

Package ‘teal.modules.general’

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Type Package

Title General Modules for 'teal' Applications

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Description Prebuilt 'shiny' modules containing tools for viewing data, visualizing data, understanding missing and outlier values within your data and performing simple data analysis. This extends 'teal' framework that supports reproducible research and analysis.

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URL <https://insightsengineering.github.io/teal.modules.general/>,
<https://github.com/insightsengineering/teal.modules.general/>

BugReports <https://github.com/insightsengineering/teal.modules.general/issues>

Depends ggmosaic (>= 0.3.0), ggplot2 (>= 3.4.0), R (>= 3.6), shiny (>= 1.6.0), teal (>= 0.15.1), teal.transform (>= 0.5.0)

Imports checkmate (>= 2.1.0), dplyr (>= 1.0.5), DT (>= 0.13),forcats (>= 1.0.0), grid, logger (>= 0.2.0), scales, shinyjs, shinyTree (>= 0.2.8), shinyvalidate, shinyWidgets (>= 0.5.1), stats, stringr (>= 1.4.1), teal.code (>= 0.5.0), teal.data (>= 0.5.0), teal.logger (>= 0.1.1), teal.reporter (>= 0.3.0), teal.widgets (>= 0.4.0), tern (>= 0.9.3), tibble (>= 2.0.0), tidyR (>= 0.8.3), tools, utils

Suggests broom (>= 0.7.10), colourpicker, ggExtra, ggpmisc (>= 0.4.3), ggpp, ggrepel, goftest, gridExtra, htmlwidgets, jsonlite, knitr (>= 1.42), lattice (>= 0.18-4), MASS, nestcolor (>= 0.1.0), rlang (>= 1.0.0), rtables (>= 0.6.6), sparkline, testthat (>= 3.0.4)

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add_facet_labels	<i>Add labels for facets to a ggplot2 object</i>
------------------	--

Description

Enhances a ggplot2 plot by adding labels that describe the faceting variables along the x and y axes.

Usage

```
add_facet_labels(p, xfacet_label = NULL, yfacet_label = NULL)
```

Arguments

p	(ggplot2) object to which facet labels will be added.
xfacet_label	(character) Label for the facet along the x-axis. If NULL, no label is added. If a vector, labels are joined with " & ".
yfacet_label	(character) Label for the facet along the y-axis. Similar behavior to xfacet_label.

Value

Returns grid or grob object (to be drawn with `grid.draw`)

Examples

```
library(ggplot2)
library(grid)

p <- ggplot(mtcars) +
  aes(x = mpg, y = disp) +
  geom_point() +
  facet_grid(gear ~ cyl)

xfacet_label <- "cylinders"
yfacet_label <- "gear"
res <- add_facet_labels(p, xfacet_label, yfacet_label)
grid.newpage()
grid.draw(res)

grid.newpage()
```

```
grid.draw(add_facet_labels(p, xfacet_label = NULL, yfacet_label))
grid.newpage()
grid.draw(add_facet_labels(p, xfacet_label, yfacet_label = NULL))
grid.newpage()
grid.draw(add_facet_labels(p, xfacet_label = NULL, yfacet_label = NULL))
```

get_scatterplotmatrix_stats*Get stats for x-y pairs in scatterplot matrix***Description**

Uses `stats::cor.test()` per default for all numerical input variables and converts results to character vector. Could be extended if different stats for different variable types are needed. Meant to be called from `lattice::panel.text()`.

Usage

```
get_scatterplotmatrix_stats(
  x,
  y,
  .f = stats::cor.test,
  .f_args = list(),
  round_stat = 2,
  round_pval = 4
)
```

Arguments

<code>x, y</code>	(numeric) vectors of data values. <code>x</code> and <code>y</code> must have the same length.
<code>.f</code>	(function) function that accepts <code>x</code> and <code>y</code> as formula input $\sim x + y$. Default <code>stats::cor.test</code> .
<code>.f_args</code>	(list) of arguments to be passed to <code>.f</code> .
<code>round_stat</code>	(integer(1)) optional, number of decimal places to use when rounding the estimate.
<code>round_pval</code>	(integer(1)) optional, number of decimal places to use when rounding the p-value.

Details

Presently we need to use a formula input for `stats::cor.test` because `na.fail` only gets evaluated when a formula is passed (see below).

```
x = c(1,3,5,7,NA)
y = c(3,6,7,8,1)
stats::cor.test(x, y, na.action = "na.fail")
stats::cor.test(~ x + y, na.action = "na.fail")
```

Value

Character with stats. For `stats::cor.test()` correlation coefficient and p-value.

Examples

```
set.seed(1)
x <- runif(25, 0, 1)
y <- runif(25, 0, 1)
x[c(3, 10, 18)] <- NA

get_scatterplotmatrix_stats(x, y, .f = stats::cor.test, .f_args = list(method = "pearson"))
get_scatterplotmatrix_stats(x, y, .f = stats::cor.test, .f_args = list(
  method = "pearson",
  na.action = na.fail
))
```

tm_a_pca

teal module: Principal component analysis

Description

Module conducts principal component analysis (PCA) on a given dataset and offers different ways of visualizing the outcomes, including elbow plot, circle plot, biplot, and eigenvector plot. Additionally, it enables dynamic customization of plot aesthetics, such as opacity, size, and font size, through UI inputs.

