the `stat_fns` argument.

When the summary type is "continuous2", pass a vector of statistics. Each element of the vector will result in a separate row in the summary table.

For both categorical and continuous variables, statistics on the number of missing and non-missing observations and their proportions are also available to display.

- `{N_obs}` total number of observations

- {N\_miss} number of missing observations
- {N\_nonmiss} number of non-missing observations
- {p\_miss} percentage of observations missing
- {p\_nonmiss} percentage of observations not missing

Note that for categorical variables, {N\_obs}, {N\_miss} and {N\_nonmiss} refer to the total number, number missing and number non missing observations in the denominator, not at each level of the categorical variable.

It is recommended to use `modify_footnote()` to properly describe the displayed statistics (see examples).

### Caution

The returned table is compatible with all `gtsummary` features applicable to a `tbl_summary` object, like `add_overall()`, `modify_footnote()` or `bold_labels()`.

However, some of them could be inappropriate in such case. In particular, `add_p()` do not take into account the type of displayed statistics and always return the p-value of a comparison test of the current variable according to the by groups, which may be incorrect if the displayed statistics refer to a third variable.

### Author(s)

Joseph Larmarange

### Examples

```
# Example 1 -----
my_stats <- function(data, ...) {
  marker_sum <- sum(data$marker, na.rm = TRUE)
  mean_age <- mean(data$age, na.rm = TRUE)
  dplyr::tibble(
    marker_sum = marker_sum,
    mean_age = mean_age
  )
}

my_stats(trial)

trial |>
tbl_custom_summary(
  include = c("stage", "grade"),
  by = "trt",
  stat_fns = everything() ~ my_stats,
  statistic = everything() ~ "A: {mean_age} - S: {marker_sum}",
  digits = everything() ~ c(1, 0),
  overall_row = TRUE,
  overall_row_label = "All stages & grades"
) |>
add_overall(last = TRUE) |>
modify_footnote(
```

```

    all_stat_cols() ~ "A: mean age - S: sum of marker"
  ) |>
  bold_labels()

# Example 2 -----
# Use `data[[variable]]` to access the current variable
mean_ci <- function(data, variable, ...) {
  test <- t.test(data[[variable]])
  dplyr::tibble(
    mean = test$estimate,
    conf.low = test$conf.int[1],
    conf.high = test$conf.int[2]
  )
}

trial |>
tbl_custom_summary(
  include = c("marker", "ttdeath"),
  by = "trt",
  stat_fns = ~ mean_ci,
  statistic = ~ "{mean} [{conf.low}; {conf.high}]"
) |>
add_overall(last = TRUE) |>
modify_footnote(
  all_stat_cols() ~ "mean [95% CI]"
)

# Example 3 -----
# Use `full_data` to access the full datasets
# Returned statistic can also be a character
diff_to_great_mean <- function(data, full_data, ...) {
  mean <- mean(data$marker, na.rm = TRUE)
  great_mean <- mean(full_data$marker, na.rm = TRUE)
  diff <- mean - great_mean
  dplyr::tibble(
    mean = mean,
    great_mean = great_mean,
    diff = diff,
    level = ifelse(diff > 0, "high", "low")
  )
}

trial |>
tbl_custom_summary(
  include = c("grade", "stage"),
  by = "trt",
  stat_fns = ~ diff_to_great_mean,
  statistic = ~ "{mean} ({level}, diff: {diff})",
  overall_row = TRUE
) |>
bold_labels()

```

---

tbl_hierarchical	<i>Hierarchical Table</i>
------------------	---------------------------

---

**Description****[Experimental]**

*This is an preview of this function. There will be changes in the coming releases, and changes will not undergo a formal deprecation cycle.*

Use these functions to generate hierarchical tables.

- `tbl_hierarchical()`: Calculates *rates* of events (e.g. adverse events) utilizing the denominator and `id` arguments to identify the rows in data to include in each rate calculation. If `variables` contains more than one variable and the last variable in `variables` is an ordered factor, then rates of events by highest level will be calculated.
- `tbl_hierarchical_count()`: Calculates *counts* of events utilizing all rows for each tabulation.

**Usage**

```
tbl_hierarchical(  
  data,  
  variables,  
  by = NULL,  
  id = NULL,  
  denominator = NULL,  
  include = everything(),  
  statistic = "{n} ({p})",  
  overall_row = FALSE,  
  label = NULL  
)
```

```
tbl_hierarchical_count(  
  data,  
  variables,  
  by = NULL,  
  denominator = NULL,  
  include = everything(),  
  overall_row = FALSE,  
  label = NULL  
)
```

**Arguments**

<code>data</code>	(data.frame) a data frame.
-------------------	-------------------------------

variables	( <a href="#">tidy-select</a> ) character vector or tidy-selector of columns in data used to create a hierarchy. Hierarchy will be built with variables in the order given.
by	( <a href="#">tidy-select</a> ) a single column from data. Summary statistics will be stratified by this variable. Default is NULL.
id	( <a href="#">tidy-select</a> ) argument used to subset data to identify rows in data to calculate event rates in <code>tbl_hierarchical()</code> .
denominator	(data.frame, integer) used to define the denominator and enhance the output. The argument is required for <code>tbl_hierarchical()</code> and optional for <code>tbl_hierarchical_count()</code> . The denominator argument must be specified when <code>id</code> is used to calculate event rates.
include	( <a href="#">tidy-select</a> ) variables from hierarchy for which summary statistics should be returned (on the variable label rows) Including the last element of hierarchy has no effect since each level has its own row for this variable. The default is <code>everything()</code> .
statistic	(string) used to specify the summary statistics to display for all variables in <code>tbl_hierarchical()</code> . The default is <code>"{n} ({p})"</code> .
overall_row	(scalar logical) whether an overall summary row should be included at the top of the table. The default is <code>FALSE</code> .
label	( <a href="#">formula-list-selector</a> ) used to override default labels in hierarchical table, e.g. <code>list(AESOC = "System Organ Class")</code> . The default for each variable is the column label attribute, <code>attr(, 'label')</code> . If no label has been set, the column name is used.

## Value

a gsummary table of class `"tbl_hierarchical"` (for `tbl_hierarchical()`) or `"tbl_hierarchical_count"` (for `tbl_hierarchical_count()`).

## Overall Row

An overall row can be added to the table as the first row by specifying `overall_row = TRUE`. Assuming that each row in data corresponds to one event record, this row will count the overall number of events recorded when used in `tbl_hierarchical_count()`, or the overall number of patients recorded with any event when used in `tbl_hierarchical()`.

A label for this overall row can be specified by passing an overall element in `label`. If no overall label has been set and `overall_row = TRUE`, `"Total number of patients with any event"` will be used by `tbl_hierarchical()` and `"Total number of events"` will be used by `tbl_hierarchical_count()`.



