Package 'tabxplor'

March 8, 2024

Title User-Friendly Tables with Color Helpers for Data Exploration

Version 1.1.3

Description Make it easy to deal with multiple cross-tables in data exploration, by creating them, manipulating them, and adding color helpers to highlight important informations (differences from totals, comparisons between lines or columns, contributions to variance, margins of error, etc.). All functions are pipe-friendly and render data frames which can be easily manipulated. In the same time, time-taking operations are done with 'data.table' to go faster with big dataframes. Tables can be exported to 'Excel' and in html with formats and colors.

URL https://github.com/BriceNocenti/tabxplor

BugReports https://github.com/BriceNocenti/tabxplor/issues

License GPL (>= 3)

Encoding UTF-8

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Suggests fansi (>= 0.5.0), htmltools (>= 0.5.0), knitr, openxlsx (>= 4.0.0), rmarkdown, rstudioapi (>= 0.1), testthat (>= 3.0.0)

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VignetteBuilder knitr

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Author Brice Nocenti [aut, cre]

Maintainer Brice Nocenti <brice.nocenti@gmail.com>

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R topics documented:

complete_partial_totals *Complete partial total rows*

Description

Complete partial total rows

Usage

```
complete_partial_totals(tabs)
```

Arguments

tabs	A table or data.framate containting tabxplor_fmt columns.

Value

The table with completed total rows, total tables, and reference rows.

dplyr_col_modify.tabxplor_grouped_tab *dplyr_col_modify method for class tabxplor_grouped_tab*

Description

dplyr_col_modify method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
dplyr_col_modify(data, cols)
```

Arguments

data	A data frame.
cols	A named list used modify columns. A NULL value should remove an existing column.

Value

An object of class tabxplor_grouped_tab.

dplyr_reconstruct.tabxplor_grouped_tab *dplyr_reconstruct method for class tabxplor_grouped_tab*

Description

dplyr_reconstruct method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
dplyr_reconstruct(data, template)
```

Arguments

data	A data frame.
template	Template to use for restoring attributes

Value

An object of class tabxplor_grouped_tab.

dplyr_row_slice.tabxplor_grouped_tab *dplyr_row_slice method for class tabxplor_grouped_tab*

Description

dplyr_row_slice method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
dplyr_row_slice(data, i, ...)
```

Arguments

data	A data frame.
i	A numeric or logical vector that indexes the rows of .data.
...	Future parameters.

Value

An object of class tabxplor_grouped_tab.

fct_recode_helper *fct_recode helper to recode multiple variables*

Description

fct_recode helper to recode multiple variables

Usage

```
fct_recode_helper(
  .data,
  .cols = -where(is.numeric),
  .data_out_name,
  cat = TRUE
)
```

Arguments

.data	The data frame.
.cols	<tidy-select> The variables to recode.
.data_out_name	The name of the output data frame, if different from the input data frame.
cat	By default the result is written in the console if there are less than 6 variables, written in a temporary file and opened otherwise. Set to false to get a data frame with a character variable instead.

Value

When the number of variables is less than 5, a text in console as a side effect. With more than 5 variables, a temporary R file. A tibble with the recode text as a character variable is returned invisibly (or as main result if cat = TRUE).

fmt *Create a vector of class formatted numbers*

Description

fmt vectors, of class tabxplor_fmt, powers **tabxplor** and tab tibbles. As a record, they stores all data necessary to calculate percentages, Chi2 metadata or confidence intervals, but also to format and color the table to help the user read it. You can access this data with vctrs::field, or change it with vctrs:field<-. A fmt vector have 13 fields : n, digits, display, wn, pct, mean, diff, ctr, var, ci, in_totrow, in_tottab, in_refrow. Other arguments are attributes, attached not to each value, but to the whole vector, like type, totcol or color. You can get them with attr and modify them with attr<-. Special functions listed below are made to facilitate programming with with **tabxplor** formatted numbers. taxplfmt vectors can use all standard operations, like +, -, sum(), or c(), using **vctrs**.

Usage

```
fmt(
  n = integer(),
  type = "n",
  digits = rep(0L, length(n)),
 display = dplyr::case_when(type == "mean" ~ "mean", type %in% c("row", "col", "all",
    "all_tabs") ~ "pct", TRUE ~ "n"),
  wn = rep(NA_real_, length(n)),
  pct = rep(NA_real_, length(n)),
  mean = rep(NA_real_, length(n)),
  diff = rep(NA_real_, length(n)),
  ctr = rep(NA_real_, length(n)),
  var = rep(NA_real_, length(n)),
  ci = rep(NA_real_, length(n)),
  in_totrow = rep(FALSE, length(n)),
  in_tottab = rep(FALSE, length(n)),
  in_refrow = rep(FALSE, length(n)),
  comp_all = NA,
  diff_type = "",
  ci_type = "",
  col_var = "",
  totcol = FALSE,
  refcol = FALSE,
  color = ""
)

is_fmt(x)

get_num(x)

set_num(x, value)

get_type(x, ...)

set_type(x, type)

is_totrow(x, ...)

as_totrow(x, in_totrow = TRUE)

is_tottab(x, ...)

as_tottab(x, in_tottab = TRUE)

is_totcol(x, ...)

as_totcol(x, totcol = TRUE)
```

```
is_refrow(x, ...)

as_refrow(x, in_refrow = TRUE)

get_comp_all(x, replace_na = TRUE)

set_comp_all(x, comp_all = FALSE)

get_diff_type(x, ...)

set_diff_type(x, diff_type)

get_ci_type(x, ...)

set_ci_type(x, ci_type)

get_col_var(x, ...)

set_col_var(x, col_var)

is_refcol(x, ...)

as_refcol(x, refcol = TRUE)

get_color(x, ...)

set_color(x, color)

get_digits(x)

set_digits(x, value)
```

Arguments

n	The underlying count, as an integer vector of length n(). It is used to calculate confidence intervals.
type	The type of the column, which defines the type of background calculation to be made (as a single string, since it's not a field but an attribute) : • "n": counts • "mean": mean column (from numeric variables) • "row": row percentages • "col": column percentages • "all": frequencies by subtable/group (i.e. by tab_vars) • "all_tabs": frequencies for the whole table
digits	The number of digits, as an integer, or an integer vector the length of n.
display	The display type : the name of the field you want to show when printing the vector. Among "n", "wn", "pct", "diff", "ctr", "mean", "var", "ci", "pct_ci"

(percentages with visible confidence interval), "mean_ci" (means with visible confidence interval). As a single string, or a character vector the length of n.

wn — The underlying weighted counts, as a double vector the length of n. It is used in certain operations on fmt, like means.

pct — The percentages, as a double vector the length of n. Calculate with tab_pct.

mean — The means, as a double vector the length of n.

diff — The differences (from totals or first cells), as a double vector the length of n. Used to set colors for means and row or col percentages. Calculate with tab_pct.

ctr — The contributions of cells to (sub)tables variances, as a double vector the length of n. Used to print colors when color = "contrib". The mean contribution of each (sub)table is written on total rows (then, colors don't print well without total rows). Calculate with tab_chi2.

var — The cells variances, as a double vector the length of n. Used with type = "mean" to calculate confidence intervals. Calculate with tab_plain.

ci — The confidence intervals, as a double vector the length of n. Used to print colors ("diff_ci", "after_ci"). Calculate with tab_ci.

in_totrow — TRUE when the cell is part of a total row

in_tottab — TRUE when the cell is part of a total table

in_refrow — TRUE when the cell is part of a reference row (cf. diff_type)

comp_all — FALSE when the comparison level is the subtable/group, TRUE when it is the whole table

diff_type — The type of difference of the vector (calculate with tab_pct) :

- "" or "no": no differences have been calculated
- "tot": the reference row (or column) is the total row (or column)
- "first": the reference row (or column) is the first row (or column)

ci_type — The type of confidence intervals of the vector (calculate with tab_ci) :

- "" or "no": no ci have been calculated
- "cell": absolute confidence intervals of cells percentages.
- "diff": confidence intervals of the difference between a cell and the relative total cell (or relative first cell when diff_type = "first").
- "auto": "diff" for means and row/col percentages, "cell" for frequencies ("all", "all_tabs").

col_var — The name of the col_var used to calculate the vector

totcol — TRUE when the vector is a total column

refcol — TRUE when the vector is a reference column

color — The type of color to print :

- "no": no colors are printed.
- "diff": color percentages and means based on cells differences from totals (or from first cells when diff = "first").

- `"diff_ci"`: color pct and means based on cells differences from totals or first cells, removing coloring when the confidence interval of this difference is higher than the difference itself.
- `"after_ci"`: idem, but cut off the confidence interval from the difference first.
- `"contrib"`: color cells based on their contribution to variance (except mean columns, from numeric variables).

`x`	The object to test, to get a field in, or to modify.
`value`	The value you want to inject in some `fmt` vector's vctrs::field or attribute using a given "set" function.
`...`	Used in methods to add arguments in the future.
`replace_na`	By default, `get_comp_all` takes NA in comparison level to be a `FALSE` (=comparison at subtables/groups level). Set to `FALSE` to avoid this behavior.

Value

A vector of class `tabxplor_fmt`.

A logical vector.

A double vector.

A modified fmt vector.

A character vector with the vectors type.

A modified fmt vector.

A logical vector with the fmt vectors totrow field.

A modified fmt vector with totrow field changed.

A logical vector with the fmt vectors tottab field.

A modified fmt vector with tottab field changed.

A logical vector with the fmt vectors totcol attribute.

A modified fmt vector with totcol attribute changed.

A logical vector with the fmt vectors in_refrow field

A modified fmt vector with in_refrom field changed.

A modified fmt vector with comp attribute changed.

A logical vector with the fmt vectors type attributes

A modified fmt vector.

A logical vector with the fmt vectors ci_type attributes

A modified fmt vector.

A logical vector with the fmt vectors col_var attributes

A modified fmt vector.

A logical vector with the fmt vectors is_refcol attributes

A modified fmt vector.

A logical vector with the fmt vectors color attributes

A modified fmt vector.

Functions

- `is_fmt()`: a test function for class fmt.
- `get_num()`: get the currently displayed field
- `set_num()`: set the currently displayed field (not changing display type)
- `get_type()`: get types of fmt columns (at `fmt` level or `tab` level)
- `set_type()`: set the column type attribute of a `fmt` vector
- `is_totrow()`: test function to detect cells in total rows (at `fmt` level or `tab` level)
- `as_totrow()`: set the "in_totrow" field (belong to total row)
- `is_tottab()`: test function to detect cells in total tables (at `fmt` level or `tab` level)
- `as_tottab()`: set the "in_tottab" field (belong to total table)
- `is_totcol()`: test function for total columns (at `fmt` level or `tab` level)
- `as_totcol()`: set the "totcol" attribute of a `fmt` vector
- `is_refrow()`: test function to detect cells in reference rows (at `fmt` level or `tab` level)
- `as_refrow()`: set the "in_refrow" field (belong to reference row)
- `get_comp_all()`: get comparison level of fmt columns
- `set_comp_all()`: set the comparison level attribute of a `fmt` vector
- `get_diff_type()`: get differences type of fmt columns (at `fmt` level or `tab` level)
- `set_diff_type()`: set the differences type attribute of a `fmt` vector
- `get_ci_type()`: get confidence intervals type of fmt columns (at `fmt` level or `tab` level)
- `set_ci_type()`: set the confidence intervals type attribute of a `fmt` vector
- `get_col_var()`: get names of column variable of fmt columns (at `fmt` level or `tab` level)
- `set_col_var()`: set the "col_var" attribute of a `fmt` vector
- `is_refcol()`: test function for reference columns (at `fmt` level or `tab` level)
- `as_refcol()`: set the "ref_col" attribute of a `fmt` vector
- `get_color()`: get color (at `fmt` level or `tab` level)
- `set_color()`: set the "color" attribute of a `fmt` vector
- `get_digits()`: get the "digits" field
- `set_digits()`: set the "digits" field

Examples