Usage

```
tm_a_pca(
  label = "Principal Component Analysis",
  dat,
  plot_height = c(600, 200, 2000),
  plot_width = NULL,
  ggtheme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void"),
  ggplot2_args = teal.widgets::ggplot2_args(),
  rotate_xaxis_labels = FALSE,
  font_size = c(12, 8, 20),
  alpha = c(1, 0, 1),
  size = c(2, 1, 8),
  pre_output = NULL,
  post_output = NULL
)
```

Arguments

label	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
dat	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) specifying columns used to compute PCA.
plot_height	(numeric) optional, specifies the plot height as a three-element vector of value, <code>min</code> , and <code>max</code> intended for use with a slider UI element.
plot_width	(numeric) optional, specifies the plot width as a three-element vector of value, <code>min</code> , and <code>max</code> for a slider encoding the plot width.
ggtheme	(character) optional, <code>ggplot2</code> theme to be used by default. Defaults to "gray".
ggplot2_args	(<code>ggplot2_args</code>) optional, object created by <code>teal.widgets::ggplot2_args()</code> with settings for all the plots or named list of <code>ggplot2_args</code> objects for plot-specific settings. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. List names should match the following: <code>c("default", "Elbow plot", "Circle plot", "Biplot", "Eigenvector plot")</code> . For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code> .
rotate_xaxis_labels	(logical) optional, whether to rotate plot X axis labels. Does not rotate by default (FALSE).
font_size	(numeric) optional, specifies font size. It controls the font size for plot titles, axis labels, and legends. <ul style="list-style-type: none"> • If vector of length == 1 then the font sizes will have a fixed size. • while vector of value, <code>min</code>, and <code>max</code> allows dynamic adjustment.
alpha	(integer(1) or integer(3)) optional, specifies point opacity. <ul style="list-style-type: none"> • When the length of alpha is one: the plot points will have a fixed opacity. • When the length of alpha is three: the plot points opacity are dynamically adjusted based on vector of value, <code>min</code>, and <code>max</code>.
size	(integer(1) or integer(3)) optional, specifies point size. <ul style="list-style-type: none"> • When the length of size is one: the plot point sizes will have a fixed size. • When the length of size is three: the plot points size are dynamically adjusted based on vector of value, <code>min</code>, and <code>max</code>.
pre_output	(<code>shiny.tag</code>) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
post_output	(<code>shiny.tag</code>) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.

Value

Object of class `teal_module` to be used in `teal` applications.

Examples

```
library(teal.widgets)

# general data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  USArests <- USArests
})
datanames(data) <- "USArests"

app <- init(
  data = data,
  modules = modules(
    tm_a_pca(
      "PCA",
      dat = data_extract_spec(
        data$name = "USArests",
        select = select_spec(
          choices = variable_choices(
            data = data[["USArests"]], c("Murder", "Assault", "UrbanPop", "Rape")
          ),
          selected = c("Murder", "Assault"),
          multiple = TRUE
        ),
        filter = NULL
      ),
      ggplot2_args = ggplot2_args(
        labs = list(subtitle = "Plot generated by PCA Module")
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

# CDISC data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  ADSL <- rADSL
})
datanames(data) <- "ADSL"
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(
    tm_a_pca(
      "PCA",
```

```

dat = data_extract_spec(
  dataname = "ADSL",
  select = select_spec(
    choices = variable_choices(
      data = data[["ADSL"]], c("BMRKR1", "AGE", "EOSDY")
    ),
    selected = c("BMRKR1", "AGE"),
    multiple = TRUE
  ),
  filter = NULL
),
ggplot2_args = ggplot2_args(
  labs = list(subtitle = "Plot generated by PCA Module")
)
)
)
)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_a_regression*teal module: Scatterplot and regression analysis***Description**

Module for visualizing regression analysis, including scatterplots and various regression diagnostics plots. It allows users to explore the relationship between a set of regressors and a response variable, visualize residuals, and identify outliers.

Usage