**Examples**

```

ADAE_subset <- cards::ADAE |>
  dplyr::filter(
    AESOC %in% unique(cards::ADAE$AESOC)[1:5],
    AETERM %in% unique(cards::ADAE$AETERM)[1:5]
  )

# Example 1 - Event Rates -----
tbl_hierarchical(
  data = ADAE_subset,
  variables = c(AESOC, AETERM),
  by = TRTA,
  denominator = cards::ADSL |> mutate(TRTA = ARM),
  id = USUBJID,
  overall_row = TRUE
)

# Example 2 - Rates by Highest Severity -----
tbl_hierarchical(
  data = ADAE_subset |> mutate(AESEV = factor(AESEV, ordered = TRUE)),
  variables = c(AESOC, AESEV),
  by = TRTA,
  id = USUBJID,
  denominator = cards::ADSL |> mutate(TRTA = ARM),
  include = AESEV,
  label = list(AESEV = "Highest Severity")
)

# Example 3 - Event Counts -----
tbl_hierarchical_count(
  data = ADAE_subset,
  variables = c(AESOC, AETERM, AESEV),
  by = TRTA,
  overall_row = TRUE
)

```

tbl\_likert

*Likert Summary***Description****[Experimental]**

Create a table of ordered categorical variables in a wide format.

**Usage**

```
tbl_likert(
  data,
```

```

  statistic = ~"{n} ({p}%)",
  label = NULL,
  digits = NULL,
  include = everything(),
  sort = c("ascending", "descending")
)

```

### Arguments

data	(data.frame) A data frame.
statistic	( <a href="#">formula-list-selector</a> ) Used to specify the summary statistics for each variable. The default is <code>everything()</code> ~ "{n} ({p}%)".
label	( <a href="#">formula-list-selector</a> ) Used to override default labels in summary table, e.g. <code>list(age = "Age, years")</code> . The default for each variable is the column label attribute, <code>attr(, 'label')</code> . If no label has been set, the column name is used.
digits	( <a href="#">formula-list-selector</a> ) Specifies how summary statistics are rounded. Values may be either integer(s) or function(s). If not specified, default formatting is assigned via <code>assign_summary_digits()</code> .
include	( <a href="#">tidy-select</a> ) Variables to include in the summary table. Default is <code>everything()</code> .
sort	(string) indicates whether levels of variables should be placed in ascending order (the default) or descending.

### Value

a 'tbl\_likert' gtsummary table

### Examples

```

levels <- c("Strongly Disagree", "Disagree", "Agree", "Strongly Agree")
df_likert <- data.frame(
  recommend_friend = sample(levels, size = 20, replace = TRUE) |> factor(levels = levels),
  regret_purchase = sample(levels, size = 20, replace = TRUE) |> factor(levels = levels)
)

# Example 1 -----
tbl_likert_ex1 <-
  df_likert |>
  tbl_likert(include = c(recommend_friend, regret_purchase)) |>
  add_n()
tbl_likert_ex1

# Example 2 -----
# Add continuous summary of the likert scores
list(

```

```
tbl_likert_ex1,
tbl_wide_summary(
  df_likert |> dplyr::mutate(dplyr::across(everything(), as.numeric)),
  statistic = c("{mean}", "{sd}"),
  type = ~"continuous",
  include = c(recommend_friend, regret_purchase)
)
)|>
tbl_merge(tab_spanner = FALSE)
```

tbl\_merge

*Merge tables***Description**

Merge gtsummary tables, e.g. tbl\_regression, tbl\_uvregression, tbl\_stack, tbl\_summary, tbl\_svysummary, etc.

**Usage**

```
tbl_merge(tbls, tab_spanner = NULL)
```

**Arguments**

tbls	(list) List of gtsummary objects to merge
tab_spanner	(character) Character vector specifying the spanning headers. Must be the same length as tbls. The strings are interpreted with gt::md. Must be same length as tbls argument. Default is NULL, and places a default spanning header. If FALSE, no header will be placed.

**Value**

A 'tbl\_merge' object

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
# Side-by-side Regression Models
library(survival)

t1 <-
  glm(response ~ trt + grade + age, trial, family = binomial) %>%
  tbl_regression(exponentiate = TRUE)
```

```

t2 <-
  coxph(Surv(ttdeath, death) ~ trt + grade + age, trial) %>%
  tbl_regression(exponentiate = TRUE)

tbl_merge(
  tbls = list(t1, t2),
  tab_spanner = c("**Tumor Response**", "**Time to Death**")
)

# Example 2 -----
# Descriptive statistics alongside univariate regression, with no spanning header
t3 <-
  trial[c("age", "grade", "response")] %>%
  tbl_summary(missing = "no") %>%
  add_n() %>%
  modify_header(stat_0 ~ "**Summary Statistics**")
t4 <-
  tbl_uvregression(
    trial[c("ttdeath", "death", "age", "grade", "response")],
    method = coxph,
    y = Surv(ttdeath, death),
    exponentiate = TRUE,
    hide_n = TRUE
  )

tbl_merge(tbls = list(t3, t4)) %>%
  modify_spanning_header(everything() ~ NA_character_)

```

---

tbl_regression	<i>Regression model summary</i>
----------------	---------------------------------

---

## Description

This function takes a regression model object and returns a formatted table that is publication-ready. The function is customizable allowing the user to create bespoke regression model summary tables. Review the [tbl\\_regression\(\) vignette](#) for detailed examples.

## Usage

```
tbl_regression(x, ...)

## Default S3 method:
tbl_regression(
  x,
  label = NULL,
  exponentiate = FALSE,
  include = everything(),
  show_single_row = NULL,

```

```

  conf.level = 0.95,
  intercept = FALSE,
  estimate_fun = ifelse(exponentiate, label_style_ratio(), label_style_sigfig()),
  pvalue_fun = label_style_pvalue(digits = 1),
  tidy_fun = broom.helpers::tidy_with_broom_or_parameters,
  add_estimate_to_reference_rows = FALSE,
  conf.int = TRUE,
  ...
)

```

## Arguments

x	(regression model) Regression model object
...	Additional arguments passed to <code>broom.helpers::tidy_plus_plus()</code> .
label	( <a href="#">formula-list-selector</a> ) Used to change variables labels, e.g. <code>list(age = "Age", stage = "Path T Stage")</code>
exponentiate	(scalar logical) Logical indicating whether to exponentiate the coefficient estimates. Default is FALSE.
include	( <a href="#">tidy-select</a> ) Variables to include in output. Default is <code>everything()</code> .
show_single_row	( <a href="#">tidy-select</a> ) By default categorical variables are printed on multiple rows. If a variable is dichotomous (e.g. Yes/No) and you wish to print the regression coefficient on a single row, include the variable name(s) here.
conf.level	(scalar real) Confidence level for confidence interval/credible interval. Defaults to 0.95.
intercept	(scalar logical) Indicates whether to include the intercept in the output. Default is FALSE
estimate_fun	(function) Function to round and format coefficient estimates. Default is <code>label_style_sigfig()</code> when the coefficients are not transformed, and <code>label_style_ratio()</code> when the coefficients have been exponentiated.
pvalue_fun	(function) Function to round and format p-values. Default is <code>label_style_pvalue()</code> .
tidy_fun	(function) Tidier function for the model. Default is to use <code>broom::tidy()</code> . If an error occurs, the tidying of the model is attempted with <code>parameters::model_parameters()</code> , if installed.
add_estimate_to_reference_rows	(scalar logical) Add a reference value. Default is FALSE.
conf.int	(scalar logical) Logical indicating whether or not to include a confidence interval in the output. Default is TRUE.

**Value**

A `tbl_regression` object

**Methods**

The default method for `tbl_regression()` model summary uses `broom::tidy(x)` to perform the initial tidying of the model object. There are, however, a few models that use [modifications](#).

- "parsnip/workflows": If the model was prepared using `parsnip/workflows`, the original model fit is extracted and the original `x=` argument is replaced with the model fit. This will typically go unnoticed; however, if you've provided a custom tidier in `tidy_fun=` the tidier will be applied to the model fit object and not the `parsnip/workflows` object.
- "survreg": The scale parameter is removed, `broom::tidy(x) %>% dplyr::filter(term != "Log(scale)")`
- "multinom": This multinomial outcome is complex, with one line per covariate per outcome (less the reference group)
- "gam": Uses the internal tidier `tidy_gam()` to print both parametric and smooth terms.
- "lmerMod", "glmerMod", "glmmTMB", "glmmadmb", "stanreg", "brmsfit": These mixed effects models use `broom.mixed::tidy(x, effects = "fixed")`. Specify `tidy_fun = broom.mixed::tidy` to print the random components.

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
glm(response ~ age + grade, trial, family = binomial()) |>
tbl_regression(exponentiate = TRUE)
```