```
library(dplyr)

f <- fmt(n = c(7, 19, 2), type = "row", pct = c(0.25, 0.679, 0.07))
f

# To get the currently displayed field :
get_num(f)

# To modify the currently displayed field :
set_num(f, c(1, 0, 0))
```

```
# See all the underlying fields of a fmt vector (a data frame with a number of rows
#  equal to the length of the vector) :
vctrs::vec_data(f)

# To get the numbers of digits :
vctrs::field(f, "digits")
f$digits

# To get the count :
vctrs::field(f, "n")
f$n

# To get the display :
vctrs::field(f, "display")
f$display

# To modify a field, you can use `dplyr::mutate` on the fmt vector,
# referring to the names of the columns of the underlying data.frame (`vctrs::vec_data`) :
vctrs::`field<-`(f, "pct", c(1, 0, 0))
mutate(f, pct = c(1, 0, 0))

# See all the attributes of a fmt vector :
attributes(f)

# To modify the "type" attribute of a fmt vector :
set_type(f, "col")

# To modify the "color" attribute of a fmt vector :
set_color(f, "contrib")




tabs <- tab(starwars, sex, hair_color, gender, na = "drop", pct = "row",
            rare_to_other = TRUE, n_min = 5)

# To identify the total columns, and work with them :
is_totcol(tabs)
tabs %>% mutate(across(where(is_totcol), ~ "total column"))

# To identify the total rows, and work with them :
is_totrow(tabs)
tabs %>%
  mutate(across(
    where(is_fmt),
    ~ if_else(is_totrow(.), true = "into_total_row", false = "normal_cell")
  ))

# To identify the total tables, and work with them :
tottabs <- is_tottab(tabs)
tabs %>% tibble::add_column(tottabs) %>%
```

```
  mutate(total = if_else(tottabs, "part of a total table", "normal cell"))

# To access the displayed numbers, as numeric vectors :
tabs %>% mutate(across(where(is_fmt), get_num))

# To access the displayed numbers, as character vectors (without colors) :
tabs %>% mutate(across(where(is_fmt), format))

# To access the (non-displayed) differences of the cells percentages from totals :
tabs %>% mutate(across(where(is_fmt), ~ vctrs::field(., "diff")))


# To do more complex operations, like creating a new column with standard deviation and
# print it with 2 decimals, use `dplyr::mutate` on all the fmt columns of a table :

tab_num(forcats::gss_cat, race, c(age, tvhours), marital, digits = 1L, comp = "all") |>
  dplyr::mutate(dplyr::across( #Mutate over the whole table.
    c(age, tvhours),
    ~ dplyr::mutate(.,         #Mutate over each fmt vector's underlying data.frame.
                    var     = sqrt(var),
                    display = "var",
                    digits  = 2L) |>
      set_color("no"),
    .names = "{.col}_sd"
  ))
```

`fmt_get_color_code` *Get HTML Color Code of a fmt vector*

Description

Get HTML Color Code of a fmt vector

Usage

```
fmt_get_color_code(x, type = "text", theme = "light", html_24_bit = NULL)
```

Arguments

`x`	The fmt vector to get the html color codes from.
`type`	The style type in `set_color_style` and `get_color_style`, `"text"` to color the text, `"bg"` to color the background.
`theme`	For `set_color_style` and `get_color_style`, is your console or html table background `"light"` or `"dark"` ? Default to RStudio theme.
`html_24_bit`	Should specific 24bits colors palettes be used for html tables ? With light themes only. Default to `getOption("tabxplor.color_html_24_bit")`

Value

A character vector with html color codes, of the length of the initial vector.

Examples

```
tabs <- tab(forcats::gss_cat, race, marital, pct = "row", color = "diff")
dplyr::mutate(tabs, across(where(is_fmt), fmt_get_color_code))
```

---

format.tabxplor_fmt *Print method for class tabxplor_fmt*

Description

Print method for class tabxplor_fmt

Usage

```
## S3 method for class 'tabxplor_fmt'
format(x, ..., html = FALSE, na = NA)
```

Arguments

x	A fmt object.
...	Other parameters.
html	Should html tags be added (to print confidence intervals as subscripts) ?
na	How NAs should be printed. Default to NA.

Value

The fmt printed in a character vector.

---

get_ci_type.data.frame *Get confidence intervals type of fmt columns*

Description

Get confidence intervals type of fmt columns

Usage

```
## S3 method for class 'data.frame'
get_ci_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A character vector with the ci_type attributes.

---

get_ci_type.default *Get confidence intervals type of fmt columns*

Description

Get confidence intervals type of fmt columns

Usage

```
## Default S3 method:
get_ci_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the ci_type attribute.

---

get_ci_type.tabxplor_fmt *Get confidence intervals type of fmt columns*

Description

Get confidence intervals type of fmt columns

Usage

```
## S3 method for class 'tabxplor_fmt'
get_ci_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the ci_type attribute.

`get_color.data.frame` *Get color*

Description

Get color

Usage

```
## S3 method for class 'data.frame'
get_color(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A character vector with the color attributes.

`get_color.default` *Get color*

Description

Get color

Usage

```
## Default S3 method:
get_color(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the color attribute.

get_color.tabxplor_fmt *Get color*

Description

Get color

Usage

```
## S3 method for class 'tabxplor_fmt'
get_color(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the color attribute.

get_col_var.data.frame *Get names of column variable of fmt columns*

Description

Get names of column variable of fmt columns

Usage

```
## S3 method for class 'data.frame'
get_col_var(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A character vector with the col_var attributes.

get_col_var.default *Get names of column variable of fmt columns*

Description

Get names of column variable of fmt columns

Usage

```
## Default S3 method:
get_col_var(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the col_var attribute.

get_col_var.tabxplor_fmt *Get names of column variable of fmt columns*

Description

Get names of column variable of fmt columns

Usage

```
## S3 method for class 'tabxplor_fmt'
get_col_var(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the col_var attribute.

get_diff_type.data.frame *Get differences type of fmt columns*

Description

Get differences type of fmt columns

Usage

```
## S3 method for class 'data.frame'
get_diff_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A character vector with the diff_type attribute.

get_diff_type.default *Get differences type of fmt columns*

Description

Get differences type of fmt columns

Usage

```
## Default S3 method:
get_diff_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the diff_type attribute.

get_diff_type.tabxplor_fmt *Get differences type of fmt columns*

Description

Get differences type of fmt columns

Usage

```
## S3 method for class 'tabxplor_fmt'
get_diff_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the diff_type attribute.

get_type.data.frame *Get types of fmt columns*

Description

Get types of fmt columns

Usage

```
## S3 method for class 'data.frame'
get_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A character vector with the data.frame column's types.

get_type.default *Get types of fmt columns*

Description

Get types of fmt columns

Usage

```
## Default S3 method:
get_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

An empty character vector.

get_type.tabxplor_fmt *Get types of fmt columns*

Description

Get types of fmt columns

Usage

```
## S3 method for class 'tabxplor_fmt'
get_type(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single string with the vector's type.

group_by.tabxplor_tab *group_by method for class tabxplor_tab*

Description

group_by method for class tabxplor_tab

Usage

```
## S3 method for class 'tabxplor_tab'
group_by(.data, ..., .add = FALSE, .drop = dplyr::group_by_drop_default(.data))
```

Arguments

.data	A tibble of class tabxplor_tab.
...	Variables or computations to group by.
.add	When FALSE, the default, group_by() will override existing groups. To add to the existing groups, use .add = TRUE.
.drop	Drop groups formed by factor levels that don't appear in the data? The default is TRUE except when .data has been previously grouped with .drop = FALSE.

Value

A tibble of class tabxplor_grouped_tab.

is_refcol.data.frame *Test function for reference columns*

Description

Test function for reference columns

Usage

```
## S3 method for class 'data.frame'
is_refcol(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A character vector with the ref_col attributes.

is_refcol.default *Test function for reference columns*

Description

Test function for reference columns

Usage

```
## Default S3 method:
is_refcol(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the ref_col attribute.

is_refcol.tabxplor_fmt *Test function for reference columns*

Description

Test function for reference columns

Usage

```
## S3 method for class 'tabxplor_fmt'
is_refcol(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single character with the ref_col attribute.

is_refrow.data.frame *Test function to detect cells in reference rows*

Description

Test function to detect cells in reference rows

Usage

```
## S3 method for class 'data.frame'
is_refrow(x, ..., partial = TRUE)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.
partial	Should partial reference rows be counted as reference rows ? Default to FALSE.

Value

A list of logical vectors with the in_refrow fields.

is_refrow.default *Test function to detect cells in reference rows*

Description

Test function to detect cells in reference rows

Usage

```
## Default S3 method:
is_refrow(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A logical vector with FALSE, the length of x.

is_refrow.tabxplor_fmt *Test function to detect cells in reference rows*

Description

Test function to detect cells in reference rows

Usage

```
## S3 method for class 'tabxplor_fmt'
is_refrow(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A logical vector with the in_refrow field.

is_totcol.data.frame *Test function for total columns*

Description

Test function for total columns

Usage

```
## S3 method for class 'data.frame'
is_totcol(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A logical vector, with the data.frame column's totcol attributes.

is_totcol.default *Test function for total columns*

Description

Test function for total columns

Usage

```
## Default S3 method:
is_totcol(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single logical vector with the totcol attribute

is_totcol.tabxplor_fmt *Test function for total columns*

Description

Test function for total columns

Usage

```
## S3 method for class 'tabxplor_fmt'
is_totcol(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A single logical vector with the totcol attribute

is_totrow.data.frame *Test function to detect cells in total rows*

Description

Test function to detect cells in total rows

Usage

```
## S3 method for class 'data.frame'
is_totrow(x, ..., partial = FALSE)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.
partial	Should partial total rows be counted as total rows ? Default to FALSE.