```

tm_a_regression(
  label = "Regression Analysis",
  regressor,
  response,
  plot_height = c(600, 200, 2000),
  plot_width = NULL,
  alpha = c(1, 0, 1),
  size = c(2, 1, 8),
  ggtheme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void"),
  ggplot2_args = teal.widgets::ggplot2_args(),
  pre_output = NULL,
  post_output = NULL,
  default_plot_type = 1,
  default_outlier_label = "USUBJID",
  label_segment_threshold = c(0.5, 0, 10)
)

```

Arguments

<code>label</code>	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
<code>regressor</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Regressor variables from an incoming dataset with filtering and selecting.
<code>response</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Response variables from an incoming dataset with filtering and selecting.
<code>plot_height</code>	(<code>numeric</code>) optional, specifies the plot height as a three-element vector of <code>value</code> , <code>min</code> , and <code>max</code> intended for use with a slider UI element.
<code>plot_width</code>	(<code>numeric</code>) optional, specifies the plot width as a three-element vector of <code>value</code> , <code>min</code> , and <code>max</code> for a slider encoding the plot width.
<code>alpha</code>	(<code>integer(1)</code> or <code>integer(3)</code>) optional, specifies point opacity. <ul style="list-style-type: none"> • When the length of <code>alpha</code> is one: the plot points will have a fixed opacity. • When the length of <code>alpha</code> is three: the plot points opacity are dynamically adjusted based on vector of <code>value</code>, <code>min</code>, and <code>max</code>.
<code>size</code>	(<code>integer(1)</code> or <code>integer(3)</code>) optional, specifies point size. <ul style="list-style-type: none"> • When the length of <code>size</code> is one: the plot point sizes will have a fixed size. • When the length of <code>size</code> is three: the plot points size are dynamically adjusted based on vector of <code>value</code>, <code>min</code>, and <code>max</code>.
<code>ggtheme</code>	(<code>character</code>) optional, <code>ggplot2</code> theme to be used by default. Defaults to "gray".
<code>ggplot2_args</code>	(<code>ggplot2_args</code>) optional, object created by <code>teal.widgets::ggplot2_args()</code> with settings for all the plots or named list of <code>ggplot2_args</code> objects for plot-specific settings. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. List names should match the following: <code>c("default", "Response vs Regressor", "Residuals vs Fitted")</code> . For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code> .
<code>pre_output</code>	(<code>shiny.tag</code>) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
<code>post_output</code>	(<code>shiny.tag</code>) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.
<code>default_plot_type</code>	(<code>numeric</code>) optional, defaults to "Response vs Regressor". <ol style="list-style-type: none"> 1. Response vs Regressor 2. Residuals vs Fitted 3. Normal Q-Q 4. Scale-Location 5. Cook's distance 6. Residuals vs Leverage 7. Cook's dist vs Leverage

default_outlier_label
 (character) optional, default column selected to label outliers.

label_segment_threshold
 (numeric(1) or numeric(3)) Minimum distance between label and point on the plot that triggers the creation of a line segment between the two. This may happen when the label cannot be placed next to the point as it overlaps another label or point. The value is used as the `min.segment.length` parameter to the `ggrepel::geom_text_repel()` function.
 It can take the following forms:

- numeric(1): Fixed value used for the minimum distance and the slider is not presented in the UI.
- numeric(3): A slider is presented in the UI (under "Plot settings") to adjust the minimum distance dynamically.

It takes the form of `c(value, min, max)` and it is passed to the `value_min_max` argument in `teal.widgets::optionalSliderInputValMinMax`.

Value

Object of class `teal_module` to be used in `teal` applications.

Note

For more examples, please see the vignette "Using regression plots" via `vignette("using-regression-plots", package = "teal.modules.general")`.

Examples

```
# general data example
library(teal.widgets)

data <- teal_data()
data <- within(data, {
  require(nestcolor)
  CO2 <- CO2
})
datanames(data) <- c("CO2")

app <- init(
  data = data,
  modules = modules(
    tm_a_regression(
      label = "Regression",
      response = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = "uptake",
          selected = "uptake",
          multiple = FALSE,
          fixed = TRUE
        )
      )
    )
  )
)
```

```
),
regressor = data_extract_spec(
  dataname = "CO2",
  select = select_spec(
    label = "Select variables:",
    choices = variable_choices(data[["CO2"]], c("conc", "Treatment")),
    selected = "conc",
    multiple = TRUE,
    fixed = FALSE
  )
),
ggplot2_args = ggplot2_args(
  labs = list(subtitle = "Plot generated by Regression Module")
)
)
)
)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

# CDISC data example
library(teal.widgets)

data <- teal_data()
data <- within(data, {
  require(nestcolor)
  ADSL <- rADSL
})
datanames(data) <- "ADSL"
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(
    tm_a_regression(
      label = "Regression",
      response = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          label = "Select variable:",
          choices = "BMRKR1",
          selected = "BMRKR1",
          multiple = FALSE,
          fixed = TRUE
        )
      ),
      regressor = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          label = "Select variables:",
          choices = variable_choices(data[["ADSL"]], c("AGE", "SEX", "RACE")),
          selected = "AGE",
        )
      )
    )
  )
)
```

```

        multiple = TRUE,
        fixed = FALSE
    )
),
ggplot2_args = ggplot2_args(
    labs = list(subtitle = "Plot generated by Regression Module")
)
)
)
)
)
if (interactive()) {
    shinyApp(app$ui, app$server)
}

```

tm_data_table*teal module: Data table viewer***Description**

Module provides a dynamic and interactive way to view `data.frames` in a `teal` application. It uses the `DT` package to display data tables in a paginated, searchable, and sortable format, which helps to enhance data exploration and analysis.

Usage

```

tm_data_table(
    label = "Data Table",
    variables_selected = list(),
    datasets_selected = character(0),
    dt_args = list(),
    dt_options = list(searching = FALSE, pageLength = 30, lengthMenu = c(5, 15, 30, 100),
                      scrollX = TRUE),
    server_rendering = FALSE,
    pre_output = NULL,
    post_output = NULL
)

```

Arguments

`label` (character(1)) Label shown in the navigation item for the module or module group. For `modules()` defaults to "root". See Details.

`variables_selected`

(named list) Character vectors of the variables (i.e. columns) which should be initially shown for each dataset. Names of list elements should correspond to the names of the datasets available in the app. If no entry is specified for a dataset, the first six variables from that dataset will initially be shown.

datasets_selected	(character) A vector of datasets which should be shown and in what order. Names in the vector have to correspond with datasets names. If vector of length == 0 (default) then all datasets are shown. Note: Only datasets of the <code>data.frame</code> class are compatible.
dt_args	(named list) Additional arguments to be passed to <code>DT::datatable()</code> (must not include data or options).
dt_options	(named list) The options argument to <code>DT::datatable</code> . By default <code>list(searching = FALSE, pageLength = 30, lengthMenu = c(5, 15, 30, 100), scrollX = TRUE)</code>
server_rendering	(logical) should the data table be rendered server side (see <code>server</code> argument of <code>DT::renderDataTable()</code>)
pre_output	(shiny.tag) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
post_output	(shiny.tag) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.

Details

The DT package has an option `DT.TOJSON_ARGS` to show Inf and NA in data tables. Configure the `DT.TOJSON_ARGS` option via `options(DT.TOJSON_ARGS = list(na = "string"))` before running the module. Note though that sorting of numeric columns with NA/Inf will be lexicographic not numerical.

Value

Object of class `teal_module` to be used in teal applications.

Examples