---

`tbl_split`

*Split gtsummary table*

---

**Description****[Experimental]**

The `tbl_split` function splits a single `gtsummary` table into multiple tables. Updates to the print method are expected.

**Usage**

```
tbl_split(x, ...)

## S3 method for class 'gtsummary'
tbl_split(x, variables, ...)

## S3 method for class 'tbl_split'
print(x, ...)
```

**Arguments**

x (gtsummary)  
gtsummary table

... These dots are for future extensions and must be empty.

variables (tidy-select)  
variables at which to split the gtsummary table rows (tables will be separated after each of these variables)

**Value**

tbl\_split object

**Examples**

```
tbl <-
  tbl_summary(trial) |>
  tbl_split(variables = c(marker, grade))
```

---

tbl\_stack

*Stack tables*


---

**Description**

Assists in patching together more complex tables. `tbl_stack()` appends two or more gtsummary tables. Column attributes, including number formatting and column footnotes, are retained from the first passed gtsummary object.

**Usage**

```
tbl_stack(tbls, group_header = NULL, quiet = FALSE)
```

**Arguments**

tbls	(list) List of gtsummary objects
group_header	(character) Character vector with table headers where length matches the length of tbls
quiet	(scalar logical) Logical indicating whether to suppress additional messaging. Default is FALSE.

**Value**

A tbl\_stack object

**Author(s)**

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
# stacking two tbl_regression objects
t1 <-
  glm(response ~ trt, trial, family = binomial) %>%
  tbl_regression(
    exponentiate = TRUE,
    label = list(trt ~ "Treatment (unadjusted)")
  )

t2 <-
  glm(response ~ trt + grade + stage + marker, trial, family = binomial) %>%
  tbl_regression(
    include = "trt",
    exponentiate = TRUE,
    label = list(trt ~ "Treatment (adjusted)")
  )

tbl_stack(list(t1, t2))

# Example 2 -----
# stacking two tbl_merge objects
library(survival)
t3 <-
  coxph(Surv(ttdeath, death) ~ trt, trial) %>%
  tbl_regression(
    exponentiate = TRUE,
    label = list(trt ~ "Treatment (unadjusted)")
  )

t4 <-
  coxph(Surv(ttdeath, death) ~ trt + grade + stage + marker, trial) %>%
  tbl_regression(
    include = "trt",
```



```

    exponentiate = TRUE,
    label = list(trt ~ "Treatment (adjusted)")
  )

# first merging, then stacking
row1 <- tbl_merge(list(t1, t3), tab_spanner = c("Tumor Response", "Death"))
row2 <- tbl_merge(list(t2, t4))

tbl_stack(list(row1, row2), group_header = c("Unadjusted Analysis", "Adjusted Analysis"))

```

tbl\_strata

*Stratified gtsummary tables***Description****[Maturing]**

Build a stratified gtsummary table. Any gtsummary table that accepts a data frame as its first argument can be stratified.

- In `tbl_strata()`, the stratified or subset data frame is passed to the function in `.tbl_fun=`, e.g. `purrr::map(data, .tbl_fun)`.
- In `tbl_strata2()`, both the stratified data frame and the strata level are passed to `.tbl_fun=`, e.g. `purrr::map2(data, strata, .tbl_fun)`

**Usage**

```

tbl_strata(
  data,
  strata,
  .tbl_fun,
  ...,
  .sep = ", ",
  .combine_with = c("tbl_merge", "tbl_stack"),
  .combine_args = NULL,
  .header = ifelse(.combine_with == "tbl_merge", "**{strata}**", "{strata}"),
  .stack_group_header = NULL,
  .quiet = NULL
)

```

```

tbl_strata2(
  data,
  strata,
  .tbl_fun,
  ...,
  .sep = ", ",
  .combine_with = c("tbl_merge", "tbl_stack"),
  .combine_args = NULL,

```

```
.header = ifelse(.combine_with == "tbl_merge", "**{strata}**", "{strata}"),
.stack_group_header = NULL,
.quiet = TRUE
)
```

### Arguments

data	(data.frame, survey.design) a data frame or survey object
strata	(tidy-select) character vector or tidy-selector of columns in data to stratify results by
.tbl_fun	(function) A function or formula. If a <i>function</i> , it is used as is. If a formula, e.g. <code>~ .x %&gt;% tbl_summary() %&gt;% add_p()</code> , it is converted to a function. The stratified data frame is passed to this function.
...	Additional arguments passed on to the <code>.tbl_fun</code> function.
.sep	(string) when more than one stratifying variable is passed, this string is used to separate the levels in the spanning header. Default is <code>" , "</code>
.combine_with	(string) One of <code>c("tbl_merge", "tbl_stack")</code> . Names the function used to combine the stratified tables.
.combine_args	(named list) named list of arguments that are passed to function specified in <code>.combine_with</code>
.header	(string) String indicating the headers that will be placed. Default is <code>"**{strata}**"</code> when <code>.combine_with = "tbl_merge"</code> and <code>"{strata}"</code> when <code>.combine_with = "tbl_stack"</code> . Items placed in curly brackets will be evaluated according to <code>glue::glue()</code> syntax. <code>- strata stratum levels - n N within stratum - N Overall N</code> The evaluated value of <code>.header</code> is also available within <code>tbl_strata2(.tbl_fun)</code>
.stack_group_header	<b>[Deprecated]</b>
.quiet	<b>[Deprecated]</b>

### Tips

- `tbl_summary()`
  - The number of digits continuous variables are rounded to is determined separately within each stratum of the data frame. Set the `digits=` argument to ensure continuous variables are rounded to the same number of decimal places.
  - If some levels of a categorical variable are unobserved within a stratum, convert the variable to a factor to ensure all levels appear in each stratum's summary table.

### Author(s)

Daniel D. Sjoberg

**Examples**

