

# Package ‘uteals’

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**Description** Provides decorators, transformers, and utility functions to extend the 'teal' framework for interactive data analysis applications. Implements methods for data visualization enhancement, statistical data transformations, and workflow integration tools. Designed to support clinical and pharmaceutical research workflows within the 'teal' ecosystem through modular and reusable components.

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uteals-package                      *Decorators and transformators for teal modules.*

---

## Description

Provides decorator and transformator modules for teal modules.

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**See Also**

Useful links:

- <https://github.com/phuse-org/uteals>

---

create\_rel\_risk\_transformator  
*Create Relative risk column*

---

**Description**

[Experimental]

**Usage**

```
create_rel_risk_transformator(dataname, column_name, control_group, label_name)
```

**Arguments**

|               |  |
|---------------|--|
| dataname      | (character(1)) the name of the dataset which columns will be used for possible transformation. dataname should be passed in quotes ex: "ADSL". |
| column_name   | (character(1)) field or variable from the dataset.   |
| control_group | (character(1)) one of the existing level from the selected column_name.  |
| label_name    | (character(1)) label for the new field or variable.  |

**Details**

This transformator allows the user to select a column and control group from the dataset and create a relative risk column.

**Value**

```
teal::teal_transform_module
```

**Examples**

```
app <- teal::init(  
  data = teal.data::teal_data(IRIS = iris, code = "IRIS <- iris"),  
  modules = teal::modules(  
    teal::example_module(  
      transformers = list(  
        create_rel_risk_transformator("IRIS",  
          label_name = "test",  
          control_group = "setosa", column_name = "Species"  
        )  
      )  
    )  
  )  
)
```

```
)  
if (interactive()) {  
  shiny::shinyApp(app$ui, app$server)  
}
```

---

extract\_modules\_to\_yaml

*Extract Non-Parent Module Labels to YAML*

---

## Description

**[Experimental]** Extracts module labels from a teal modules object, filters out parent modules (grouping containers), and generates a YAML file with the functional modules.

## Usage

```
extract_modules_to_yaml(mods, filepath, verbose = FALSE)
```

## Arguments

|          |  |
|----------|--|
| mods     | (teal_module or teal_modules) a teal modules object containing the module structure. |
| filepath | (character(1)) character string specifying the output YAML file path.                |
| verbose  | (logical(1)) whether to print informational messages. Default is FALSE.              |

## Value

Character vector of non-parent module labels

## Examples

```
# Extract modules from mods object to YAML file  
mods <- teal::modules(  
  teal::example_module("mod1"),  
  teal::example_module("mod2")  
)  
labels <- extract_modules_to_yaml(mods, "panel_str_modules.yaml")  
unlink("panel_str_modules.yaml")  
  
# Clean up  
if (file.exists("panel_str_modules.yaml")) {  
  file.remove("panel_str_modules.yaml")  
}
```

---

```
forestplot_x_decorator
Change X-axis scaling decorator for
teal.modules.clinical::tm_g_forest_rsp.
```

---

## Description

**[Experimental]** A function to create a UI component for selecting a transform function for the forest plot x axis.

## Usage

```
forestplot_x_decorator()
```

## Details

The module creates a UI with a radio control for selecting the transform function. The selected transformation function is applied to the forest plot to update the plot's axis and annotations accordingly.

## Value

`teal::teal_transform_module` Returns a modified plot object with the transformation applied.