```
# general data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  iris <- iris
})
datanames(data) <- c("iris")

app <- init(
  data = data,
  modules = modules(
    tm_data_table(
      variables_selected = list(
        iris = c("Sepal.Length", "Sepal.Width", "Petal.Length", "Petal.Width", "Species")
      ),
      dt_args = list(caption = "ADSL Table Caption")
    )
  )
)
```

```

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

# CDISC data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  ADSL <- rADSL
})
datanames(data) <- "ADSL"
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(
    tm_data_table(
      variables_selected = list(ADSL = c("STUDYID", "USUBJID", "SUBJID", "SITEID", "AGE", "SEX")),
      dt_args = list(caption = "ADSL Table Caption")
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_file_viewer *teal module: File viewer*

Description

The file viewer module provides a tool to view static files. Supported formats include text formats, PDF, PNG APNG, JPEG SVG, WEBP, GIF and BMP.

Usage

```
tm_file_viewer(
  label = "File Viewer Module",
  input_path = list(`Current Working Directory` = ".")
)
```

Arguments

label	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
input_path	(list) of the input paths, optional. Each element can be: Paths can be specified as absolute paths or relative to the running directory of the application. Default to the current working directory if not supplied.

Value

Object of class `teal_module` to be used in `teal` applications.

Examples

```
data <- teal_data()
data <- within(data, {
  data <- data.frame(1)
})
datanames(data) <- c("data")

app <- init(
  data = data,
  modules = modules(
    tm_file_viewer(
      input_path = list(
        folder = system.file("sample_files", package = "teal.modules.general"),
        png = system.file("sample_files/sample_file.png", package = "teal.modules.general"),
        txt = system.file("sample_files/sample_file.txt", package = "teal.modules.general"),
        url = "https://fda.gov/files/drugs/published/Portable-Document-Format-Specifications.pdf"
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}
```

tm_front_page

teal module: Front page

Description

Creates a simple front page for `teal` applications, displaying introductory text, tables, additional html or shiny tags, and footnotes.

Usage

```
tm_front_page(
  label = "Front page",
  header_text = character(0),
  tables = list(),
  additional_tags = tagList(),
  footnotes = character(0),
  show_metadata = FALSE
)
```

Arguments

<code>label</code>	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
<code>header_text</code>	(character vector) text to be shown at the top of the module, for each element, if named the name is shown first in bold as a header followed by the value. The first element's header is displayed larger than the others.
<code>tables</code>	(named list of <code>data.frames</code>) tables to be shown in the module.
<code>additional_tags</code>	(<code>shiny.tag.list</code> or <code>html</code>) additional shiny tags or html to be included after the table, for example to include an image, <code>tagList(tags\$img(src = "image.png"))</code> or to include further <code>html</code> , <code>HTML("html text here")</code> .
<code>footnotes</code>	(character vector) of text to be shown at the bottom of the module, for each element, if named the name is shown first in bold, followed by the value.
<code>show_metadata</code>	(logical) indicating whether the metadata of the datasets be available on the module.

Value

Object of class `teal_module` to be used in `teal` applications.

Examples

```

data <- teal_data()
data <- within(data, {
  require(nestcolor)
  ADSL <- rADSL
  attr(ADSL, "metadata") <- list("Author" = "NEST team", "data_source" = "synthetic data")
})
datanames(data) <- "ADSL"
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

table_1 <- data.frame(Info = c("A", "B"), Text = c("A", "B"))
table_2 <- data.frame(`Column 1` = c("C", "D"), `Column 2` = c(5.5, 6.6), `Column 3` = c("A", "B"))
table_3 <- data.frame(Info = c("E", "F"), Text = c("G", "H"))

table_input <- list(
  "Table 1" = table_1,
  "Table 2" = table_2,
  "Table 3" = table_3
)

app <- init(
  data = data,
  modules = modules(
    tm_front_page(
      header_text = c(
        "Important information" = "It can go here.",
        "Other information" = "Can go here."
      ),
      ...
    )
  )
)
  
```

```

    tables = table_input,
    additional_tags = HTML("Additional HTML or shiny tags go here <br>"),
    footnotes = c("X" = "is the first footnote", "Y is the second footnote"),
    show_metadata = TRUE
  )
),
header = tags$h1("Sample Application"),
footer = tags$p("Application footer"),
)

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

```

tm_g_association

teal module: Stack plots of variables and show association with reference variable

Description

Module provides functionality for visualizing the distribution of variables and their association with a reference variable. It supports configuring the appearance of the plots, including themes and whether to show associations.