```
# Example 1 -----
trial |>
  select(age, grade, stage, trt) |>
  mutate(grade = paste("Grade", grade)) |>
  tbl_strata(
    strata = grade,
    .tbl_fun =
      ~ .x |>
        tbl_summary(by = trt, missing = "no") |>
        add_n(),
    .header = "**{strata}**, N = {n}"
  )

# Example 2 -----
trial |>
  select(grade, response) |>
  mutate(grade = paste("Grade", grade)) |>
  tbl_strata2(
    strata = grade,
    .tbl_fun =
      ~ .x %>%
        tbl_summary(
          label = list(response = .y),
          missing = "no",
          statistic = response ~ "{p}%"
        ) |>
        add_ci(pattern = "{stat} ({ci})") |>
        modify_header(stat_0 = "**Rate (95% CI)**") |>
        modify_footnote(stat_0 = NA),
    .combine_with = "tbl_stack",
    .combine_args = list(group_header = NULL),
    .quiet = TRUE
  ) |>
  modify_caption("**Response Rate by Grade**")
```

tbl\_summary

*Summary table***Description**

The `tbl_summary()` function calculates descriptive statistics for continuous, categorical, and dichotomous variables. Review the [tbl\\_summary vignette](#) for detailed examples.

**Usage**

```
tbl_summary(
  data,
  by = NULL,
```

```

label = NULL,
statistic = list(all_continuous() ~ "{median} ({p25}, {p75})", all_categorical() ~
  "{n} ({p}%)",
digits = NULL,
type = NULL,
value = NULL,
missing = c("ifany", "no", "always"),
missing_text = "Unknown",
missing_stat = "{N_miss}",
sort = all_categorical(FALSE) ~ "alphanumeric",
percent = c("column", "row", "cell"),
include = everything()
)

```

### Arguments

data	(data.frame) A data frame.
by	(tidy-select) A single column from data. Summary statistics will be stratified by this variable. Default is NULL.
label	(formula-list-selector) Used to override default labels in summary table, e.g. <code>list(age = "Age, years")</code> . The default for each variable is the column label attribute, <code>attr(, 'label')</code> . If no label has been set, the column name is used.
statistic	(formula-list-selector) Specifies summary statistics to display for each variable. The default is <code>list(all_continuous() ~ "{median} ({p25}, {p75})", all_categorical() ~ "{n} ({p}%)"</code> . See below for details.
digits	(formula-list-selector) Specifies how summary statistics are rounded. Values may be either integer(s) or function(s). If not specified, default formatting is assigned via <code>assign_summary_digits()</code> . See below for details.
type	(formula-list-selector) Specifies the summary type. Accepted values are <code>c("continuous", "continuous2", "categorical", "dichotomous")</code> . If not specified, default type is assigned via <code>assign_summary_type()</code> . See below for details.
value	(formula-list-selector) Specifies the level of a variable to display on a single row. The <code>gtsummary</code> type selectors, e.g. <code>all_dichotomous()</code> , cannot be used with this argument. Default is NULL. See below for details.
missing, missing_text, missing_stat	Arguments dictating how and if missing values are presented: <ul style="list-style-type: none"> <li>• <code>missing</code>: must be one of <code>c("ifany", "no", "always")</code></li> <li>• <code>missing_text</code>: string indicating text shown on missing row. Default is "Unknown"</li> </ul>

	<ul style="list-style-type: none"> <li>• <code>missing_stat</code>: statistic to show on missing row. Default is "<code>{N_miss}</code>". Possible values are <code>N_miss</code>, <code>N_obs</code>, <code>N_nonmiss</code>, <code>p_miss</code>, <code>p_nonmiss</code>.</li> </ul>
<code>sort</code>	<p>(<a href="#">formula-list-selector</a>)          Specifies sorting to perform for categorical variables. Values must be one of <code>c("alphanumeric", "frequency")</code>. Default is <code>all_categorical(FALSE) ~ "alphanumeric"</code>.</p>
<code>percent</code>	<p>(string)          Indicates the type of percentage to return. Must be one of <code>c("column", "row", "cell")</code>. Default is "column".</p>
<code>include</code>	<p>(<a href="#">tidy-select</a>)          Variables to include in the summary table. Default is <code>everything()</code>.</p>

### Value

a gtsummary table of class "tbl\_summary"

A table of class `c('tbl_summary', 'gtsummary')`

### statistic argument

The statistic argument specifies the statistics presented in the table. The input dictates the summary statistics presented in the table. For example, `statistic = list(age ~ "{mean} ({sd})")` would report the mean and standard deviation for age; `statistic = list(all_continuous() ~ "{mean} ({sd})")` would report the mean and standard deviation for all continuous variables.

The values are interpreted using `glue::glue()` syntax: a name that appears between curly brackets will be interpreted as a function name and the formatted result of that function will be placed in the table.

For categorical variables, the following statistics are available to display: `{n}` (frequency), `{N}` (denominator), `{p}` (percent).

For continuous variables, **any univariate function may be used**. The most commonly used functions are `{median}`, `{mean}`, `{sd}`, `{min}`, and `{max}`. Additionally, `{p##}` is available for percentiles, where `##` is an integer from 0 to 100. For example, `p25: quantile(probs=0.25, type=2)`.

When the summary type is "continuous2", pass a vector of statistics. Each element of the vector will result in a separate row in the summary table.

For both categorical and continuous variables, statistics on the number of missing and non-missing observations and their proportions are available to display.

- `{N_obs}` total number of observations
- `{N_miss}` number of missing observations
- `{N_nonmiss}` number of non-missing observations
- `{p_miss}` percentage of observations missing
- `{p_nonmiss}` percentage of observations not missing

**digits argument**

The digits argument specifies the the number of digits (or formatting function) statistics are rounded to.

The values passed can either be a single integer, a vector of integers, a function, or a list of functions. If a single integer or function is passed, it is recycled to the length of the number of statistics presented. For example, if the statistic is "{mean} ({sd})", it is equivalent to pass 1, c(1, 1), label\_style\_number(digits=1), and list(label\_style\_number(digits=1), label\_style\_number(digits=1)).

Named lists are also accepted to change the default formatting for a single statistic, e.g. list(sd = label\_style\_number(digits=1)).

**type and value arguments**

There are four summary types. Use the type argument to change the default summary types.

- "continuous" summaries are shown on a *single row*. Most numeric variables default to summary type continuous.
- "continuous2" summaries are shown on *2 or more rows*
- "categorical" *multi-line* summaries of nominal data. Character variables, factor variables, and numeric variables with fewer than 10 unique levels default to type categorical. To change a numeric variable to continuous that defaulted to categorical, use type = list(varname ~ "continuous")
- "dichotomous" categorical variables that are displayed on a *single row*, rather than one row per level of the variable. Variables coded as TRUE/FALSE, 0/1, or yes/no are assumed to be dichotomous, and the TRUE, 1, and yes rows are displayed. Otherwise, the value to display must be specified in the value argument, e.g. value = list(varname ~ "level to show")

**Author(s)**

Daniel D. Sjoberg

**See Also**

See [tbl\\_summary vignette](#) for detailed tutorial

See [table gallery](#) for additional examples

Review [list](#), [formula](#), and [selector syntax](#) used throughout gtsummary

**Examples**