Value

A list of logical vectors, with the data.frame column's totrow fields.

is_totrow.default *Test function to detect cells in total rows*

Description

Test function to detect cells in total rows

Usage

```
## Default S3 method:
is_totrow(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A logical vector with FALSE.

is_totrow.tabxplor_fmt *Test function to detect cells in total rows*

Description

Test function to detect cells in total rows

Usage

```
## S3 method for class 'tabxplor_fmt'
is_totrow(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A logical vector with the totrow field.

is_tottab.data.frame *Test function to detect cells in total tables*

Description

Test function to detect cells in total tables

Usage

```
## S3 method for class 'data.frame'
is_tottab(x, ..., partial = FALSE)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.
partial	Should partial total tabs be counted as total tabs ? Default to FALSE.

Value

A list of logical vectors, with the data.frame column's tottab fields.

is_tottab.default *Test function to detect cells in total tables*

Description

Test function to detect cells in total tables

Usage

```
## Default S3 method:
is_tottab(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A logical vector with FALSE.

is_tottab.tabxplor_fmt *Test function to detect cells in total tables*

Description

Test function to detect cells in total tables

Usage

```
## S3 method for class 'tabxplor_fmt'
is_tottab(x, ...)
```

Arguments

x	The object to test, to get a field in, or to modify.
...	Used in methods to add arguments in the future.

Value

A logical vector with the tottab field.

mutate.tabxplor_fmt *mutate method to access vctrs::fields of tabxplor_fmt vectors*

Description

mutate method to access vctrs::fields of tabxplor_fmt vectors

Usage

```
## S3 method for class 'tabxplor_fmt'
mutate(.data, ...)
```

Arguments

.data	A tabxplor_fmt column.
...	<data-masking> Name-value pairs. The name gives the name of the column in the output (do not change it). The value can be: • A vector of length 1, which will be recycled to the correct length. • A vector the same length as the current group (or the whole data frame if ungrouped).

Value

An object of class `tabxplor_fmt`.

new_tab *A constructor for class tabxplor_tab*

Description

A constructor for class tabxplor_tab

Usage

```
new_tab(
  tabs = tibble::tibble(),
  subtext = "",
 chi2 = tibble::tibble(tables = character(), pvalue = double(), df = integer(), cells =
    integer(), variance = double(), count = integer()),
  ...,
  class = character()
)

new_grouped_tab(
```

```
  tabs = tibble::tibble(),
  groups,
  subtext = "",
 chi2 = tibble::tibble(tables = character(), pvalue = double(), df = integer(), cells =
    integer(), variance = double(), count = integer()),
  ...,
  class = character()
)
```

Arguments

tabs	A table, stored into a tibble data.frame. It is generally made with tab, tab_many or tab_plain.
subtext	A character vector to print legend lines under the table.
chi2	A tibble storing information about pvalues and variances, to fill with tab_chi2.
...	Needed to implement subclasses.
class	Needed to implement subclasses.
groups	The grouping data.

Value

A tibble of class tabxplor_tab.

A tibble of class tabxplor_grouped_tab.

---

pillar_shaft.tabxplor_fmt

Pillar_shaft method to print class fmt in a tibble column

Description

Pillar_shaft method to print class fmt in a tibble column

Usage

```
## S3 method for class 'tabxplor_fmt'
pillar_shaft(x, ...)
```

Arguments

x	A fmt object.
...	Other parameter.

Value

A fmt printed in a pillar.

pillar_shaft.tab_chi2_fmt *Print Chi2 tables columns*

Description

Print Chi2 tables columns

Usage

```
## S3 method for class 'tab_chi2_fmt'
pillar_shaft(x, ...)
```

Arguments

x	A fmt object.
...	Other parameter.

Value

A Chi2 table column printed in a pillar.

print.tabxplor_grouped_tab *Printing method for class tabxplor_grouped_tab*

Description

Printing method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
print(
  x,
  width = NULL,
  ...,
  n = 100,
  max_extra_cols = NULL,
  max_footer_lines = NULL,
  min_row_var = 30
)
```

Arguments

x	Object to format or print.
width	Width of text output to generate.
...	Passed on to tbl_format_setup().
n	Number of rows to show.
max_extra_cols	Number of extra columns to print abbreviated information for, if the width is too small for the entire tibble.
max_footer_lines	Maximum number of footer lines.
min_row_var	Minimum number of characters for the row variable. Default to 30.

Value

A printed grouped table.

---

print.tabxplor_tab *Printing method for class tabxplor_tab*

Description

Printing method for class tabxplor_tab

Usage

```
## S3 method for class 'tabxplor_tab'
print(
  x,
  width = NULL,
  ...,
  n = 100,
  max_extra_cols = NULL,
  max_footer_lines = NULL,
  min_row_var = 30
)
```

Arguments

x	Object to format or print.
width	Width of text output to generate.
...	Passed on to tbl_format_setup().
n	Number of rows to show.
max_extra_cols	Number of extra columns to print abbreviated information for, if the width is too small for the entire tibble.
max_footer_lines	Maximum number of footer lines.
min_row_var	Minimum number of characters for the row variable. Default to 30.

Value

A printed table.

`relocate.tabxplor_grouped_tab`
relocate method for class tabxplor_grouped_tab

Description

relocate method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
relocate(.data, ...)
```

Arguments

`.data`	A tibble of class `tabxplor_tab`.
`...`	Columns to move. will move columns to the left-hand side; specifying both is an error.

Value

An object of class `tabxplor_grouped_tab`.

`rename.tabxplor_grouped_tab`
rename method for class tabxplor_grouped_tab

Description

rename method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
rename(.data, ...)
```

Arguments

`.data`	A tibble of class `tabxplor_tab`.
`...`	Use `new_name = old_name` to rename selected variables.

Value

An object of class `tabxplor_grouped_tab`.

rename_with.tabxplor_grouped_tab *rename_with method for class tabxplor_grouped_tab*

Description

rename_with method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
rename_with(.data, .fn, .cols = dplyr::everything(), ...)
```

Arguments

.data	A tibble of class tabxplor_tab.
.fn	A function used to transform the selected .cols. Should return a character vector the same length as the input.
.cols	Columns to rename; defaults to all columns.
...	Additional arguments passed onto .fn.

Value

An object of class tabxplor_grouped_tab.

rowwise.tabxplor_grouped_tab *rowwise method for class tabxplor_grouped_tab*

Description

rowwise method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
rowwise(data, ...)
```

Arguments

data	A tibble of class tabxplor_tab.
...	Variables to be preserved when calling summarise(). This is typically a set of variables whose combination uniquely identify each row.

Value

An object of class tabxplor_grouped_tab and rowwise_df.

rowwise.tabxplor_tab *rowwise method for class tabxplor_tab*

Description

rowwise method for class tabxplor_tab

Usage

```
## S3 method for class 'tabxplor_tab'
rowwise(data, ...)
```

Arguments

data	A tibble of class tabxplor_tab.
...	Variables to be preserved when calling summarise(). This is typically a set of variables whose combination uniquely identify each row.

Value

A tibble of class tabxplor_grouped_tab and rowwise_df.

select.tabxplor_grouped_tab *select method for class tabxplor_grouped_tab*

Description

select method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
select(.data, ...)
```

Arguments

.data	A tibble of class tabxplor_tab.
...	One or more unquoted expressions separated by commas. Variable names can be used as if they were positions in the data frame, so expressions like x:y can be used to select a range of variables.

Value

An object of class tabxplor_grouped_tab.

summarise.tabxplor_grouped_tab *summarise method for class tabxplor_grouped_tab*

Description

summarise method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
summarise(.data, ..., .groups = NULL)
```

Arguments

.data	A tibble of class tabxplor_tab.
...	Name-value pairs of summary functions. The name will be the name of the variable in the result.
.groups	Grouping structure of the result.

Value

An object of class tabxplor_grouped_tab.

tab *Single cross-table, with color helpers*

Description

A full-featured function to create, manipulate and format single cross-tables, using colors to make the printed tab more easily readable (in R terminal or exported to Excel with tab_xl). Since objects of class tab are also of class tibble, you can then use all **dplyr** verbs to modify the result, like select, like arrange, filter or mutate. Wrapper around the more powerful tab_many.

Usage

```
tab(
  data,
  row_var,
  col_var,
  tab_vars,
  wt,
  sup_cols,
  na = "keep",
  digits = 0,
```