## Examples

```
library(teal.modules.clinical)
data <- teal.data::teal_data()
data <- within(data, {
  ADSL <- teal.modules.clinical::tmc_ex_adsl
  ADRS <- teal.modules.clinical::tmc_ex_adrs |>
  dplyr::mutate(AVALC = tern::d_onco_rsp_label(AVALC) |>
    formatters::with_label("Character Result/Finding")) |>
  dplyr::filter(PARAMCD != "OVRINV" | AVISIT == "FOLLOW UP")
})
teal.data::join_keys(data) <- teal.data::default_cdisc_join_keys[names(data)]

ADSL <- data[["ADSL"]]
ADRS <- data[["ADRS"]]

arm_ref_comp <- list(
  ARM = list(
    ref = "B: Placebo",
    comp = c("A: Drug X", "C: Combination")
  ),
  ARMCD = list(
    ref = "ARM B",
    comp = c("ARM A", "ARM C")
  )
)
```

```

)

app <- teal::init(
  data = data,
  modules = teal::modules(
    teal.modules.clinical::tm_g_forest_rsp(
      label = "Forest Response",
      dataname = "ADRS",
      arm_var = choices_selected(
        variable_choices(ADSL, c("ARM", "ARMCD")),
        "ARMCD"
      ),
      arm_ref_comp = arm_ref_comp,
      paramcd = choices_selected(
        value_choices(ADRS, "PARAMCD", "PARAM"),
        "INVET"
      ),
      subgroup_var = choices_selected(
        variable_choices(ADSL, names(ADSL)),
        c("BMRKR2", "SEX")
      ),
      strata_var = choices_selected(
        variable_choices(ADSL, c("STRATA1", "STRATA2")),
        "STRATA2"
      ),
      plot_height = c(600L, 200L, 2000L),
      decorators = list(
        plot = forestplot_x_decorator()
      ),
      default_responses = list(
        BESRSPI = list(
          rsp = c("Stable Disease (SD)", "Not Evaluable (NE)"),
          levels = c(
            "Complete Response (CR)", "Partial Response (PR)", "Stable Disease (SD)",
            "Progressive Disease (PD)", "Not Evaluable (NE)"
          )
        ),
        INVET = list(
          rsp = c("Complete Response (CR)", "Partial Response (PR)"),
          levels = c(
            "Complete Response (CR)", "Not Evaluable (NE)", "Partial Response (PR)",
            "Progressive Disease (PD)", "Stable Disease (SD)"
          )
        ),
        OVRINV = list(
          rsp = c("Progressive Disease (PD)", "Stable Disease (SD)"),
          levels = c("Progressive Disease (PD)", "Stable Disease (SD)", "Not Evaluable (NE)")
        )
      )
    )
  )
)

if (interactive()) {

```

```

  shinyApp(app$ui, app$server)
}

```

---

ggplot\_decorator      ggplot *decorator*

---

## Description

**[Experimental]** Decorator function to update various settings for ggplot plot objects

## Usage

```

ggplot_decorator(
  output_name,
  label_text = "decorator",
  render_ui = c(),
  plot_options = list(title = "", footnote = "", y_breaks = "", y_limits_max = "",
    y_limits_min = "", x_breaks = "", x_labels_discrete = "", x_labels_cont = "",
    y_labels_discrete = "", y_labels_cont = "", font_size_geom_text = "",
    font_size_plot_title = "", font_size_axis_title = "", font_size_axis_text = "")
)

```

## Arguments

|              |  |
|--------------|--|
| output_name  | a name for the output plot object.   |
| label_text   | customized label text for the decorator  |
| render_ui    | vector of ggplot_options from the following list: "title" - Title of the plot, "footnote" - Footnote of the plot, "y_breaks" - Value of breaks(numeric) for y-axis. Note: y_limits_max and y_limits_min should also be provided, "y_limits_max" - Value of y-axis maximum limit(numeric). Note: y_limits_max and y_breaks should also be provided, "y_limits_min" - Value of y-axis minimum limit(numeric). Note: y_breaks and y_limits_min should also be provided, "x_breaks"- Value of breaks for continuous x-axis(numeric). Note: should be comma separated, "x_labels_discrete" - Values of labels for discrete x-axis. Note: should be comma separated, "x_labels_cont" - Values of labels for continuous x-axis. Note: should be comma separated, "y_labels_discrete" - Values of labels for discrete y-axis. Note: should be comma separated, "y_labels_cont" - Values of labels for continuous y-axis. Note: should be comma separated, "font_size_geom_text" - Font size of geom_text. Note: numeric value should be provided, "font_size_plot_title"- Font size of plot title text. Note: numeric value should be provided, "font_size_axis_title"- Font size of axis title text. Note: numeric value should be provided, "font_size_axis_text"- Font size of axis labels text. Note: numeric value should be provided |
| plot_options | named list with the list of values for the ggplot options. The app developer can specify the required list of options while calling the decorator.   |