```
# Example 1 -----
trial |>
  select(age, grade, response) |>
  tbl_summary()

# Example 2 -----
trial |>
  select(age, grade, response, trt) |>
  tbl_summary(
    by = trt,
```

```

    label = list(age = "Patient Age"),
    statistic = list(all_continuous() ~ "{mean} ({sd})"),
    digits = list(age = c(0, 1))
  )

# Example 3 -----
trial |>
  select(age, marker) |>
  tbl_summary(
    type = all_continuous() ~ "continuous2",
    statistic = all_continuous() ~ c("{median} ({p25}, {p75})", "{min}, {max}"),
    missing = "no"
  )

```

tbl\_survfit

*Survival table***Description**

Function takes a survfit object as an argument, and provides a formatted summary table of the results

**Usage**

```

tbl_survfit(x, ...)

## S3 method for class 'survfit'
tbl_survfit(x, ...)

## S3 method for class 'data.frame'
tbl_survfit(x, y, include = everything(), conf.level = 0.95, ...)

## S3 method for class 'list'
tbl_survfit(
  x,
  times = NULL,
  probs = NULL,
  statistic = "{estimate} ({conf.low}, {conf.high})",
  label = NULL,
  label_header = ifelse(!is.null(times), "**Time {time}**",
    "**{style_sigfig(prob, scale=100)}% Percentile**"),
  estimate_fun = ifelse(!is.null(times), label_style_percent(suffix = "%"),
    label_style_sigfig()),
  missing = "--",
  type = NULL,
  reverse = FALSE,
  quiet = TRUE,

```

```
    ...
  )
```

### Arguments

x	(survfit, list, data.frame) a survfit object, list of survfit objects, or a data frame. If a data frame is passed, a list of survfit objects is constructed using each variable as a stratifying variable.								
...	For <code>tbl_survfit.data.frame()</code> and <code>tbl_survfit.survfit()</code> the arguments are passed to <code>tbl_survfit.list()</code> . They are not used when <code>tbl_survfit.list()</code> is called directly.								
y	outcome call, e.g. <code>y = Surv(ttdeath, death)</code>								
include	Variable to include as stratifying variables.								
conf.level	(scalar numeric) ] Confidence level for confidence intervals. Default is 0.95								
times	(numeric) a vector of times for which to return survival probabilities.								
probs	(numeric) a vector of probabilities with values in (0,1) specifying the survival quantiles to return.								
statistic	(string) string defining the statistics to present in the table. Default is "{estimate} ({conf.low}, {conf.high})"								
label	( <a href="#">formula-list-selector</a> ) List of formulas specifying variables labels, e.g. <code>list(age = "Age, yrs", stage = "Path T Stage")</code> , or a string for a single variable table.								
label_header	(string) string specifying column labels above statistics. Default is "{prob} Percentile" for survival percentiles, and "Time {time}" for n-year survival estimates								
estimate_fun	(function) function to format the Kaplan-Meier estimates. Default is <a href="#">label_style_percent()</a> for survival probabilities and <a href="#">label_style_sigfig()</a> for survival times								
missing	(string) text to fill when estimate is not estimable. Default is "--"								
type	(string or NULL) type of statistic to report. Available for Kaplan-Meier time estimates only, otherwise type is ignored. Default is NULL. Must be one of the following: <table style="margin-left: 40px;"> <thead> <tr> <th>type</th> <th>transformation</th> </tr> </thead> <tbody> <tr> <td>"survival"</td> <td>x</td> </tr> <tr> <td>"risk"</td> <td>1 - x</td> </tr> <tr> <td>"cumhaz"</td> <td>-log(x)</td> </tr> </tbody> </table>	type	transformation	"survival"	x	"risk"	1 - x	"cumhaz"	-log(x)
type	transformation								
"survival"	x								
"risk"	1 - x								
"cumhaz"	-log(x)								
reverse	<b>[Deprecated]</b>								
quiet	<b>[Deprecated]</b>								



**Author(s)**

Daniel D. Sjoberg

**Examples**

```

library(survival)

# Example 1 -----
# Pass single survfit() object
tbl_survfit(
  survfit(Surv(ttdeath, death) ~ trt, trial),
  times = c(12, 24),
  label_header = "**{time} Month**"
)

# Example 2 -----
# Pass a data frame
tbl_survfit(
  trial,
  y = "Surv(ttdeath, death)",
  include = c(trt, grade),
  probs = 0.5,
  label_header = "**Median Survival**"
)

# Example 3 -----
# Pass a list of survfit() objects
list(survfit(Surv(ttdeath, death) ~ 1, trial),
      survfit(Surv(ttdeath, death) ~ trt, trial)) |>
  tbl_survfit(times = c(12, 24))

# Example 4 Competing Events Example -----
# adding a competing event for death (cancer vs other causes)
set.seed(1123)
library(dplyr, warn.conflicts = FALSE, quietly = TRUE)
trial2 <- trial |>
  dplyr::mutate(
    death_cr =
      dplyr::case_when(
        death == 0 ~ "censor",
        runif(n()) < 0.5 ~ "death from cancer",
        TRUE ~ "death other causes"
      ) |>
    factor()
  )

survfit(Surv(ttdeath, death_cr) ~ grade, data = trial2) |>
  tbl_survfit(times = c(12, 24), label = "Tumor Grade")

```

---

tbl_svysummary	<i>Create a table of summary statistics from a survey object</i>
----------------	--

---

### Description

The `tbl_svysummary()` function calculates descriptive statistics for continuous, categorical, and dichotomous variables taking into account survey weights and design.

### Usage