```
  pct = "no",
  color = "no",
  diff = "tot",
  comp = "tab",
  totaltab = "line",
  totaltab_name = "Ensemble",
  tot = c("row", "col"),
  total_names = "Total",
  chi2 = FALSE,
  ci = "no",
  conf_level = 0.95,
  subtext = "",
  cleannames = NULL,
  rare_to_other = FALSE,
  n_min = 30,
  other_level = "Others",
  filter
)
```

Arguments

`data`	A data frame.
`row_var, col_var`	The row variable, which will be printed with one level per line, and the column variable, which will be printed with one level per column. For numeric variables means are calculated, in a single column.
`tab_vars`	<tidy-select> Tab variables : a subtable is made for each combination of levels of the selected variables. Leave empty to make a simple cross-table. All `tab_vars` are converted to factor.
`wt`	A weight variable, of class numeric. Leave empty for unweighted results.
`sup_cols`	<tidy-select> Supplementary columns variables, with only the first level printed, and row percentages (for numeric variables, a mean will be calculated for each `row_var`). To pass many variables you may use syntax `sup_cols = c(sup_col1, sup_col2, ...)`. To keep all levels of other `col_vars`, or other types of percentages, use `tab_many` instead.
`na`	The policy to adopt for missing values, as a single string : • `"keep"`: by default, `NA`'s of row, col and tab variables are printed as an explicit `"NA"` level. • `"drop"`: remove `NA`'s in row, col and tab variables before calculations are done. Supplementary columns are then calculated for observations with no `NA` in any of the row, col and tab variables.
`digits`	The number of digits to print, as a single integer. To print a different number of digits for each `sup_cols`, an integer vector of length 1 + `sup_cols` (the first being the number of digits for the base table).
`pct`	The type of percentages to calculate : • `"row"`: row percentages.

- "col": column percentages.
- "all": frequencies for each subtable/group, if there is tab_vars.
- "all_tabs": frequencies for the whole (set of) table(s).

color The type of colors to print, as a single string :

- "no": by default, no colors are printed.
- "diff": color percentages and means based on cells differences from totals (or from first cells when diff = "first").
- "diff_ci": color pct and means based on cells differences from totals or first cells, removing coloring when the confidence interval of this difference is higher than the difference itself.
- "after_ci": idem, but cut off the confidence interval from the difference first.
- "contrib": color cells based on their contribution to variance (except mean columns, from numeric variables).
- "auto": frequencies (pct = "all", pct = "all_tabs") and counts are colored with "contrib". When ci = "diff", row and col percentages are colored with "after_ci" ; otherwise they are colored with "diff".

diff The reference cell to calculate differences (used to print colors) :

- "tot": by default, cells differences from total rows are calculated with pct = "row", and cells differences from total columns with pct = "col".
- "first": calculate cells differences from the first cell of the row or column (useful to color temporal developments).
- "no": not use diffs to gain calculation time.

comp The comparison level : by subtables/groups, or for the whole table.

- "tab": by default, contributions to variance, row differences from totals/first cells, and row confidence intervals for these differences, are calculated for each tab_vars group.
- "all": compare cells to the general total line (provided there is a total table with a total row), or with the first line of the total table when diff = "first".

totaltab The total table, if there are subtables/groups (i.e. when tab_vars is provided) :

- "line": by default, add a general total line (necessary for calculations with comp = "all")
- "table": add a complete total table (i.e. row_var by col_vars without tab_vars).
- "no": not to draw any total table.

totaltab_name The name of the total table, as a single string.

tot The totals :

- c("col", "row") or "both" : by default, both total rows and total columns.
- "row": only total rows.
- "col": only total column.
- "no": remove all totals (after calculations if needed).

`total_names` The names of the totals, as a character vector of length one or two. Use syntax of type `c("Total row", "Total column")` to set different names for rows and cols.

`chi2` Set to `TRUE` to calculate Chi2 summaries with `tab_chi2`. Useful to print metadata, and to color cells based on their contribution to variance (`color = "contrib"`). Automatically added if needed for `color`.

`ci` The type of confidence intervals to calculate, passed to `tab_ci` (automatically added if needed for `color`).

- `"cell"`: absolute confidence intervals of cells percentages.
- `"diff"`: confidence intervals of the difference between a cell and the relative total cell (or relative first cell when `diff = "first"`).
- `"auto"`: `ci = "diff"` for means and row/col percentages, `ci = "cell"` for frequencies ("all", "all_tabs").

By default, for percentages, with `ci = "cell"` Wilson's method is used, and with `ci = "diff"` Wald's method along Agresti and Caffo's adjustment. Means use classic method. This can be changed in `tab_ci`.

`conf_level` The confidence level, as a single numeric between 0 and 1. Default to 0.95 (95%).

`subtext` A character vector to print rows of legend under the table.

`cleannames` Set to `TRUE` to clean levels names, by removing prefix numbers like "1-", and text in parenthesis. All data formatting arguments are passed to `tab_prepare`.

`rare_to_other` When set to `TRUE`, levels with less count than `n_min` will be merged into an "Other" level.

`n_min` The count under which a level is aggregated in the "Other" level.

`other_level` The name of the "Other" level, as a single string.

`filter` A `dplyr::filter` to apply to the data frame first, as a single string (which will be converted to code, i.e. to a call). Useful when printing multiples tabs with `tibble::tribble`, to use different filters for similar tables or simply make the field of observation more visible into the code.

Value

A `tibble` of class `tab`, possibly with colored reading helpers. All non-text columns are of class `fmt`, storing all the data necessary to print formats and colors. Columns with `row_var` and `tab_vars` are of class `factor` : every added `factor` will be considered as a `tab_vars` and used for grouping. To add text columns without using them in calculations, be sure they are of class `character`.

Examples

```
# A simple cross-table:
tab(forcats::gss_cat, marital, race)


# With more variables provided, `tab` makes a subtables for each combination of levels:

tab(forcats::gss_cat, marital, tab_vars = c(year, race))
```

```
# You can also add supplementary columns, text or numeric:

tab(dplyr::storms, category, status, sup_cols = c("pressure", "wind"))


# Colors to help the user read the table:
data <- forcats::gss_cat %>%
  dplyr::filter(year %in% c(2000, 2006, 2012), !marital %in% c("No answer", "Widowed"))
gss  <- "Source: General social survey 2000-2014"
gss2 <- "Source: General social survey 2000, 2006 and 2012"

# Differences between the cell and it's subtable's total cell:

tab(data, race, marital, year, subtext = gss2, pct = "row", color = "diff")


# Differences between the cell and the whole table's general total cell:

tab(data, race, marital, year, subtext = gss2, pct = "row", color = "diff",
  comp = "all")


# Historical differences:

data2 <- data %>% dplyr::mutate(year = as.factor(year))
tab(data2, year, marital, race, subtext = gss2, pct = "row",
    color = "diff", diff = "first", tot = "col")


# Differences with the total, except if their confidences intervals are superior to them:
tab(forcats::gss_cat, race, marital, subtext = gss, pct = "row", color = "diff_ci")

# Same differences, minus their confidence intervals:
tab(forcats::gss_cat, race, marital, subtext = gss, pct = "row", color = "after_ci")

# Contribution of cells to table's variance, like in a correspondence analysis:
tab(forcats::gss_cat, race, marital, subtext = gss, color = "contrib")


# Since the result is a tibble, you can use all dplyr verbs to modify it :

library(dplyr)
tab(dplyr::storms, category, status, sup_cols = c("pressure", "wind")) %>%
  dplyr::filter(category != "-1") %>%
  dplyr::select(-`tropical depression`) %>%
  dplyr::arrange(is_totrow(.), desc(category))



# With `dplyr::arrange`, don't forget to keep the order of tab variables and total rows:
tab(data, race, marital, year, pct = "row") %>%
```

```
  dplyr::arrange(year, is_totrow(.), desc(Married))
```

---

tab_chi2 *Add Chi2 summaries to a* tab

Description

Add Chi2 summaries to a tab

Usage

```
tab_chi2(
  tabs,
  calc = c("ctr", "p", "var", "counts"),
  comp = NULL,
  color = c("no", "auto", "all", "all_pct")
)
```

Arguments

tabs	A tibble of class tab, made with tab_plain or tab_many.
calc	By default all elements of the Chi2 summary are calculated : contributions to variance, pvalue, variance and unweighted count. You can choose which are computed by selecting elements in the vector c("ctr", "p", "var", "counts").
comp	Comparison level. When tab_vars are present, should the contributions to variance be calculated for each subtable/group (by default, comp = "tab") ? Should they be calculated for the whole table (comp = "all") ? comp must be set once and for all the first time you use tab_plain, tab_num or tab_chi2 with rows, or tab_ci.
color	The type of colors to print, as a single string. • "no": by default, no colors are printed • "all": color all cells based on their contribution to variance (except for mean columns, from numeric variables) • "all_pct": color all percentages cells based on their contribution to variance • "auto": only color columns with counts, pct = "all" or pct = "all_tabs"

Value

A tibble of class tab, with Chi2 summaries as metadata, possibly colored based on contributions of cells to variance.

tab_ci *Add confidence intervals to a* tab

Description

Add confidence intervals to a tab

Usage

```
tab_ci(
  tabs,
  ci = "auto",
  comp = NULL,
  conf_level = 0.95,
  color = "no",
  visible = FALSE,
  method_cell = "wilson",
  method_diff = "ac"
)
```

Arguments

tabs	A tibble of class tab made with tab_plain or tab_many.
ci	The type of ci to calculate. Set to "cell" to calculate absolute confidence intervals. Set to "diff" to calculate the confidence intervals of the difference between a cell and the relative total cell (or the reference cell, when diff is not "tot" in tab_plain or tab_num). By default, "diff" ci are calculated for means and row and col percentages, "cell" ci for frequencies ("all", "all_tabs").
comp	Comparison level. When tab_vars are present, should the contributions to variance be calculated for each subtable/group (by default, comp = "tab") ? Should they be calculated for the whole table (comp = "all") ? comp must be set once and for all the first time you use tab_plain, tab_num or tab_chi2 with rows, or tab_ci.
conf_level	The confidence level, as a single numeric between 0 and 1. Default to 0.95 (95%).
color	The type of colors to print, as a single string. • "no": by default, no colors are printed • "diff_ci": color pct and means based on cells differences from totals or first cells, removing coloring when the confidence interval of this difference is higher than the difference itself • "after_ci": idem, but cut off the confidence interval from the difference
visible	By default confidence intervals are calculated and used to set colors, but not printed. Set to TRUE to print them in the result.

`method_cell` Character string specifying which method to use with percentages for `ci = "cell"`. This can be one out of: "wald", "wilson", "wilsoncc", "agresti-coull", "jeffreys", "modified wilson", "modified jeffreys", "clopper-pearson", "arcsine", "logit", "witting", "pratt", "midp", "lik" and "blaker". Defaults to "wilson". See `BinomCI`.

`method_diff` Character string specifying which method to use with percentages for `ci = "diff"`. This can be one out of: "wald", "waldcc", "ac", "score", "scorecc", "mn", "mee", "blj", "ha", "hal", "jp". Defaults to "ac", Wald interval with the adjustment according to Agresti, Caffo for difference in proportions and independent samples. See `BinomDiffCI`.

Value

A `tibble` of class `tab`, colored based on differences (from totals/first cells) and confidence intervals.

Examples

```
# A typical workflow with tabxplor step-by-step functions :

data <- dplyr::starwars %>%
  tab_prepare(sex, hair_color, gender, rare_to_other = TRUE,
              n_min = 5, na_drop_all = sex)

data %>%
  tab_plain(sex, hair_color, gender, tot = c("row", "col"),
    pct = "row", comp = "all") %>%
    tab_ci("diff", color = "after_ci")
```

---

tab_kable *Print a tabxplor table in html*

Description

Print a tabxplor table in html

Usage

```
tab_kable(
  tabs,
  theme = c("light", "dark"),
  color_type = NULL,
  html_24_bit = NULL,
  tooltips = TRUE,
  popover = NULL,
  color_legend = TRUE,
  caption = NULL,
  ...
)
```

Arguments

tabs	A table made with tab or tab_many.
theme	By default, a white table with black text, Set to "dark" for a black table with white text.
color_type	Set to "text" to color the text, "bg" to color the background. By default it takes getOption("tabxplor.color_style_type").
html_24_bit	Should specific 24bits colors palettes be used ? Default to getOption("tabxplor.color_html_24_bit"
tooltips	By default, html tooltips are used to display additional informations at mouse hover. Set to FALSE to discard.
popover	By default, takes getOption("tabxplor.kable_popover"). When FALSE, html tooltips are of the base kind : they can't be used with floating table of content in **rmarkdown** documents. Set to TRUE to use **kableExtra** html popovers instead, which are compatible with floating toc. Remember to enable the popover module by copying the following code into your document : <script> $(document).ready(function() $('[data-toggle="popover"]').popover(); }); </script>
color_legend	Print colors legend below the table ? You can then use a css chunk in rmarkdown to change popovers colors.
caption	The table caption. For formatting, you need to use a css with caption{}in rmarkdown.
...	Other arguments to pass to kableExtra::kable_styling.

Value

A html table (opened in the viewer in RStudio). Differences from totals, confidence intervals, contribution to variance, and unweighted counts, are available in an html tooltip at cells hover.

Examples