## Details

The module creates a UI with text controls for specifying the list of ggplot options given in the `plot_options` parameter value. The entered ggplot options are applied to ggplot plot object.

## Value

`teal::teal_transform_module` Returns a modified plot object with the transformation applied.

## Examples

```
data <- teal.data::teal_data()
data <- within(data, {
  ADSL <- teal.data::rADSL
})
teal.data::join_keys(data) <- teal.data::default_cdisc_join_keys[names(data)]

# teal.modules.general >= 0.6.0
app <- teal::init(
  data = data,
  modules = teal::modules(
    teal.modules.general::tm_g_scatterplot(
      label = "Scatterplot Choices",
      x = teal.transform::data_extract_spec(
        dataname = "ADSL",
        select = teal.transform::select_spec(
          label = "Select variable:",
          choices = teal.transform::variable_choices(data[["ADSL"]], c("AGE", "BMRKR1", "BMRKR2")),
          selected = "AGE",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
    ),
    y = teal.transform::data_extract_spec(
      dataname = "ADSL",
      select = teal.transform::select_spec(
        label = "Select variable:",
        choices = teal.transform::variable_choices(data[["ADSL"]], c("AGE", "BMRKR1", "BMRKR2")),
        selected = "BMRKR1",
        multiple = FALSE,
        fixed = FALSE
      )
    ),
    decorators = list(
      plot = ggplot_decorator(
        output_name = "plot", render_ui = c("title", "footnote", "font_size_axis_title")
      )
    )
  )
)

if (interactive()) {
```

```

    shiny::shinyApp(app$sui, app$server)
  }

```

---

g\_forest\_with\_transform

*Create a forest plot from an rtable with x transform settings*

---

## Description

**[Experimental]** Given a `rtables::rtable()` object with at least one column with a single value and one column with 2 values, converts table to a `ggplot2::ggplot()` object and generates an accompanying forest plot. The table and forest plot are printed side-by-side.

## Usage

```

g_forest_with_transform(
  tbl,
  col_x = attr(tbl, "col_x"),
  col_ci = attr(tbl, "col_ci"),
  vline = 1,
  forest_header = attr(tbl, "forest_header"),
  xlim = c(0.1, 10),
  transform_x = NA,
  x_at = c(0.1, 1, 10),
  width_columns = NULL,
  lbl_col_padding = 0,
  rel_width_forest = 0.25,
  font_size = 12,
  col_symbol_size = attr(tbl, "col_symbol_size"),
  col = getOption("ggplot2.discrete.colour")[1],
  ggtheme = NULL,
  as_list = FALSE
)

```

## Arguments

|        |   |
|--------|---|
| tbl    | (VTableTree)<br>rtables table with at least one column with a single value and one column with 2 values.  |
| col_x  | (integer(1) or NULL)<br>column index with estimator. By default tries to get this from tbl attribute col_x, otherwise needs to be manually specified. If NULL, points will be excluded from forest plot.            |
| col_ci | (integer(1) or NULL)<br>column index with confidence intervals. By default tries to get this from tbl attribute col_ci, otherwise needs to be manually specified. If NULL, lines will be excluded from forest plot. |