```
tbl_svysummary(
  data,
  by = NULL,
  label = NULL,
  statistic = list(all_continuous() ~ "{median} ({p25}, {p75})", all_categorical() ~
    "{n} ({p}%)" ),
  digits = NULL,
  type = NULL,
  value = NULL,
  missing = c("ifany", "no", "always"),
  missing_text = "Unknown",
  missing_stat = "{N_miss}",
  sort = all_categorical(FALSE) ~ "alphanumeric",
  percent = c("column", "row", "cell"),
  include = everything()
)
```

### Arguments

<code>data</code>	( <code>survey.design</code> ) A survey object created with <code>survey::svydesign()</code>
<code>by</code>	( <a href="#">tidy-select</a> ) A single column from data. Summary statistics will be stratified by this variable. Default is <code>NULL</code> .
<code>label</code>	( <a href="#">formula-list-selector</a> ) Used to override default labels in summary table, e.g. <code>list(age = "Age, years")</code> . The default for each variable is the column label attribute, <code>attr(., 'label')</code> . If no label has been set, the column name is used.
<code>statistic</code>	( <a href="#">formula-list-selector</a> ) Specifies summary statistics to display for each variable. The default is <code>list(all_continuous() ~ "{median} ({p25}, {p75})", all_categorical() ~ "{n} ({p}%)"</code> . See below for details.
<code>digits</code>	( <a href="#">formula-list-selector</a> ) Specifies how summary statistics are rounded. Values may be either integer(s) or function(s). If not specified, default formatting is assigned via <code>assign_summary_digits()</code> . See below for details.

type	( <a href="#">formula-list-selector</a> ) Specifies the summary type. Accepted values are <code>c("continuous", "continuous2", "categorical", "dichotomous")</code> . If not specified, default type is assigned via <code>assign_summary_type()</code> . See below for details.
value	( <a href="#">formula-list-selector</a> ) Specifies the level of a variable to display on a single row. The <code>gtsummary</code> type selectors, e.g. <code>all_dichotomous()</code> , cannot be used with this argument. Default is <code>NULL</code> . See below for details.
missing, missing_text, missing_stat	Arguments dictating how and if missing values are presented: <ul style="list-style-type: none"> <li>• <code>missing</code>: must be one of <code>c("ifany", "no", "always")</code></li> <li>• <code>missing_text</code>: string indicating text shown on missing row. Default is <code>"Unknown"</code></li> <li>• <code>missing_stat</code>: statistic to show on missing row. Default is <code>"{N_miss}"</code>. Possible values are <code>N_miss</code>, <code>N_obs</code>, <code>N_nonmiss</code>, <code>p_miss</code>, <code>p_nonmiss</code>.</li> </ul>
sort	( <a href="#">formula-list-selector</a> ) Specifies sorting to perform for categorical variables. Values must be one of <code>c("alphanumeric", "frequency")</code> . Default is <code>all_categorical(FALSE) ~ "alphanumeric"</code> .
percent	(string) Indicates the type of percentage to return. Must be one of <code>c("column", "row", "cell")</code> . Default is <code>"column"</code> .
include	( <a href="#">tidy-select</a> ) Variables to include in the summary table. Default is <code>everything()</code> .

**Value**

A 'tbl\_svysummary' object

**statistic argument**

The statistic argument specifies the statistics presented in the table. The input is a list of formulas that specify the statistics to report. For example, `statistic = list(age ~ "{mean} ({sd})")` would report the mean and standard deviation for age; `statistic = list(all_continuous() ~ "{mean} ({sd})")` would report the mean and standard deviation for all continuous variables. A statistic name that appears between curly brackets will be replaced with the numeric statistic (see [glue::glue\(\)](#)).

For categorical variables the following statistics are available to display.

- `{n}` frequency
- `{N}` denominator, or cohort size
- `{p}` proportion
- `{p.std.error}` standard error of the sample proportion (on the 0 to 1 scale) computed with [survey::svymean\(\)](#)
- `{deff}` design effect of the sample proportion computed with [survey::svymean\(\)](#)
- `{n_unweighted}` unweighted frequency

- {N\_unweighted} unweighted denominator
- {p\_unweighted} unweighted formatted percentage

For continuous variables the following statistics are available to display.

- {median} median
- {mean} mean
- {mean.std.error} standard error of the sample mean computed with `survey::svymean()`
- {deff} design effect of the sample mean computed with `survey::svymean()`
- {sd} standard deviation
- {var} variance
- {min} minimum
- {max} maximum
- {p##} any integer percentile, where ## is an integer from 0 to 100
- {sum} sum

Unlike `tbl_summary()`, it is not possible to pass a custom function.

For both categorical and continuous variables, statistics on the number of missing and non-missing observations and their proportions are available to display.

- {N\_obs} total number of observations
- {N\_miss} number of missing observations
- {N\_nonmiss} number of non-missing observations
- {p\_miss} percentage of observations missing
- {p\_nonmiss} percentage of observations not missing
- {N\_obs\_unweighted} unweighted total number of observations
- {N\_miss\_unweighted} unweighted number of missing observations
- {N\_nonmiss\_unweighted} unweighted number of non-missing observations
- {p\_miss\_unweighted} unweighted percentage of observations missing
- {p\_nonmiss\_unweighted} unweighted percentage of observations not missing

Note that for categorical variables, {N\_obs}, {N\_miss} and {N\_nonmiss} refer to the total number, number missing and number non missing observations in the denominator, not at each level of the categorical variable.

### type and value arguments

There are four summary types. Use the type argument to change the default summary types.

- "continuous" summaries are shown on a *single row*. Most numeric variables default to summary type continuous.
- "continuous2" summaries are shown on *2 or more rows*

- "categorical" *multi-line* summaries of nominal data. Character variables, factor variables, and numeric variables with fewer than 10 unique levels default to type categorical. To change a numeric variable to continuous that defaulted to categorical, use `type = list(varname ~ "continuous")`
- "dichotomous" categorical variables that are displayed on a *single row*, rather than one row per level of the variable. Variables coded as TRUE/FALSE, 0/1, or yes/no are assumed to be dichotomous, and the TRUE, 1, and yes rows are displayed. Otherwise, the value to display must be specified in the value argument, e.g. `value = list(varname ~ "level to show")`

### Author(s)

Joseph Larmarange

### Examples

```
# Example 1 -----
survey::svydesign(~1, data = as.data.frame(Titanic), weights = ~Freq) |>
  tbl_svsummary(by = Survived, percent = "row", include = c(Class, Age))

# Example 2 -----
# A dataset with a complex design
data(api, package = "survey")
survey::svydesign(id = ~dnum, weights = ~pw, data = apiclus1, fpc = ~fpc) |>
  tbl_svsummary(by = "both", include = c(api00, stype)) |>
  modify_spanning_header(all_stat_cols() ~ "**Survived**")
```

---

tbl\_uvregression

*Univariable regression model summary*


---

### Description

This function estimates univariable regression models and returns them in a publication-ready table. It can create regression models holding either a covariate or an outcome constant.

### Usage

```
tbl_uvregression(data, ...)