```
tabs <- tab(forcats::gss_cat, race, marital, year, pct = "row", color = "diff")
tab_kable(tabs, theme = "light", color_type = "text")
```

---

tab_many *Many cross-tables as one, with color helpers*

Description

A full-featured function to create, manipulate and format many cross-tables as one, using colors to make the printed tab more easily readable (in R terminal or exported to Excel with tab_xl). Since objects of class tab are also of class tibble, you can then use all **dplyr** verbs to modify the result, like select, arrange, filter or mutate.

Only breaks for attractions/over-representations (in green) should be given, as a vector of positive doubles, with length between 1 and 5. Breaks for aversions/under-representations (in orange/red) will simply be the opposite.

Usage

```
tab_many(
  data,
  row_vars,
  col_vars,
  tab_vars,
  wt,
  levels = "all",
  na = "keep",
  na_drop_all,
  digits = 0,
  totaltab = "line",
  totaltab_name = "Ensemble",
  totrow = TRUE,
  totcol = "last",
  total_names = "Total",
  pct = "no",
  diff = "tot",
  comp = "tab",
  chi2 = FALSE,
  ci = "no",
  conf_level = 0.95,
  method_cell = "wilson",
  method_diff = "ac",
  color = "no",
  subtext = "",
  cleannames = NULL,
  rare_to_other = FALSE,
  n_min = 30,
  other_level = "Others",
  filter
)

tab_get_vars(tabs, vars = c("row_var", "col_vars", "tab_vars"))

is_tab(x)

set_color_style(
  type = c("text", "bg"),
  theme = NULL,
  html_24_bit = c("blue_red", "green_red", "no"),
  custom_palette = NULL
)

get_color_style(
  mode = c("crayon", "color_code"),
  type = NULL,
  theme = NULL,
```

```
  html_24_bit = NULL
)

set_color_breaks(pct_breaks, mean_breaks, contrib_breaks)

get_color_breaks(brk, type = c("positive", "all"))
```

Arguments

data	A data frame.
row_vars	The row variable, which will be printed with one level per line. If numeric, it will be converted to factor. If more than one row_var if provided, a different table is made for each of them.
col_vars	<tidy-select> One column is printed for each level of each column variable. For numeric variables means are calculated, in a single column. To pass many variables you may use syntax `col_vars = c(col_var1, col_var2, ...)`.
tab_vars	<tidy-select> One subtable is made for each combination of levels of the tab variables. To pass many variables you may use syntax `tab_vars = c(tab_var1, tab_var2, ...)`. All tab variables are converted to factor. Leave empty to make a simple table.
wt	A weight variable, of class numeric. Leave empty for unweighted results.
levels	The levels of `col_vars` to keep (for more complex selections use `dplyr::select`). The argument is vectorised over `col_vars`. • `"all"`: by default, all levels are kept. • `"first"`: only keep the first level of each `col_vars`
na	The policy to adopt with missing values. It must be a single string. • `na = "keep"`: by default, prints `NA`'s as explicit `"NA"` level. • `na = "drop"`: removes `NA` levels before making each table (tabs made with different column variables may have a different number of observations, and won't exactly have the same total columns).
na_drop_all	<tidy-select> Removes all observations with a `NA` in any of the chosen variables, for all tables (tabs for each column variable will have the same number of observations).
digits	The number of digits to print, as a single integer, or an integer vector the same length as `col_vars`. The argument is vectorisez over `col_vars`.
totaltab	The total table, if there are subtables/groups (i.e. when `tab_vars` is provided). Vectorised over `row_vars`. • `"line"`: by default, add a general total line (necessary for calculations with `comp = "all"`) • `"table"`: add a complete total table (i.e. `row_var` by `col_vars` without `tab_vars`). • `"no"`: not to draw any total table.
totaltab_name	The name of the total table, as a single string.

totrow — By default, total rows are printed. Set to FALSE to remove them (after calculations if needed). Vectorised over row_vars.

totcol — The policy with total columns. Vectorised over col_vars.

- "last": by default, only prints a total column for the last column variable (of class factor, not numeric).
- "each": print a total column for each column variable.
- "no": remove all total columns (after calculations if needed).

total_names — The names of the totals, as a character vector of length one or two. Use syntax of type c("Total row", "Total column") to set different names for rows and cols.

pct — The type of percentages to calculate :

- "row": row percentages.
- "col": column percentages.
- "all": frequencies for each subtable/group, if there is tab_vars.
- "all_tabs": frequencies for the whole (set of) table(s).

The argument is vectorised over both row_vars and col_vars. You can then write as the following : pct = list(row_var1 = list("row", "col", "col"), row_var2 = list("col", "row", "row"))

diff — The reference cell to calculate differences (used to print colors). Vectorisez over row_vars.

- "tot": by default, cells differences from total rows are calculated with pct = "row", and cells differences from total columns with pct = "col".
- "first": calculate cells differences from the first cell of the row or column (useful to color temporal developments).
- n: when diff is an integer, the nth row (or column) is used for comparison.
- "regex": when diff is a string, it it used as a regular expression, to match with the names of the rows (or columns). Be precise enough to match only one column or row, otherwise you get a warning message.
- "no": not use diffs.

comp — The comparison level : by subtables/groups, or for the whole table. Vectorised over row_vars.

- "tab": by default, contributions to variance, row differences from totals/first cells, and row confidence intervals for these differences, are calculated for each tab_vars group.
- "all": compare cells to the general total line (provided there is a total table with a total row), or with the reference line of the total table when diff = "first", an integer or a regular expression.

chi2 — Set to TRUE to calculate Chi2 summaries with tab_chi2. Useful to print metadata, and to color cells based on their contribution to variance (color = "contrib"). Vectorised over row_vars.

ci — The type of confidence intervals to calculate, passed to tab_ci. Vectorised over row_vars.

- "cell": absolute confidence intervals of cells percentages.

- "diff": confidence intervals of the difference between a cell and the relative total cell (or relative first cell when diff = "first").
- "auto": ci = "diff" for means and row/col percentages, ci = "cell" for frequencies ("all", "all_tabs").

By default, for percentages, with ci = "cell" Wilson's method is used, and with ci = "diff" Wald's method along Agresti and Caffo's adjustment. Means use classic method. This can be changed in tab_ci.

conf_level — The confidence level, as a single numeric between 0 and 1. Default to 0.95 (95%).

method_cell — Character string specifying which method to use with percentages for ci = "cell". This can be one out of: "wald", "wilson", "wilsoncc", "agresti-coull", "jeffreys", "modified wilson", "modified jeffreys", "clopper-pearson", "arcsine", "logit", "witting", "pratt", "midp", "lik" and "blaker". Defaults to "wilson". See BinomCI.

method_diff — Character string specifying which method to use with percentages for ci = "diff". This can be one out of: "wald", "waldcc", "ac", "score", "scorecc", "mn", "mee", "blj", "ha", "hal", "jp". Defaults to "ac", Wald interval with the adjustment according to Agresti, Caffo for difference in proportions and independent samples. See BinomDiffCI.

color — The type of colors to print, as a single string. Vectorised over row_vars.

- "no": by default, no colors are printed.
- "diff": color percentages and means based on cells differences from totals (or from first cells when diff = "first").
- "diff_ci": color pct and means based on cells differences from totals or first cells, removing coloring when the confidence interval of this difference is higher than the difference itself.
- "after_ci": idem, but cut off the confidence interval from the difference first.
- "contrib": color cells based on their contribution to variance (except mean columns, from numeric variables).
- "auto": frequencies (pct = "all", pct = "all_tabs") and counts are colored with "contrib". When ci = "diff", row and col percentages are colored with "after_ci" ; otherwise they are colored with "diff".

subtext — A character vector to print rows of legend under the table.

cleannames — Set to TRUE to clean levels names, by removing prefix numbers like "1-", and text in parenthesis. All data formatting arguments are passed to tab_prepare.

rare_to_other — When set to TRUE, levels with less count than n_min will be merged into an "Other" level.

n_min — The count under which a level is aggregated in the "Other" level.

other_level — The name of the "Other" level, as a single string.

filter — A dplyr::filter to apply to the data frame first, as a single string (which will be converted to code, i.e. to a call). Useful when printing multiples tabs with tibble::tribble, to use different filters for similar tables or simply make the field of observation more visible into the code.

tabs — A tibble of class tab, made with tab, tab_many or tab_plain.

vars	In tab_get_vars, a character vector containing the wanted vars names: "row_var", "col_vars" or "tab_vars".
x	A object to test with is_tab.
type	Default to "positive", which just print breaks for positive spreads. Set to all to get breaks for negative spreads as well.
theme	For set_color_style and get_color_style, is your console or html table background "light" or "dark" ? Default to RStudio theme.
html_24_bit	Should specific 24bits colors palettes be used for html tables ? With light themes only. Default to getOption("tabxplor.color_html_24_bit")
custom_palette	Possibility to provide a custom color styles, as a character vector of 10 html color codes (the five first for over-represented numbers, the five last for under-represented ones). The result is saved to options("tabxplor.color_style"). To discard, relaunch the function with custom_palette = NULL.
mode	By default, get_color_style returns a list of **crayon** coloring functions. Set to "color_code" to return html color codes.
pct_breaks	If they are to be changed, the breaks used for percentages. Default to c(0.05, 0.1, 0.2, 0.3) : first color used when the pct of a cell is +5% superior to the pct of the related total ; second color used when it is +10% superior ; third +20% superior ; fourth +30% superior. The opposite for cells inferior to the total. With color = "after_ci", the first break is subtracted from all breaks (default becomes c(0, 0.05, 0.15, 0.25) : +0%, +5%, +15%, +25%).
mean_breaks	If they are to be changed, the breaks used for means. Default to c(1.15, 1.5, 2, 4) : first color used when the mean of a cell is superior to 1.15 times the mean of the related total row ; second color used when it is superior to 1.5 times ; etc. The opposite for cells inferior to the total. With color = "after_ci", the first break is divided from all breaks (default becomes c(1, 1.3, 1.7, 3.5)).
contrib_breaks	If they are to be changed, the breaks used for contributions to variance. Default to c(1, 2, 5, 10) : first color used when the contribution of a cell is superior to the mean contribution ; second color used when it is superior to 2 times the mean contribution ; etc. The global color (for example green or red/orange) is given by the sign of the spread.
brk	When missing, return all color breaks. Specify to return a given color break, among "pct", "mean", "contrib", "pct_ci" and "mean_ci".

Value

A tibble of class tab, possibly with colored reading helpers. When there are two row_vars or more, a list of tibble of class tab. All non-text columns are of class fmt, storing all the data necessary to print formats and colors. Columns with row_var and tab_vars are of class factor : every added factor will be considered as a tab_vars and used for grouping. To add text columns without using them in calculations, be sure they are of class character.

A list with the variables names.

A single logical.

Set global options "tabxplor.color_style_type" and "tabxplor.color_style_theme", used when printing tab objects.

A vector of crayon color functions, or a vector of color html codes.

Set the global option "tabxplor.color_breaks" as a list different double vectors, and also returns it invisibly.

The color breaks as a double vector, or list of double vectors.

Functions

- `tab_get_vars()`: Get the variables names of a **tabxplor** `tab`
- `is_tab()`: a test function for class tabxplor_tab
- `set_color_style()`: define the color style used to print `tab`.
- `get_color_style()`: get color styles as **crayon** functions or html codes.
- `set_color_breaks()`: set the breaks used to print colors
- `get_color_breaks()`: get the breaks currently used to print colors

Examples