|                               |  |
|-------------------------------|--|
| <code>vline</code>            | (numeric)<br>position of the vertical reference line on the plot. like 0 and 1 in forest plot.   |
| <code>forest_header</code>    | (character(2))<br>text displayed to the left and right of <code>vline</code> , respectively. If <code>vline = NULL</code> then <code>forest_header</code> is not printed. By default tries to get this from <code>tbl</code> attribute <code>forest_header</code> . If <code>NULL</code> , defaults will be extracted from the table if possible, and set to "Comparison\nBetter" and "Treatment\nBetter" if not.  |
| <code>xlim</code>             | (numeric)<br>range of x-axis limits. Example: <code>c(0.1, 10)</code>  |
| <code>transform_x</code>      | (character)<br>function for x-values transformation  |
| <code>x_at</code>             | (numeric)<br>specifies the tick marks on the axis. The value of <code>union(xlim, vline)</code> or Example: <code>c(0.1, 1, 10)</code> .   |
| <code>width_columns</code>    | (numeric)<br>a vector of column widths. Each element's position in <code>colwidths</code> corresponds to the column of <code>tbl</code> in the same position. If <code>NULL</code> , column widths are calculated according to maximum number of characters per column.  |
| <code>lbl_col_padding</code>  | (numeric)<br>padding between label and columns value. Default is 0.  |
| <code>rel_width_forest</code> | (proportion)<br>proportion of total width to allocate to the forest plot. Relative width of table is then <code>1 - rel_width_forest</code> . If <code>as_list = TRUE</code> , this parameter is ignored.  |
| <code>font_size</code>        | (numeric(1))<br>font size.   |
| <code>col_symbol_size</code>  | (numeric or <code>NULL</code> )<br>column index from <code>tbl</code> containing data to be used to determine relative size for estimator plot symbol. Typically, the symbol size is proportional to the sample size used to calculate the estimator. If <code>NULL</code> , the same symbol size is used for all subgroups. By default tries to get this from <code>tbl</code> attribute <code>col_symbol_size</code> , otherwise needs to be manually specified. |
| <code>col</code>              | (character)<br>color(s).   |
| <code>ggtheme</code>          | (theme)<br>a graphical theme as provided by <code>ggplot2</code> to control styling of the plot.   |
| <code>as_list</code>          | (flag)<br>whether the two <code>ggplot</code> objects should be returned as a list. If <code>TRUE</code> , a named list with two elements, <code>table</code> and <code>plot</code> , will be returned. If <code>FALSE</code> (default) the table and forest plot are printed side-by-side via <code>cowplot::plot_grid()</code> .   |

**Value**

`ggplot` forest plot and table.

---

|               |                                     |
|---------------|-------------------------------------|
| keep_by_label | <i>Filter Teal Modules by Label</i> |
|---------------|-------------------------------------|

---

### Description

**[Experimental]** Recursively filters a teal modules object to keep only modules whose labels match the specified labels. Removes modules that don't match and empty parent containers.

### Usage

```
keep_by_label(x, label)
```

### Arguments

x (teal\_module or teal\_modules) the object to filter.  
label (character(1)) character vector of module labels to keep.

### Value

Filtered teal\_modules or teal\_module object, or NULL if none matches.

### Examples

```
# Keep only specific modules by label
mods <- teal::modules(
  teal::example_module("mod1"),
  teal::example_module("mod2")
)
filtered_mods <- keep_by_label(mods, c("Data Table", "Disposition"))
```

---

|                            |  |
|----------------------------|--|
| merge_levels_transformator | <i>Combine levels of a variable into one level</i> |
|----------------------------|--|

---

### Description

**[Experimental]**

### Usage

```
merge_levels_transformator(dataname, predefined = list())
```

**Arguments**

|            |  |
|------------|--|
| dataname   | (character(1)) the name of the dataset which columns will be used for possible transformation. |
| predefined | (list) the list which has variable name, levels and new label                                  |

**Details**

This transformator allows the user to select a column from the dataset and merge levels of this column into a new level. Only selected levels are affected.