Usage

```

tm_g_association(
  label = "Association",
  ref,
  vars,
  show_association = TRUE,
  plot_height = c(600, 400, 5000),
  plot_width = NULL,
  distribution_theme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic",
    "void"),
  association_theme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic",
    "void"),
  pre_output = NULL,
  post_output = NULL,
  ggplot2_args = teal.widgets::ggplot2_args()
)

```

Arguments

label	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
-------	---

<code>ref</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Reference variable, must accept a <code>data_extract_spec</code> with <code>select_spec(multiple = FALSE)</code> to ensure single selection option.
<code>vars</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Variables to be associated with the reference variable.
<code>show_association</code>	(logical) optional, whether show association of <code>vars</code> with reference variable. Defaults to TRUE.
<code>plot_height</code>	(numeric) optional, specifies the plot height as a three-element vector of value, <code>min</code> , and <code>max</code> intended for use with a slider UI element.
<code>plot_width</code>	(numeric) optional, specifies the plot width as a three-element vector of value, <code>min</code> , and <code>max</code> for a slider encoding the plot width.
<code>distribution_theme, association_theme</code>	(character) optional, <code>ggplot2</code> themes to be used by default. Default to "gray".
<code>pre_output</code>	(<code>shiny.tag</code>) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
<code>post_output</code>	(<code>shiny.tag</code>) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.
<code>ggplot2_args</code>	(<code>ggplot2_args</code>) optional, object created by <code>teal.widgets::ggplot2_args()</code> with settings for all the plots or named list of <code>ggplot2_args</code> objects for plot-specific settings. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. List names should match the following: <code>c("default", "Bivariate1", "Bivariate2")</code> . For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code> .

Value

Object of class `teal_module` to be used in `teal` applications.

Note

For more examples, please see the vignette "Using association plot" via `vignette("using-association-plot", package = "teal.modules.general")`.

Examples

```
library(teal.widgets)

# general data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  CO2 <- CO2
  factors <- names(Filter(isTRUE, vapply(CO2, is.factor, logical(1L))))
```

```
CO2[factors] <- lapply(CO2[factors], as.character)
})
datanames(data) <- c("CO2")

app <- init(
  data = data,
  modules = modules(
    tm_g_association(
      ref = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["CO2"]], c("Plant", "Type", "Treatment")),
          selected = "Plant",
          fixed = FALSE
        )
      ),
      vars = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variables:",
          choices = variable_choices(data[["CO2"]], c("Plant", "Type", "Treatment")),
          selected = "Treatment",
          multiple = TRUE,
          fixed = FALSE
        )
      ),
      ggplot2_args = ggplot2_args(
        labs = list(subtitle = "Plot generated by Association Module")
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

# CDISC data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  ADSL <- rADSL
})
datanames(data) <- "ADSL"
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(
    tm_g_association(
      ref = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
```

```

label = "Select variable:",
choices = variable_choices(
  data[["ADSL"]],
  c("SEX", "RACE", "COUNTRY", "ARM", "STRATA1", "STRATA2", "ITTFL", "BMRKR2")
),
selected = "RACE",
fixed = FALSE
)
),
vars = data_extract_spec(
  dataname = "ADSL",
  select = select_spec(
    label = "Select variables:",
    choices = variable_choices(
      data[["ADSL"]],
      c("SEX", "RACE", "COUNTRY", "ARM", "STRATA1", "STRATA2", "ITTFL", "BMRKR2")
    ),
    selected = "BMRKR2",
    multiple = TRUE,
    fixed = FALSE
  )
),
ggplot2_args = ggplot2_args(
  labs = list(subtitle = "Plot generated by Association Module")
)
)
)
)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_g_bivariate*teal module: Univariate and bivariate visualizations*

Description

Module enables the creation of univariate and bivariate plots, facilitating the exploration of data distributions and relationships between two variables.

Usage

```

tm_g_bivariate(
  label = "Bivariate Plots",
  x,
  y,
  row_facet = NULL,
  col_facet = NULL,
  facet = !is.null(row_facet) || !is.null(col_facet),

```

```

    color = NULL,
    fill = NULL,
    size = NULL,
    use_density = FALSE,
    color_settings = FALSE,
    free_x_scales = FALSE,
    free_y_scales = FALSE,
    plot_height = c(600, 200, 2000),
    plot_width = NULL,
    rotate_xaxis_labels = FALSE,
    swap_axes = FALSE,
    ggtheme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void"),
    ggplot2_args = teal.widgets::ggplot2_args(),
    pre_output = NULL,
    post_output = NULL
)

```

Arguments

label	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
x	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Variable names selected to plot along the x-axis by default. Can be numeric, factor or character. No empty selections are allowed.
y	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Variable names selected to plot along the y-axis by default. Can be numeric, factor or character.
row_facet	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, specification of the data variable(s) to use for faceting rows.
col_facet	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, specification of the data variable(s) to use for faceting columns.
facet	(logical) optional, specifies whether the facet encodings ui elements are toggled on and shown to the user by default. Defaults to TRUE if either <code>row_facet</code> or <code>column_facet</code> are supplied.
color	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, specification of the data variable(s) selected for the outline color inside the coloring settings. It will be applied when <code>color_settings</code> is set to TRUE.
fill	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, specification of the data variable(s) selected for the fill color inside the coloring settings. It will be applied when <code>color_settings</code> is set to TRUE.
size	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, specification of the data variable(s) selected for the size of <code>geom_point</code> plots inside the coloring settings. It will be applied when <code>color_settings</code> is set to TRUE.
use_density	(logical) optional, indicates whether to plot density (TRUE) or frequency (FALSE). Defaults to frequency (FALSE).
color_settings	(logical) Whether coloring, filling and size should be applied and UI tool offered to the user.

<code>free_x_scales</code>	(logical) optional, whether X scaling shall be changeable. Does not allow scaling to be changed by default (FALSE).
<code>free_y_scales</code>	(logical) optional, whether Y scaling shall be changeable. Does not allow scaling to be changed by default (FALSE).
<code>plot_height</code>	(numeric) optional, specifies the plot height as a three-element vector of value, min, and max intended for use with a slider UI element.
<code>plot_width</code>	(numeric) optional, specifies the plot width as a three-element vector of value, min, and max for a slider encoding the plot width.
<code>rotate_xaxis_labels</code>	(logical) optional, whether to rotate plot X axis labels. Does not rotate by default (FALSE).
<code>swap_axes</code>	(logical) optional, whether to swap X and Y axes. Defaults to FALSE.
<code>ggtheme</code>	(character) optional, ggplot2 theme to be used by default. Defaults to "gray".
<code>ggplot2_args</code>	(ggplot2_args) object created by <code>teal.widgets::ggplot2_args()</code> with settings for the module plot. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code>
<code>pre_output</code>	(shiny.tag) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
<code>post_output</code>	(shiny.tag) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.

Details

This is a general module to visualize 1 & 2 dimensional data.

Value

Object of class `teal_module` to be used in `teal` applications.

Note

For more examples, please see the vignette "Using bivariate plot" via `vignette("using-bivariate-plot", package = "teal.modules.general")`.

Examples