## S3 method for class 'data.frame'
tbl_uvregression(
  data,
  y = NULL,
  x = NULL,
  method,
  method.args = list(),
  exponentiate = FALSE,
  label = NULL,
```

```

include = everything(),
tidy_fun = broom.helpers::tidy_with_broom_or_parameters,
hide_n = FALSE,
show_single_row = NULL,
conf.level = 0.95,
estimate_fun = ifelse(exponentiate, label_style_ratio(), label_style_sigfig()),
pvalue_fun = label_style_pvalue(digits = 1),
formula = "{y} ~ {x}",
add_estimate_to_reference_rows = FALSE,
conf.int = TRUE,
...
)

## S3 method for class 'survey.design'
tbl_uvregression(
  data,
  y = NULL,
  x = NULL,
  method,
  method.args = list(),
  exponentiate = FALSE,
  label = NULL,
  include = everything(),
  tidy_fun = broom.helpers::tidy_with_broom_or_parameters,
  hide_n = FALSE,
  show_single_row = NULL,
  conf.level = 0.95,
  estimate_fun = ifelse(exponentiate, label_style_ratio(), label_style_sigfig()),
  pvalue_fun = label_style_pvalue(digits = 1),
  formula = "{y} ~ {x}",
  add_estimate_to_reference_rows = FALSE,
  conf.int = TRUE,
  ...
)

```

## Arguments

data	(data.frame, survey.design) A data frame or a survey design object.
...	Additional arguments passed to <a href="#">broom.helpers::tidy_plus_plus()</a> .
y, x	(expression, string) Model outcome (e.g. <code>y=recurrence</code> or <code>y=Surv(time, recur)</code> ) or covariate (e.g. <code>x=trt</code> . All other column specified in <code>include</code> will be regressed against the constant y or x. Specify one and only one of y or x.
method	(string/function) Regression method or function, e.g. <a href="#">lm</a> , <a href="#">glm</a> , <a href="#">survival::coxph</a> , <code>survey::svyglm</code> , etc. Methods may be passed as functions ( <code>method=lm</code> ) or as strings ( <code>method='lm'</code> ).

method.args	(named list) Named list of arguments passed to method.
exponentiate	(scalar logical) Logical indicating whether to exponentiate the coefficient estimates. Default is FALSE.
label	( <a href="#">formula-list-selector</a> ) Used to change variables labels, e.g. <code>list(age = "Age", stage = "Path T Stage")</code>
include	( <a href="#">tidy-select</a> ) Variables to include in output. Default is <code>everything()</code> .
tidy_fun	(function) Tidier function for the model. Default is to use <code>broom::tidy()</code> . If an error occurs, the tidying of the model is attempted with parameters <code>:model_parameters()</code> , if installed.
hide_n	(scalar logical) Hide N column. Default is FALSE
show_single_row	( <a href="#">tidy-select</a> ) By default categorical variables are printed on multiple rows. If a variable is dichotomous (e.g. Yes/No) and you wish to print the regression coefficient on a single row, include the variable name(s) here.
conf.level	(scalar real) Confidence level for confidence interval/credible interval. Defaults to 0.95.
estimate_fun	(function) Function to round and format coefficient estimates. Default is <a href="#">label_style_sigfig()</a> when the coefficients are not transformed, and <a href="#">label_style_ratio()</a> when the coefficients have been exponentiated.
pvalue_fun	(function) Function to round and format p-values. Default is <a href="#">label_style_pvalue()</a> .
formula	(string) String of the model formula. Uses <a href="#">glue::glue()</a> syntax. Default is " <code>{y} ~ {x}</code> ", where <code>{y}</code> is the dependent variable, and <code>{x}</code> represents a single covariate. For a random intercept model, the formula may be <code>formula = "{y} ~ {x} + (1   gear)"</code> .
add_estimate_to_reference_rows	(scalar logical) Add a reference value. Default is FALSE.
conf.int	(scalar logical) Logical indicating whether or not to include a confidence interval in the output. Default is TRUE.

## Value

A `tbl_uvregression` object

### x and y arguments

For models holding outcome constant, the function takes as arguments a data frame, the type of regression model, and the outcome variable `y`. Each column in the data frame is regressed on the specified outcome. The `tbl_uvregression()` function arguments are similar to the `tbl_regression()` arguments. Review the [tbl\\_uvregression vignette](#) for detailed examples.

You may alternatively hold a single covariate constant. For this, pass a data frame, the type of regression model, and a single covariate in the `x` argument. Each column of the data frame will serve as the outcome in a univariate regression model. Take care using the `x` argument that each of the columns in the data frame are appropriate for the same type of model, e.g. they are all continuous variables appropriate for `lm`, or dichotomous variables appropriate for logistic regression with `glm`.

### Methods

The default method for `tbl_regression()` model summary uses `broom::tidy(x)` to perform the initial tidying of the model object. There are, however, a few models that use [modifications](#).

- "parsnip/workflows": If the model was prepared using `parsnip/workflows`, the original model fit is extracted and the original `x` argument is replaced with the model fit. This will typically go unnoticed; however, if you've provided a custom tidier in `tidy_fun` the tidier will be applied to the model fit object and not the `parsnip/workflows` object.
- "survreg": The scale parameter is removed, `broom::tidy(x) %>% dplyr::filter(term != "Log(scale)")`
- "multinom": This multinomial outcome is complex, with one line per covariate per outcome (less the reference group)
- "gam": Uses the internal tidier `tidy_gam()` to print both parametric and smooth terms.
- "lmerMod", "glmerMod", "glmmTMB", "glmmadmb", "stanreg", "brmsfit": These mixed effects models use `broom.mixed::tidy(x, effects = "fixed")`. Specify `tidy_fun = broom.mixed::tidy` to print the random components.

### Author(s)

Daniel D. Sjoberg

### See Also

See `tbl_regression` [vignette](#) for detailed examples

### Examples

```
# Example 1 -----
tbl_uvregression(
  trial,
  method = glm,
  y = response,
  method.args = list(family = binomial),
  exponentiate = TRUE,
  include = c("age", "grade")
)
```



```
# Example 2 -----
# rounding pvalues to 2 decimal places
library(survival)

tbl_uvregression(
  trial,
  method = coxph,
  y = Surv(ttdeath, death),
  exponentiate = TRUE,
  include = c("age", "grade", "response"),
  pvalue_fun = label_style_pvalue(digits = 2)
)
```

---

tbl_wide_summary	<i>Wide summary table</i>
------------------	---------------------------

---

## Description

### [Experimental]

This function is similar to `tbl_summary()`, but places summary statistics wide, in separate columns. All included variables must be of the same summary type, e.g. all continuous summaries or all categorical summaries (which encompasses dichotomous variables).

## Usage

```
tbl_wide_summary(
  data,
  label = NULL,
  statistic = switch(type[[1]], continuous = c("{median}", "{p25}", "{p75}"), c("{n}",
    "{p}%")),
  digits = NULL,
  type = NULL,
  value = NULL,
  sort = all_categorical(FALSE) ~ "alphanumeric",
  include = everything()
)
```

## Arguments

data	(data.frame) A data frame.
label	(formula-list-selector) Used to override default labels in summary table, e.g. <code>list(age = "Age, years")</code> . The default for each variable is the column label attribute, <code>attr(., 'label')</code> . If no label has been set, the column name is used.

statistic	(character) character vector of the statistics to present. Each element of the vector will result in a column in the summary table. Default is <code>c("{median}", "{p25}", "{p75}")</code> for continuous summaries, and <code>c("{n}", "{p}%")</code> for categorical/dichotomous summaries
digits	( <a href="#">formula-list-selector</a> ) Specifies how summary statistics are rounded. Values may be either integer(s) or function(s). If not specified, default formatting is assigned via <code>assign_summary_digits()</code> . See below for details.
type	( <a href="#">formula-list-selector</a> ) Specifies the summary type. Accepted values are <code>c("continuous", "continuous2", "categorical", "dichotomous")</code> . If not specified, default type is assigned via <code>assign_summary_type()</code> . See below for details.
value	( <a href="#">formula-list-selector</a> ) Specifies the level of a variable to display on a single row. The gtsummary type selectors, e.g. <code>all_dichotomous()</code> , cannot be used with this argument. Default is <code>NULL</code> . See below for details.
sort	( <a href="#">formula-list-selector</a> ) Specifies sorting to perform for categorical variables. Values must be one of <code>c("alphanumeric", "frequency")</code> . Default is <code>all_categorical(FALSE) ~ "alphanumeric"</code> .
include	( <a href="#">tidy-select</a> ) Variables to include in the summary table. Default is <code>everything()</code> .

**Value**

a gtsummary table of class `'tbl_wide_summary'`

**Examples**