**Value**

teal::teal\_transform\_module

**Examples**

```
app <- teal::init(
  data = teal.data::teal_data(IRIS = iris, code = "IRIS <- iris"),
  modules = teal::modules(
    teal::example_module(
      transformers = list(merge_levels_transformator(
        dataname = "IRIS",
        predefined = list(
          list("Species", "setosa", "SETOSA_WITHIN_FIX"),
          list("Petal.Width", c(0.2, 0.3, 0.5), 12)
        )
      )
    )
  )
)
if (interactive()) {
  shiny::shinyApp(app$ui, app$server)
}
```

---

or\_filtering\_transformator

*Apply Logical AND/OR filter transformations*

---

**Description**

**[Experimental]** This transformator provides users with flexible, dynamic filtering capabilities for datasets.

Each instance of the transformator operates on a **single dataset**. Users can define multiple filter blocks, where:

- Conditions within each block are combined with **logical AND**.
- Multiple blocks are combined with **logical OR**.

This allows creating complex filter expressions like:

(Condition1 AND Condition2) OR (Condition3 AND Condition4)

To apply filtering across **multiple datasets**, users can instantiate multiple instances of this module, each configured for a different dataset. Each module call is independent and manages filters for its specific dataset.

- **Supported Data Types & Expressions:**

- Supports filtering on character, factor, and numeric columns.
- Conditions can use operators: ==, !=, <, >, <=, >=.
- Conditions are specified as simple expressions, e.g., columnA == 'value' columnB != 5 columnC >= 10
- Each block's conditions are combined with **AND**.
- Multiple blocks are combined with **OR**.

- **Features:**

- **Add Multiple OR Blocks:** Dynamically add new blocks for alternative conditions.
- **Add Conditions:** Within each block, add multiple conditions, with duplicate prevention.
- **Preview Filter Expression:** Generate and display the current combined filter expression.
- **Remove Conditions:** Remove individual conditions within a block.
- **Expression Generation:** The resulting expression can be directly used with `dplyr::filter()` or similar functions.

- **Usage Pattern:**

- Call the module multiple times with different dataset names to filter multiple datasets independently.
- Each call manages its own filter state and expression.
- Users can build complex filters per dataset and apply or combine them as needed.

## Usage

```
or_filtering_transformator(dataname)
```

## Arguments

dataname (character(1)) Name of the dataset to filter. Pass a single dataset name as a string.

## Value

```
teal::teal_transform_module
```

## Examples

```
app <- teal::init(
  data = teal.data::teal_data(IRIS = iris),
  modules = teal::modules(teal::example_module(
    transformers = list(or_filtering_transformator("IRIS"))
  ))
)
```

```

)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

---

patchwork\_plot\_decorator

*Patchwork Decorator*

---

### Description

**[Experimental]** Decorator function to add plot title and footnote to patchwork plots

### Usage

```
patchwork_plot_decorator(output_name, label_text = "decorator")
```

### Arguments

output\_name (character(1)) A name for the output plot object.  
 label\_text (character(1)) A customized label text for the decorator.

### Details

The module creates a UI with text controls for plot title and footnote. The entered title and footnote text are applied to the patchwork plots.

### Value

(teal.data : :qenv) Returns a modified plot object with the transformation applied.

---

remove\_by\_label

*Remove Teal Modules by Label*

---

### Description

**[Experimental]** Recursively removes modules from a teal modules structure that match the specified labels.

### Usage

```
remove_by_label(x, label)
```

### Arguments

x (teal\_module or teal\_modules) The object to filter.  
 label (character(1)) character vector of module labels to remove.

**Value**

The filtered teal modules object with matching modules removed, or NULL if all modules are removed.

**Examples**

```
mods <- teal::modules(
  teal::example_module("mod1"),
  teal::example_module("mod2")
)
# Remove a single module
filtered_mods <- remove_by_label(mods, "Deaths")