```
library(teal.widgets)

# general data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  C02 <- data.frame(C02)
```

```
})
datanames(data) <- c("CO2")
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(
    tm_g_bivariate(
      x = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["CO2"]]),
          selected = "conc",
          fixed = FALSE
        )
      ),
      y = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["CO2"]]),
          selected = "uptake",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      row_facet = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["CO2"]]),
          selected = "Type",
          fixed = FALSE
        )
      ),
      col_facet = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["CO2"]]),
          selected = "Treatment",
          fixed = FALSE
        )
      ),
      ggplot2_args = ggplot2_args(
        labs = list(subtitle = "Plot generated by Bivariate Module")
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
```

```

}

# CDISC data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  ADSL <- rADSL
})
datanames(data) <- c("ADSL")
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(
    tm_g_bivariate(
      x = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["ADSL"]]),
          selected = "AGE",
          fixed = FALSE
        )
      ),
      y = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["ADSL"]]),
          selected = "SEX",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      row_facet = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["ADSL"]]),
          selected = "ARM",
          fixed = FALSE
        )
      ),
      col_facet = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["ADSL"]]),
          selected = "COUNTRY",
          fixed = FALSE
        )
      ),
    )
  )
),

```

```

        ggplot2_args = ggplot2_args(
            labs = list(subtitle = "Plot generated by Bivariate Module")
        )
    )
)
)
if (interactive()) {
    shinyApp(app$ui, app$server)
}

```

tm_g_distribution*teal module: Distribution analysis*

Description

Module is designed to explore the distribution of a single variable within a given dataset. It offers several tools, such as histograms, Q-Q plots, and various statistical tests to visually and statistically analyze the variable's distribution.

Usage

```

tm_g_distribution(
    label = "Distribution Module",
    dist_var,
    strata_var = NULL,
    group_var = NULL,
    freq = FALSE,
    ggtheme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void"),
    ggplot2_args = teal.widgets::ggplot2_args(),
    bins = c(30L, 1L, 100L),
    plot_height = c(600, 200, 2000),
    plot_width = NULL,
    pre_output = NULL,
    post_output = NULL
)

```

Arguments

<code>label</code>	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
<code>dist_var</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Variable(s) for which the distribution will be analyzed.
<code>strata_var</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Categorical variable used to split the distribution analysis.
<code>group_var</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Variable used for faceting plot into multiple panels.

<code>freq</code>	(logical) optional, whether to display frequency (TRUE) or density (FALSE). Defaults to density (FALSE).
<code>ggtheme</code>	(character) optional, ggplot2 theme to be used by default. Defaults to "gray".
<code>ggplot2_args</code>	(<code>ggplot2_args</code>) optional, object created by <code>teal.widgets::ggplot2_args()</code> with settings for all the plots or named list of <code>ggplot2_args</code> objects for plot-specific settings. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. List names should match the following: <code>c("default", "Histogram", "QQplot")</code> . For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code> .
<code>bins</code>	(<code>integer(1)</code> or <code>integer(3)</code>) optional, specifies the number of bins for the histogram. <ul style="list-style-type: none"> When the length of <code>bins</code> is one: The histogram bins will have a fixed size based on the <code>bins</code> provided. When the length of <code>bins</code> is three: The histogram bins are dynamically adjusted based on vector of <code>value</code>, <code>min</code>, and <code>max</code>. Defaults to <code>c(30L, 1L, 100L)</code>.
<code>plot_height</code>	(<code>numeric</code>) optional, specifies the plot height as a three-element vector of <code>value</code> , <code>min</code> , and <code>max</code> intended for use with a slider UI element.
<code>plot_width</code>	(<code>numeric</code>) optional, specifies the plot width as a three-element vector of <code>value</code> , <code>min</code> , and <code>max</code> for a slider encoding the plot width.
<code>pre_output</code>	(<code>shiny.tag</code> , optional) with text placed before the output to put the output into context. For example a title.
<code>post_output</code>	(<code>shiny.tag</code> , optional) with text placed after the output to put the output into context. For example the <code>shiny::helpText()</code> elements are useful.

Value

Object of class `teal_module` to be used in `teal` applications.

Examples

```
library(teal.widgets)

# general data example
data <- teal_data()
data <- within(data, {
  iris <- iris
})
datanames(data) <- "iris"

app <- init(
  data = data,
  modules = list(
    tm_g_distribution(
      dist_var = data_extract_spec(
```