```
# Make a summary table with many col_vars, showing only one specific level :

library(dplyr)
first_lvs <- c("Married", "$25000 or more", "Strong republican", "Protestant")
data <- forcats::gss_cat %>% mutate(across(
  where(is.factor),
  ~ forcats::fct_relevel(., first_lvs[first_lvs %in% levels(.)])
))
tab_many(data, race, c(marital, rincome, partyid, relig, age, tvhours),
         levels = "first", pct = "row", chi2 = TRUE, color = "auto")


# Can be used with map and tribble to program several tables with different parameters
#  all at once, in a readable way:

library(purrr)
library(tibble)
pmap(
  tribble(
    ~row_var, ~col_vars       , ~pct , ~filter              , ~subtext               ,
    "race"  , "marital"       , "row", NULL                 , "Source: GSS 2000-2014",
    "relig" , c("race", "age"), "row", "year %in% 2000:2010", "Source: GSS 2000-2010",
    NA_character_, "race"     , "no" , NULL                 , "Source: GSS 2000-2014",
  ),
  .f = tab_many,
  data = forcats::gss_cat, color = "auto", chi2 = TRUE)

set_color_style(type = "bg")
set_color_breaks(
  pct_breaks = c(0.05, 0.15, 0.3),
  mean_breaks = c(1.15, 2, 4),
  contrib_breaks = c(1, 2, 5)
)
```

tab_num *Means table*

Description

Cross categorical variables with numeric variables, and get a table of means and standard deviations.

Usage

```
tab_num(
  data,
  row_var,
  col_vars,
  tab_vars,
  wt,
  diff = "tot",
  ci = NULL,
  conf_level = 0.95,
  comp = c("tab", "all"),
  color = c("auto", "diff", "diff_ci", "after_ci"),
  digits = 0,
  na = c("keep", "drop", "drop_fct", "drop_num"),
  totaltab = "line",
  totaltab_name = "Ensemble",
  tot = NULL,
  total_names = "Total",
  subtext = "",
  num = FALSE,
  df = FALSE
)
```

Arguments

data	A data frame.
row_var	The row variable, which will be printed with one level per line. If numeric, it will be used as a factor.
col_vars	The numeric variables, which will appear in columns : means and standard deviation are calculated for each levels of row_var and tab_vars.
tab_vars	<tidy-select> Tab variables : a subtable is made for each combination of levels of the selected variables. Leave empty to make a simple cross-table. All tab variables are converted to factor.
wt	A weight variable, of class numeric. Leave empty for unweighted results.
diff	The reference cell to calculate differences (used to print colors) :

- "tot": by default, cells differences from total rows are calculated with pct = "row", and cells differences from total columns with pct = "col".

- "first": calculate cells differences from the first cell of the row or column (useful to color temporal developments).
- "no": not use diffs to gain calculation time.

ci — The type of confidence intervals to calculate, passed to tab_ci (automatically added if needed for color).

- "cell": absolute confidence intervals of cells percentages.
- "diff": confidence intervals of the difference between a cell and the relative total cell (or relative first cell when diff = "first").
- "auto": ci = "diff" for means and row/col percentages, ci = "cell" for frequencies ("all", "all_tabs").

By default, for percentages, with ci = "cell" Wilson's method is used, and with ci = "diff" Wald's method along Agresti and Caffo's adjustment. Means use classic method. This can be changed in tab_ci.

conf_level — The confidence level for the confidence intervals, as a single numeric between 0 and 1. Default to 0.95 (95%).

comp — Comparison level. When tab_vars are present, should the contributions to variance be calculated for each subtable/group (by default, comp = "tab") ? Should they be calculated for the whole table (comp = "all") ? comp must be set once and for all the first time you use tab_plain, tab_num or tab_chi2 with rows, or tab_ci.

color — TRUE print the color percentages and means based on cells differences from totals or reference cell, as provided by diff. Default to FALSE, no colors.

digits — The number of digits to print, as a single integer.

na — The policy to adopt for missing values in row and tab variables (factors), as a single string.

- "keep": by default, NA's of row and tab variables are printed as an explicit "NA" level.
- "drop": remove NA's in row and tab variables.

NAs in numeric variables are always removed when calculating means. For that reason the n field of each resulting fmt column, used to calculate confidence intervals, only takes into account the complete observations (without NA). To drop all rows with NA in any numeric variable first, use tab_prepare or tab_many with the na_drop_all argument.

totaltab — The total table, if there are subtables/groups (i.e. when tab_vars is provided) :

- "line": by default, add a general total line (necessary for calculations with comp = "all")
- "table": add a complete total table (i.e. row_var by col_vars without tab_vars).
- "no": not to draw any total table.

totaltab_name — The name of the total table, as a single string.

tot — The totals :

- c("col", "row") or "both" : by default, both total rows and total columns.
- "row": only total rows.

- "col": only total column.
- "no": remove all totals (after calculations if needed).

total_names	The names of the totals, as a character vector of length one or two. Use syntax of type c("Total row", "Total column") to set different names for rows and cols.
subtext	A character vector to print rows of legend under the table.
num	Set to TRUE to obtain a table with normal numeric vectors (not fmt).
df	Set to TRUE to obtain a plain data.frame (not a tibble), with normal numeric vectors (not fmt). Useful, for example, to pass the table to correspondence analysis with **FactoMineR**.

Value

A tibble of class tabxplor_tab. If ... (tab_vars) are provided, a tab of class tabxplor_grouped_tab. All non-text columns are fmt vectors of class tabxplor_fmt, storing all the data necessary to print formats and colors. Columns with row_var and tab_vars are of class factor : every added factor will be considered as a tab_vars and used for grouping. To add text columns without using them in calculations, be sure they are of class character.

Examples

```
data <- dplyr::storms %>% tab_prepare(category, wind, na_drop_all = wind)
tab_num(data, category, wind, tot = "row", color = "after_ci")
```

---

tab_pct *Add percentages and diffs to a tab*

Description

Add percentages and diffs to a tab

Usage

```
tab_pct(
  tabs,
  pct = "row",
  digits = NULL,
  diff = c("tot", "first", "no"),
  comp = NULL,
  color = FALSE,
  just_diff = FALSE
)
```

Arguments

tabs	A tibble of class tab made with tab_plain or tab_many.
pct	The type of percentages to calculate. "row" draw row percentages. Set to "col" for column percentages. Set to "all" for frequencies (based on each subtable/group if tab_vars is provided). Set to "all_tabs" to calculate frequencies based on the whole (set of) table(s).
digits	The number of digits to print for percentages. As a single integer, or an integer vector the same length than col_vars.
diff	By default, with pct = "row", differences from total rows are calculated, and with pct = "col" differences from total columns. Set to diff = "first" to calculate differences with the first cell of the row/col (useful to color temporal developments). When not using diffs for colors, set to diff = "no" to gain calculation time. Diffs are also calculated for mean columns (made from numeric variables).
comp	Comparison level. When tab_vars are present, should the row differences be calculated for each subtable/group (by default comp = "tab" : comparison of each cell to the relative total row) ? Should they be calculated for the whole table (comp = "all" : comparison of each cell to the total row of the total table) ? When comp = "all" and diff = "first", cells are compared to the first cell of the total table instead. This parameter doesn't affect column percentages. comp must be set once and for all the first time you use tab_chi2, tab_pct with rows, or tab_ci.
color	Set to TRUE to color the resulting tab based on differences (from totals or from the first cell).
just_diff	If percentages are already calculated and you just want to recalculate differences.

Value

A tibble of class tab, with percentages displayed, possibly colored based on differences from totals or first cell.

---

tab_plain *Plain single cross-table*

Description

Plain single cross-table

Usage