```
trial |>
  tbl_wide_summary(include = c(response, grade))

trial |>
  tbl_strata(
    strata = trt,
    ~tbl_wide_summary(.x, include = c(age, marker))
  )
```

---

 theme\_gtsummary

*Available gtsummary themes*


---

**Description**

The following themes are available to use within the gtsummary package. Print theme elements with `theme_gtsummary_journal(set_theme = FALSE) |> print()`. Review the [themes vignette](#) for details.

**Usage**

```

theme_gtsummary_journal(
  journal = c("jama", "lancet", "nejm", "qjecon"),
  set_theme = TRUE
)

theme_gtsummary_compact(set_theme = TRUE, font_size = NULL)

theme_gtsummary_printer(
  print_engine = c("gt", "kable", "kable_extra", "flextable", "huxtable", "tibble"),
  set_theme = TRUE
)

theme_gtsummary_language(
  language = c("de", "en", "es", "fr", "gu", "hi", "is", "ja", "kr", "mr", "nl", "no",
    "pt", "se", "zh-cn", "zh-tw"),
  decimal.mark = NULL,
  big.mark = NULL,
  iqr.sep = NULL,
  ci.sep = NULL,
  set_theme = TRUE
)

theme_gtsummary_continuous2(
  statistic = "{median} ({p25}, {p75})",
  set_theme = TRUE
)

theme_gtsummary_mean_sd(set_theme = TRUE)

theme_gtsummary_eda(set_theme = TRUE)

```

**Arguments**

journal	String indicating the journal theme to follow. One of c("jama", "lancet", "nejm", "qjecon"). Details below.
set_theme	(scalar logical) Logical indicating whether to set the theme. Default is TRUE. When FALSE the named list of theme elements is returned invisibly
font_size	(scalar numeric) Numeric font size for compact theme. Default is 13 for gt tables, and 8 for all other output types
print_engine	String indicating the print method. Must be one of "gt", "kable", "kable_extra", "flextable", "tibble"
language	(string) String indicating language. Must be one of "de" (German), "en" (English), "es" (Spanish), "fr" (French), "gu" (Gujarati), "hi" (Hindi), "is" (Icelandic), "ja"

(Japanese), "kr" (Korean), "nl" (Dutch), "mr" (Marathi), "no" (Norwegian), "pt" (Portuguese), "se" (Swedish), "zh-cn" (Chinese Simplified), "zh-tw" (Chinese Traditional)

If a language is missing a translation for a word or phrase, please feel free to reach out on [GitHub](#) with the translated text.

decimal.mark	(string) The character to be used to indicate the numeric decimal point. Default is "." or getOption("OutDec")
big.mark	(string) Character used between every 3 digits to separate hundreds/thousands/millions/etc. Default is ",", except when decimal.mark = " " when the default is a space.
iqr.sep	(string) String indicating separator for the default IQR in tbl_summary(). If decimal.mark= is NULL, iqr.sep= is ", ". The comma separator, however, can look odd when decimal.mark = " ". In this case the argument will default to an en dash
ci.sep	(string) String indicating separator for confidence intervals. If decimal.mark= is NULL, ci.sep= is ", ". The comma separator, however, can look odd when decimal.mark = " ". In this case the argument will default to an en dash
statistic	Default statistic continuous variables

## Themes

- theme\_gtsummary\_journal(journal)
  - "jama" *The Journal of the American Medical Association*
    - \* Round large p-values to 2 decimal places; separate confidence intervals with "l1 to u1".
    - \* tbl\_summary() Doesn't show percent symbol; use em-dash to separate IQR; run add\_stat\_label()
    - \* tbl\_regression()/tbl\_uvregression() show coefficient and CI in same column
  - "lancet" *The Lancet*
    - \* Use mid-point as decimal separator; round large p-values to 2 decimal places; separate confidence intervals with "l1 to u1".
    - \* tbl\_summary() Doesn't show percent symbol; use em-dash to separate IQR
  - "nejm" *The New England Journal of Medicine*
    - \* Round large p-values to 2 decimal places; separate confidence intervals with "l1 to u1".
    - \* tbl\_summary() Doesn't show percent symbol; use em-dash to separate IQR
  - "qjecon" *The Quarterly Journal of Economics*
    - \* tbl\_summary() all percentages rounded to one decimal place
    - \* tbl\_regression(),tbl\_uvregression() add significance stars with add\_significance\_stars(); hides CI and p-value from output
      - For flextable and huxtable output, the coefficients' standard error is placed below. For gt, it is placed to the right.

- `theme_gtsummary_compact()`
  - tables printed with `gt`, `flextable`, `kableExtra`, or `huxtable` will be compact with smaller font size and reduced cell padding
- `theme_gtsummary_printer(print_engine)`
  - Use this theme to permanently change the default printer.
- `theme_gtsummary_continuous2()`
  - Set all continuous variables to summary type "continuous2" by default
- `theme_gtsummary_mean_sd()`
  - Set default summary statistics to mean and standard deviation in `tbl_summary()`
  - Set default continuous tests in `add_p()` to t-test and ANOVA
- `theme_gtsummary_eda()`
  - Set all continuous variables to summary type "continuous2" by default
  - In `tbl_summary()` show the median, mean, IQR, SD, and Range by default

Use `reset_gtsummary_theme()` to restore the default settings

Review the [themes vignette](#) to create your own themes.

## See Also

[Themes vignette](#)

[set\\_gtsummary\\_theme\(\)](#), [reset\\_gtsummary\\_theme\(\)](#)

## Examples

```
# Setting JAMA theme for gtsummary
theme_gtsummary_journal("jama")
# Themes can be combined by including more than one
theme_gtsummary_compact()

trial |>
  select(age, grade, trt) |>
  tbl_summary(by = trt) |>
  as_gt()

# reset gtsummary themes
reset_gtsummary_theme()
```

---

trial

*Results from a simulated study of two chemotherapy agents*

---

## Description

A dataset containing the baseline characteristics of 200 patients who received Drug A or Drug B. Dataset also contains the outcome of tumor response to the treatment.

**Usage**`trial`**Format**

A data frame with 200 rows—one row per patient

**trt** Chemotherapy Treatment

**age** Age

**marker** Marker Level (ng/mL)

**stage** T Stage

**grade** Grade

**response** Tumor Response

**death** Patient Died

**ttdeath** Months to Death/Censor

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