```
    dataname = "iris",
    select = select_spec(variable_choices("iris"), "Petal.Length")
),
ggplot2_args = ggplot2_args(
    labs = list(subtitle = "Plot generated by Distribution Module")
)
)
)
)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

# CDISC data example
data <- teal_data()
data <- within(data, {
  ADSL <- rADSL
})
datanames(data) <- c("ADSL")
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

vars1 <- choices_selected(
  variable_choices(data[["ADSL"]], c("ARM", "COUNTRY", "SEX")),
  selected = NULL
)

app <- init(
  data = data,
  modules = modules(
    tm_g_distribution(
      dist_var = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          choices = variable_choices(data[["ADSL"]], c("AGE", "BMRKR1")),
          selected = "BMRKR1",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      strata_var = data_extract_spec(
        dataname = "ADSL",
        filter = filter_spec(
          vars = vars1,
          multiple = TRUE
        )
      ),
      group_var = data_extract_spec(
        dataname = "ADSL",
        filter = filter_spec(
          vars = vars1,
          multiple = TRUE
        )
      )
),
```

```

ggplot2_args = ggplot2_args(
  labs = list(subtitle = "Plot generated by Distribution Module")
)
)
)
)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_g_response *teal module: Response plot*

Description

Generates a response plot for a given response and x variables. This module allows users customize and add annotations to the plot depending on the module's arguments. It supports showing the counts grouped by other variable facets (by row / column), swapping the coordinates, show count annotations and displaying the response plot as frequency or density.

Usage

```

tm_g_response(
  label = "Response Plot",
  response,
  x,
  row_facet = NULL,
  col_facet = NULL,
  coord_flip = FALSE,
  count_labels = TRUE,
  rotate_xaxis_labels = FALSE,
  freq = FALSE,
  plot_height = c(600, 400, 5000),
  plot_width = NULL,
  ggtheme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void"),
  ggplot2_args = teal.widgets::ggplot2_args(),
  pre_output = NULL,
  post_output = NULL
)

```

Arguments

- | | |
|----------|---|
| label | (character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details. |
| response | (<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Which variable to use as the response. You can define one fixed column by setting <code>fixed = TRUE</code> inside the <code>select_spec</code> .
The <code>data_extract_spec</code> must not allow multiple selection in this case. |

x	(data_extract_spec or list of multiple data_extract_spec) Specifies which variable to use on the X-axis of the response plot. Allow the user to select multiple columns from the data allowed in teal. The data_extract_spec must not allow multiple selection in this case.
row_facet	(data_extract_spec or list of multiple data_extract_spec) optional specification of the data variable(s) to use for faceting rows.
col_facet	(data_extract_spec or list of multiple data_extract_spec) optional specification of the data variable(s) to use for faceting columns.
coord_flip	(logical(1)) Indicates whether to flip coordinates between x and response. The default value is FALSE and it will show the x variable on the x-axis and the response variable on the y-axis.
count_labels	(logical(1)) Indicates whether to show count labels. Defaults to TRUE.
rotate_xaxis_labels	(logical) optional, whether to rotate plot X axis labels. Does not rotate by default (FALSE).
freq	(logical(1)) Indicates whether to display frequency (TRUE) or density (FALSE). Defaults to density (FALSE).
plot_height	(numeric) optional, specifies the plot height as a three-element vector of value, min, and max intended for use with a slider UI element.
plot_width	(numeric) optional, specifies the plot width as a three-element vector of value, min, and max for a slider encoding the plot width.
ggtheme	(character) optional, ggplot2 theme to be used by default. Defaults to "gray".
ggplot2_args	(ggplot2_args) object created by <code>teal.widgets::ggplot2_args()</code> with settings for the module plot. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code>
pre_output	(shiny.tag) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
post_output	(shiny.tag) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.

Value

Object of class `teal_module` to be used in `teal` applications.

Note

For more examples, please see the vignette "Using response plot" via `vignette("using-response-plot", package = "teal.modules.general")`.

Examples

```

# general data example
library(teal.widgets)

data <- teal_data()
data <- within(data, {
  require(nestcolor)
  mtcars <- mtcars
  for (v in c("cyl", "vs", "am", "gear")) {
    mtcars[[v]] <- as.factor(mtcars[[v]])
  }
})
datanames(data) <- "mtcars"

app <- init(
  data = data,
  modules = modules(
    tm_g_response(
      label = "Response Plots",
      response = data_extract_spec(
        dataname = "mtcars",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["mtcars"]], c("cyl", "gear")),
          selected = "cyl",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      x = data_extract_spec(
        dataname = "mtcars",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["mtcars"]], c("vs", "am")),
          selected = "vs",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      ggplot2_args = ggplot2_args(
        labs = list(subtitle = "Plot generated by Response Module")
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

# CDISC data example
library(teal.widgets)

```

```

data <- teal_data()
data <- within(data, {
  require(nestcolor)
  ADSL <- rADSL
})
datanames(data) <- c("ADSL")
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(
    tm_g_response(
      label = "Response Plots",
      response = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["ADSL"]], c("BMRKR2", "COUNTRY")),
          selected = "BMRKR2",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      x = data_extract_spec(
        dataname = "ADSL",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["ADSL"]], c("SEX", "RACE")),
          selected = "RACE",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      ggplot2_args = ggplot2_args(
        labs = list(subtitle = "Plot generated by Response Module")
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_g_scatterplot teal module: Scatterplot

Description

Generates a customizable scatterplot using ggplot2. This module allows users to select variables for the x and y axes, color and size encodings, faceting options, and more. It supports log trans-

formations, trend line additions, and dynamic adjustments of point opacity and size through UI controls.

Usage