```
tab_plain(
  data,
  row_var,
  col_var,
  tab_vars,
```

```
  wt,
  pct = "no",
  diff = "tot",
  comp = "tab",
  color = TRUE,
  digits = 0,
  na = "keep",
  subtext = "",
  totaltab = "line",
  totaltab_name = "Ensemble",
  tot = NULL,
  total_names = "Total",
  num = FALSE,
  df = FALSE
)
```

Arguments

`data`	A data frame.
`row_var, col_var`	The row variable, which will be printed with one level per line, and the column variable, which will be printed with one level per column. Numeric variables will be used as factors. To calculate means, use `tab_num`.
`tab_vars`	<tidy-select> Tab variables : a subtable is made for each combination of levels of the selected variables. Leave empty to make a simple cross-table. All tab variables are converted to factor.
`wt`	A weight variable, of class numeric. Leave empty for unweighted results.
`pct`	The type of percentages to calculate : • `"row"`: row percentages. • `"col"`: column percentages. • `"all"`: frequencies for each subtable/group, if there is `tab_vars`. • `"all_tabs"`: frequencies for the whole (set of) table(s).
`diff`	The reference cell to calculate differences (used to print `colors`) : • `"tot"`: by default, cells differences from total rows are calculated with `pct = "row"`, and cells differences from total columns with `pct = "col"`. • `"first"`: calculate cells differences from the first cell of the row or column (useful to color temporal developments). • `"no"`: not use diffs to gain calculation time.
`comp`	Comparison level. When `tab_vars` are present, should the contributions to variance be calculated for each subtable/group (by default, `comp = "tab"`) ? Should they be calculated for the whole table (`comp = "all"`) ? `comp` must be set once and for all the first time you use `tab_plain`, `tab_num` or `tab_chi2` with rows, or `tab_ci`.
`color`	`TRUE` print the color percentages and means based on cells differences from totals or reference cell, as provided by `diff`. Default to `FALSE`, no colors.

digits — The number of digits to print, as a single integer.

na — The policy to adopt with missing values, as a single string.

- "keep": by default, NA's of row, col and tab variables are printed as explicit "NA" level.
- "drop": removes NA of row, col and tab variables.

subtext — A character vector to print rows of legend under the table.

totaltab — The total table, if there are subtables/groups (i.e. when tab_vars is provided) :

- "line": by default, add a general total line (necessary for calculations with comp = "all")
- "table": add a complete total table (i.e. row_var by col_vars without tab_vars).
- "no": not to draw any total table.

totaltab_name — The name of the total table, as a single string.

tot — The totals :

- c("col", "row") or "both" : by default, both total rows and total columns.
- "row": only total rows.
- "col": only total column.
- "no": remove all totals (after calculations if needed).

total_names — The names of the totals, as a character vector of length one or two. Use syntax of type c("Total row", "Total column") to set different names for rows and cols.

num — Set to TRUE to obtain a table with normal numeric vectors (not fmt).

df — Set to TRUE to obtain a plain data.frame (not a tibble), with normal numeric vectors (not fmt). Useful, for example, to pass the table to correspondence analysis with **FactoMineR**.

Value

A tibble of class tabxplor_tab. If ... (tab_vars) are provided, a tab of class tabxplor_grouped_tab. All non-text columns are fmt vectors of class tabxplor_fmt, storing all the data necessary to print formats and colors. Columns with row_var and tab_vars are of class factor : every added factor will be considered as a tab_vars and used for grouping. To add text columns without using them in calculations, be sure they are of class character.

Examples

```
# A typical workflow with tabxplor step-by-step functions :

data <- dplyr::starwars %>% tab_prepare(sex, hair_color)

data %>%
  tab_plain(sex, hair_color, tot = c("row", "col"), pct = "row") %>%
  tab_chi2() %>%
  tab_ci(color = "after_ci")
```

tab_prepare *Prepare data for* tab_plain.

Description

Prepare data for tab_plain.

Usage

```
tab_prepare(
  data,
  ...,
  na_drop_all,
  cleannames = NULL,
  rare_to_other = FALSE,
  n_min = 30,
  other_level = "Others"
)
```

Arguments

data	A dataframe.
...	Variables then to be passed in tab_plain.
na_drop_all	<tidy-select> Removes all observation with a NA in any of the chosen variables.
cleannames	Set to TRUE to clean levels names, by removing prefix numbers like "1-", and text in parentheses.
rare_to_other	When set to TRUE, levels with less count than n_min will be merged into an "Other" level.
n_min	The count under which a level is aggregated in the "Other" level.
other_level	The name of the "Other" level, as a character vector of length one.

Value

A modified data.frame.

Examples

```
data <- dplyr::starwars %>%
tab_prepare(sex, hair_color, gender, rare_to_other = TRUE,
            n_min = 5, na_drop_all = sex)
data
```

tab_spread *Spread a tab, passing a tab variable to column*

Description

Spread a tab, passing a tab variable to column

Usage

```
tab_spread(
  tabs,
  spread_vars,
  names_prefix,
  names_sort = FALSE,
  totname = "Total"
)
```

Arguments

tabs	A tibble of class tab, made with tab, tab_many or tab_plain.
spread_vars	<tidy-select> The tab variables to pass to column, with a syntax of type c(var1, var2, ...).
names_prefix	String added to the start of every variable name.
names_sort	If no names_prefix is given, new names takes the form spread_var_col_var_level. Should then the column names be sorted ? If FALSE, the default, column names are ordered by first appearance.
totname	The new name of the total rows, as a single string.

Value

A tibble of class tab, with less rows and more columns.

Examples

```
 data <- forcats::gss_cat %>% dplyr::filter(year %in% c(2000, 2014))

tabs <-
  tab(data, relig, marital, c(year, race), pct = "row", totaltab = "no",
      color = "diff", tot = "row", rare_to_other = TRUE)

tabs %>%
  dplyr::select(year, race, relig, Married) %>%
  tab_spread(race)
```

tab_tot *Add totals to a* tab

Description

Add totals to a tab

Usage

```
tab_tot(
  tabs,
  tot = c("row", "col"),
  name = "Total",
  totcol = "last",
  data = NULL
)
```

Arguments

tabs	A tibble of class tab, made with tab_plain or tab_many.
tot	c("col", "row") and "both" print total rows and total columns. Set to "row" or "col" to print only one type. Set to "no" to remove all totals.
name	The names of the totals, as a character vector of length one or two. Use c("Total_row", "Total_column") to set different names for rows and cols.
totcol	"last" only prints a total column for the last factor column variable. Set to "each" to print a total column for each column variable.
data	The original database used to calculate the tab : it is only useful for mean columns (of numeric variables), in order to calculate the variances of total rows, necessary to calculate confidence intervals with tab_ci.

Value

A tibble of class tab. Total rows can then be detected using is_totrow, and total columns using is_totcol.

Examples

```
data <- dplyr::starwars %>% tab_prepare(sex, hair_color)

data %>%
  tab_plain(sex, hair_color) %>%
  tab_tot("col", totcol = "each")
```

tab_totaltab *Add total table to a* tab

Description

Add total table to a tab

Usage

```
tab_totaltab(
  tabs,
  totaltab = c("table", "line", "no"),
  name = "Ensemble",
  data = NULL
)
```

Arguments

tabs	A tibble of class tab, made with tab_plain or tab_many.
totaltab	If there are subtables, corresponding to the levels of tab_vars, totaltab = "table" add a complete total table. totaltab = "line" add a total table of only one row with the general total. totaltab = "no" remove any existing total table.
name	The name of the total table, as a single string.
data	The original database used to calculate the tab : it is only useful for mean columns (of numeric variables), in order to calculate the variances necessary to calculate confidence intervals with tab_ci.

Value

A tibble of class tab. Rows belonging to the total table can then be detected using is_tottab.

Examples

```
 data <- dplyr::starwars %>%
tab_prepare(sex, hair_color, gender, rare_to_other = TRUE,
            n_min = 5, na_drop_all = sex)

data %>%
  tab_plain(sex, hair_color, gender) %>%
  tab_totaltab("line")
```

tab_xl *Excel output for tabxplor tables, with formatting and colors*

Description

To modify the colors used into the Excel table, you can change the global options with set_color_style and set_color_breaks.

Usage

```
tab_xl(
  tabs,
  path = NULL,
  replace = FALSE,
  open = rlang::is_interactive(),
  colnames_rotation = 0,
  remove_tab_vars = TRUE,
  colwidth = "auto",
  print_ci = FALSE,
  print_color_legend = TRUE,
  sheets = "tabs",
  n_min = 0,
  titles,
  hide_near_zero = "auto",
  color_type = "text"
)
```

Arguments

tabs	A table made with tab, tab_many or tab_plain, or a list of such tables.
path, replace, open	The name, and possibly the path, of the Excel file to create (possibly without the .xlsx extension). Default path to temporary directory. Set global option "tabxplor.export_dir" with link[base:options]{options} to change default directory. By default replace is TRUE when path is provided, FALSE when path is not provided. Use replace = TRUE to overwrite existing files. Use open = FALSE if you don't want to automatically open the tables in Excel (or another software associated with .xlsx files).
colnames_rotation	Rotate the names of columns to an angle (in degrees).
remove_tab_vars	By default, tab_vars columns are removed to gain space. Set to FALSE to keep them.
colwidth	The standard width for numeric columns, as a number. Default to "auto".
print_ci	Set to TRUE to print confidence intervals in another table, at the left of the base table.

print_color_legend
: Should the color legends be printed with the subtexts ?

sheets
: The Excel sheets options :
  - "tabs": a new sheet is created for each table
  - "unique": all tables are on the same sheet
  - "auto": subsequent tables with the same columns are printed on the same sheets

n_min
: The total count under which a column or row is turned pale grey because there is not enough observation for it to be significant. Default to 0 (not used).

titles
: The titles of the different tables, as a character vector. When missing titles are given based on the names of the variables.

hide_near_zero
: By default all cells displayed as 0 (even rounded) turn pale grey, to make the distribution of empty cells (and other cells) more visible. Provide a number to turn grey every cell below it. Set to Inf not to use this feature.

color_type
: By default, the text is colored. Set to "bg" to color the background instead.

Value

The table(s) with formatting and colors in an Excel file, as a side effect. Invisibly returns tabs.

Examples

```
forcats::gss_cat %>%
  tab(marital, race, pct = "row", color = "diff") %>%
  tab_xl()
```

---

tab_xl_confidential *Excel output for tabxplor tables with confidentiality rules.*

Description

Excel output for tabxplor tables, with colors to show if counts and percentages respect statistical confidentiality rules. Don't forget to provide subtext = c("Source : description of the source of the data") in tab or tab_many, otherwise it is not possible to assess, for your reader, which confidentiality rules applies. For the same reason, you must supply a description of all variables in var_labels.

Usage

```
tab_xl_confidential(
  tabs,
  path = NULL,
  replace = NULL,
  open = rlang::is_interactive(),
```

```
  n_min = 5,
  pct_max = 0.95,
  recalculate_totcols = NULL,
  var_labels = character(),
  colnames_rotation = 0,
  colwidth = 10,
  sheets = "unique",
  print_color_legend = TRUE,
  titles,
  hide_near_zero = "auto",
  color_type = "text"
)
```

Arguments

tabs	A table made with tab, tab_many or tab_plain, or a list of such tables.
path, replace, open	The name, and possibly the path, of the Excel file to create (possibly without the .xlsx extension). Default path to temporary directory. Set global option "tabxplor.export_dir" with link[base:options]{options} to change default directory. By default replace is TRUE when path is provided, FALSE when path is not provided. Use replace = TRUE to overwrite existing files. Use open = FALSE if you don't want to automatically open the tables in Excel (or another software associated with .xlsx files).
n_min	The total count under which a column or row doesnt respect statistical confidentiality. Default to 5.
pct_max	The row or column percentage above which, knowing the column category, it becomes possible to guess the row category, or the other way round. Default to 0.95 (95%).
recalculate_totcols	By default, total columns are recalculated from counts if there are many col_vars but only one total column. Provide a logical vector the length of the number of tables, or a single logical, to choose the wanted behavior. The fastest way to do it is to use tab_many() with totcol = "each" before.
var_labels	The description of all the variables, necessary to assess that the tables don't break confidentiality rules, as a character vector of the type c('variable1' = 'description of the variable', 'variable2' = ...)
colnames_rotation	Rotate the names of columns to an angle (in degrees).
colwidth	The standard width for numeric columns, as a number. Default to 10.
sheets	The Excel sheets options : • "unique": all tables are on the same sheet • "tabs": a new sheet is created for each table • "auto": subsequent tables with the same columns are printed on the same sheets
print_color_legend	Should the color legends be printed with the subtexts ?

titles	The titles of the different tables, as a character vector. When missing titles are given based on the names of the variables.
hide_near_zero	By default all cells displayed as 0 (even rounded) turn pale grey, to make the distribution of empty cells (and other cells) more visible. Provide a number to turn grey every cell below it. Set to Inf not to use this feature.
color_type	By default, the text is colored. Set to "bg" to color the background instead.

Value

The table(s) with formatting and colors in an Excel file, as a side effect. Invisibly returns tabs.

Examples