# Remove multiple modules
filtered_mods <- remove_by_label(mods, c("Deaths", "Lab Summary Table"))
```

---

title\_footer\_decorator

*Title and Footer Decorator*

---

**Description**

**[Experimental]** A function to create a UI component for selecting a title and footer for tables or plots. It reads title information from a specified Excel file and allows users to choose a title from the provided options. It also provides user with flexibility to customize the title and footer according to their specific needs.

**Usage**

```
title_footer_decorator(
  output_name,
  titles_file,
  choices = NULL,
  selected = NULL
)
```

**Arguments**

|             |  |
|-------------|--|
| output_name | (character(1)) a name for the output object (e.g., a plot or table).   |
| titles_file | (character(1)) the path to an Excel file containing title and footer information. The function expects the titles to be in the first sheet named Sheet1. |
| choices     | (character) an array of titles and footers, which are available for selection. Default NULL, indicates all titles and footers are available.             |
| selected    | (character(1)) the selected title or footer. Default NULL, indicates no title or footer is selected.   |

**Details**

The module creates a UI with a dropdown for selecting a title. Once a title is selected, it updates the output by either adding titles to a table or modifying a plot's title and caption accordingly. Additionally, it includes a checkbox that user can check to enable customization, allowing them to enter their own values for the title and footer in the designated input fields.

**Value**

```
teal::teal_transform_module()
```

**See Also**

For the exact Excel workbook layout expected by this function, see the package vignette: `vignette("title-footer-decora package = "uteals")`

**Examples**

```
library(openxlsx)
library(teal.modules.general)

example_excel <- data.frame(
  `TABLE ID` = c(
    "DO_NOT_DELETE",
    "TSFAE01A", "TSFAE01A", "TSFAE01A",
    "TSFAE01B", "TSFAE01B"
  ),
  IDENTIFIER = c(
    "DO_NOT_DELETE",
    "TITLE", "FOOTNOTE1", "FOOTNOTE2",
    "TITLE", "FOOTNOTE1"
  ),
  TEXT = c(
    "DO_NOT_DELETE",
    "Adverse Events Summary A", "Source: Clinical Study Report", "Confidential",
    "Adverse Events Summary B", "Draft Version"
  ),
  stringsAsFactors = FALSE,
  check.names = FALSE
)

temp_titles <- tempfile(fileext = ".xlsx")
write.xlsx(example_excel, temp_titles, sheetName = "Sheet1", asTable = TRUE)
plot_module <- tm_g_scatterplot(
  label = "Scatter Plot",
  x = data_extract_spec(
    dataname = "IRIS",
    select = select_spec(
      choices = variable_choices(
        "IRIS", c("Sepal.Length", "Sepal.Width")
      ), selected = "Sepal.Length"
    )
  )
)
```

```

),
y = data_extract_spec(
  dataname = "IRIS",
  select = select_spec(
    choices = variable_choices(
      "IRIS", c("Petal.Length", "Petal.Width")
    ), selected = "Petal.Length"
  )
),
decorators = list(
  plot = title_footer_decorator(
    "plot", temp_titles,
    choices = c("TSFAE01A", "TSFAE01B"), selected = NULL
  )
)
)

# Initialize the teal app
app <- init(
  data = teal_data(IRIS = iris),
  modules = list(plot_module)
)

# Run the app
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

---

watermark\_decorator     *Watermark Decorator*

---

## Description

**[Experimental]** A function to create a UI component for selecting watermark text for plots. Note: Currently tables are not supported

## Usage

```
watermark_decorator(output_name, watermark_text = "", font_size = 90)
```

## Arguments

`output_name`     (character(1)) a name for the output object (e.g., a plot or table).

`watermark_text` (character(1)) text to display for the watermark.

`font_size`       (character(1)) font size for the watermark text.

**Details**

The module creates a UI with `textInput` for specifying watermark text and font size. the entered watermark text is displayed with a default `gridify` layout.

**Value**

```
teal::teal_transform_module()
```

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