```
forcats::gss_cat |>
  tab(race, marital, year, pct = "row", color = "diff",
     subtext = c('Source : National Opinion Research Center, General Social Survey.')) |>
  tab_xl_confidential(titles = "Marital status by race",
     var_labels = c("marital" = "marital status", "race" = "race",
     "year" = "year of survey"))
```

---

tbl_format_body.tabxplor_tab

Table body for class tab

Description

Table body for class tab

Usage

```
## S3 method for class 'tabxplor_tab'
tbl_format_body(x, setup, ...)
```

Arguments

x	An object of class tabxplor_tab
setup	A setup object from the table
...	Other parameters.

Value

A character vector.

tbl_format_footer.tabxplor_tab

Table footer for class tab

Description

Table footer for class tab

Usage

```
## S3 method for class 'tabxplor_tab'
tbl_format_footer(x, setup, ...)
```

Arguments

x	An object of class tabxplor_tab
setup	A setup object from the table
...	Other parameters.

Value

A character vector.

tbl_sum.tabxplor_grouped_tab

Table headers for class grouped tab

Description

Table headers for class grouped tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
tbl_sum(x, ...)
```

Arguments

x	An object of class tabxplor_tab
...	Other parameters.

Value

A table header

tbl_sum.tabxplor_tab *Table headers for class tab*

Description

Table headers for class tab

Usage

```
## S3 method for class 'tabxplor_tab'
tbl_sum(x, ...)
```

Arguments

x	An object of class tabxplor_tab
...	Other parameters.

Value

A table header

ungroup.tabxplor_grouped_tab *ungroup method for class tabxplor_grouped_tab*

Description

ungroup method for class tabxplor_grouped_tab

Usage

```
## S3 method for class 'tabxplor_grouped_tab'
ungroup(x, ...)
```

Arguments

x	A tibble of class tabxplor_grouped_tab.
...	Variables to remove from the grouping.

Value

An object of class tabxplor_tab or tabxplor_grouped_tab.

`vec_arith.tabxplor_fmt` *Vec_arith method for fmt*

Description

Vec_arith method for fmt

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_arith(op, x, y, ...)

## Default S3 method:
vec_arith.tabxplor_fmt(op, x, y, ...)

## S3 method for class 'tabxplor_fmt'
vec_arith.tabxplor_fmt(op, x, y, ...)

## S3 method for class 'numeric'
vec_arith.tabxplor_fmt(op, x, y, ...)

## S3 method for class 'tabxplor_fmt'
vec_arith.numeric(op, x, y, ...)

## S3 method for class 'MISSING'
vec_arith.tabxplor_fmt(op, x, y, ...)
```

Arguments

op	Operation to do.
x	fmt object.
y	Second object.
...	Other parameter.

Value

A fmt vector

A fmt vector

A fmt vector

A fmt vector

A fmt vector

A fmt vector

Methods (by class)

- `vec_arith.tabxplor_fmt(default)`: default vec_arith method for fmt
- `vec_arith.tabxplor_fmt(tabxplor_fmt)`: vec_arith method for fmt + fmt
- `vec_arith.tabxplor_fmt(numeric)`: vec_arith method for fmt + numeric
- `vec_arith.tabxplor_fmt(MISSING)`: vec_arith method for -fmt

Functions

- `vec_arith.numeric(tabxplor_fmt)`: vec_arith method for numeric + fmt

`vec_cast.character.tabxplor_fmt` *Convert fmt into character*

Description

Convert fmt into character

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_cast.character(x, to, ...)
```

Arguments

`x`	A fmt vector
`to`	A character vector
`...`	Other parameter

Value

A character vector

`vec_cast.double.tabxplor_fmt` *Convert fmt into double*

Description

Convert fmt into double

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_cast.double(x, to, ...)
```

Arguments

`x`	A fmt vector
`to`	A double vector
`...`	Other parameter.

Value

A double vector

`vec_cast.integer.tabxplor_fmt` *Convert fmt into integer*

Description

Convert fmt into integer

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_cast.integer(x, to, ...)
```

Arguments

`x`	A integer vector
`to`	A fmt vector
`...`	Other parameter.

Value

An integer vector

vec_cast.tabxplor_fmt.double

Convert double into fmt

Description

Convert double into fmt

Usage

```
## S3 method for class 'tabxplor_fmt.double'
vec_cast(x, to, ...)
```

Arguments

x	A double vector
to	A fmt vector
...	Other parameter.

Value

A fmt vector

vec_cast.tabxplor_fmt.integer

Convert integer into fmt

Description

Convert integer into fmt

Usage

```
## S3 method for class 'tabxplor_fmt.integer'
vec_cast(x, to, ...)
```

Arguments

x	A integer vector
to	A fmt vector
...	Other parameter.

Value

A fmt vector

vec_cast.tabxplor_fmt.tabxplor_fmt *Convert fmt into fmt*

Description

Convert fmt into fmt

Usage

```
## S3 method for class 'tabxplor_fmt.tabxplor_fmt'
vec_cast(x, to, ...)
```

Arguments

x	A fmt vector
to	A fmt vector
...	Other parameter.

Value

A fmt vector

vec_math.tabxplor_fmt *Vec_math method for class fmt*

Description

Vec_math method for class fmt

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_math(.fn, .x, ...)
```

Arguments

.fn	A function
.x	A fmt object
...	Other parameter

Value

A fmt vector

vec_proxy_compare.tabxplor_fmt *Compare with fmt vector*

Description

Compare with fmt vector

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_proxy_compare(x, ...)
```

Arguments

x	A fmt vector
...	Other parameter

Value

A double vector

vec_proxy_equal.tabxplor_fmt *Test equality with fmt vector*

Description

Test equality with fmt vector

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_proxy_equal(x, ...)
```

Arguments

x	A fmt vector
...	Other parameter

Value

A double vector

`vec_ptype2.double.tabxplor_fmt`

Find common ptype between double and fmt

Description

Find common ptype between double and fmt

Usage

```
## S3 method for class 'double.tabxplor_fmt'
vec_ptype2(x, y, ...)
```

Arguments

`x`	A double vector
`y`	A fmt vector
`...`	Other parameter.

Value

A fmt vector

`vec_ptype2.integer.tabxplor_fmt`

Find common ptype between integer and fmt

Description

Find common ptype between integer and fmt

Usage

```
## S3 method for class 'integer.tabxplor_fmt'
vec_ptype2(x, y, ...)
```

Arguments

`x`	An integer vector
`y`	A fmt vector
`...`	Other parameter.

Value

A fmt vector

vec_ptype2.tabxplor_fmt.double

Find common ptype between fmt and double

Description

Find common ptype between fmt and double

Usage

```
## S3 method for class 'tabxplor_fmt.double'
vec_ptype2(x, y, ...)
```

Arguments

x	A fmt vector
y	A double vector
...	Other parameter.

Value

A fmt vector

vec_ptype2.tabxplor_fmt.integer

Find common ptype between fmt and integer

Description

Find common ptype between fmt and integer

Usage

```
## S3 method for class 'tabxplor_fmt.integer'
vec_ptype2(x, y, ...)
```

Arguments

x	A fmt vector
y	An integer vector
...	Other parameter.

Value

A fmt vector

vec_ptype2.tabxplor_fmt.tabxplor_fmt

Find common ptype between fmt and fmt

Description

Find common ptype between fmt and fmt

Usage

```
## S3 method for class 'tabxplor_fmt.tabxplor_fmt'
vec_ptype2(x, y, ...)
```

Arguments

x	A fmt object.
y	A fmt object.
...	Other parameter.

Value

A fmt vector

vec_ptype_abbr.tabxplor_fmt

Abbreviated display name for class fmt in tibbles

Description

Abbreviated display name for class fmt in tibbles

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_ptype_abbr(x, ...)
```

Arguments

x	A fmt object.
...	Other parameter.

Value

A single string with abbreviated fmt type.

vec_ptype_full.tabxplor_fmt *Printed type for class fmt*

Description

Printed type for class fmt

Usage

```
## S3 method for class 'tabxplor_fmt'
vec_ptype_full(x, ...)
```

Arguments

x	A fmt object.
...	Other parameter.

Value

A single string with full fmt type.

[.tabxplor_grouped_tab *subset method for class tabxplor_grouped_tab*

Description

subset method for class tabxplor_grouped_tab

Usage

```
"x[i]  ;  x[i, j, ... , drop = TRUE]"
```

Arguments

x	A tabxplor_grouped_tab object.
i, j, ...	Indices
drop	For matrices and arrays. If TRUE the result is coerced to the lowest possible dimension (see the examples). This only works for extracting elements, not for the replacement.

Value

An object of class tabxplor_grouped_tab.

`[<-.tabxplor_grouped_tab`

set subset method for class tabxplor_grouped_tab

Description

set subset method for class tabxplor_grouped_tab

Usage

```
"x[i] <- value  ;  x[i, j, ...] <- value"
```

Arguments

`x`	A tabxplor_grouped_tab object.
`i, j, ...`	Indices.
`value`	The new value.

Value

An object of class `tabxplor_grouped_tab`.

`[[<-.tabxplor_grouped_tab`

set sub-subset method for class tabxplor_grouped_tab

Description

set sub-subset method for class tabxplor_grouped_tab

Usage

```
"x[[...]] <- value"
```

Arguments

`x`	A tabxplor_grouped_tab object.
`...`	Indices
`value`	The new value.

Value

An object of class `tabxplor_grouped_tab`.

`$.tabxplor_fmt` *$ method for class tabxplor_fmt*

Description

$ method for class tabxplor_fmt

Usage

```
## S3 method for class 'tabxplor_fmt'
x$name
```

Arguments

`x`	A tabxplor_fmt object.
`name`	The name of the field to extract.

Value

The relevant field of the tabxplor_fmt.

Index