```
tm_g_scatterplot(
  label = "Scatterplot",
  x,
  y,
  color_by = NULL,
  size_by = NULL,
  row_facet = NULL,
  col_facet = NULL,
  plot_height = c(600, 200, 2000),
  plot_width = NULL,
  alpha = c(1, 0, 1),
  shape = shape_names,
  size = c(5, 1, 15),
  max_deg = 5L,
  rotate_xaxis_labels = FALSE,
  ggtheme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void"),
  pre_output = NULL,
  post_output = NULL,
  table_dec = 4,
  ggplot2_args = teal.widgets::ggplot2_args()
)
```

Arguments

<code>label</code>	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
<code>x</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Specifies variable names selected to plot along the x-axis by default.
<code>y</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Specifies variable names selected to plot along the y-axis by default.
<code>color_by</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, defines the color encoding. If <code>NULL</code> then no color encoding option will be displayed.
<code>size_by</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, defines the point size encoding. If <code>NULL</code> then no size encoding option will be displayed.
<code>row_facet</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, specifies the variable(s) for faceting rows.
<code>col_facet</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) optional, specifies the variable(s) for faceting columns.
<code>plot_height</code>	(<code>numeric</code>) optional, specifies the plot height as a three-element vector of value, <code>min</code> , and <code>max</code> intended for use with a slider UI element.

plot_width	(numeric) optional, specifies the plot width as a three-element vector of value, min, and max for a slider encoding the plot width.
alpha	(integer(1) or integer(3)) optional, specifies point opacity. <ul style="list-style-type: none"> • When the length of alpha is one: the plot points will have a fixed opacity. • When the length of alpha is three: the plot points opacity are dynamically adjusted based on vector of value, min, and max.
shape	(character) optional, character vector with the names of the shape, e.g. c("triangle", "square", "circle"). It defaults to shape_names. This is a complete list from vignette("ggplot2-specs", package="ggplot2").
size	(integer(1) or integer(3)) optional, specifies point size. <ul style="list-style-type: none"> • When the length of size is one: the plot point sizes will have a fixed size. • When the length of size is three: the plot points size are dynamically adjusted based on vector of value, min, and max.
max_deg	(integer) optional, maximum degree for the polynomial trend line. Must not be less than 1.
rotate_xaxis_labels	(logical) optional, whether to rotate plot X axis labels. Does not rotate by default (FALSE).
ggtheme	(character) optional, ggplot2 theme to be used by default. Defaults to "gray".
pre_output	(shiny.tag) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
post_output	(shiny.tag) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like shiny::helpText() are useful.
table_dec	(integer) optional, number of decimal places used to round numeric values in the table.
ggplot2_args	(ggplot2_args) object created by <code>teal.widgets::ggplot2_args()</code> with settings for the module plot. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code>

Value

Object of class `teal_module` to be used in `teal` applications.

Note

For more examples, please see the vignette "Using scatterplot" via `vignette("using-scatterplot", package = "teal.modules.general")`.

Examples

```

library(teal.widgets)

# general data example
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  CO2 <- CO2
})
datanames(data) <- "CO2"

app <- init(
  data = data,
  modules = modules(
    tm_g_scatterplot(
      label = "Scatterplot Choices",
      x = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["CO2"]], c("conc", "uptake")),
          selected = "conc",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      y = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["CO2"]], c("conc", "uptake")),
          selected = "uptake",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      color_by = data_extract_spec(
        dataname = "CO2",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(
            data[["CO2"]],
            c("Plant", "Type", "Treatment", "conc", "uptake")
          ),
          selected = NULL,
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      size_by = data_extract_spec(
        dataname = "CO2",
        select = select_spec(

```

```
label = "Select variable:",
choices = variable_choices(data[["CO2"]], c("conc", "uptake")),
selected = "uptake",
multiple = FALSE,
fixed = FALSE
)
),
row_facet = data_extract_spec(
dataname = "CO2",
select = select_spec(
label = "Select variable:",
choices = variable_choices(data[["CO2"]], c("Plant", "Type", "Treatment")),
selected = NULL,
multiple = FALSE,
fixed = FALSE
)
),
col_facet = data_extract_spec(
dataname = "CO2",
select = select_spec(
label = "Select variable:",
choices = variable_choices(data[["CO2"]], c("Plant", "Type", "Treatment")),
selected = NULL,
multiple = FALSE,
fixed = FALSE
)
),
ggplot2_args = ggplot2_args(
labs = list(subtitle = "Plot generated by Scatterplot Module")
)
)
)
)
)
if (interactive()) {
shinyApp(app$ui, app$server)
}

# CDISC data example
data <- teal_data()
data <- within(data, {
require(nestcolor)
ADSL <- rADSL
})
datanames(data) <- c("ADSL")
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
data = data,
modules = modules(
tm_g_scatterplot(
label = "Scatterplot Choices",
x = data_extract_spec(
dataname = "ADSL",
```

```

select = select_spec(
  label = "Select variable:",
  choices = variable_choices(data[["ADSL"]], c("AGE", "BMRKR1", "BMRKR2")),
  selected = "AGE",
  multiple = FALSE,
  fixed = FALSE
)
),
y = data_extract_spec(
  dataname = "ADSL",
  select = select_spec(
    label = "Select variable:",
    choices = variable_choices(data[["ADSL"]], c("AGE", "BMRKR1", "BMRKR2")),
    selected = "BMRKR1",
    multiple = FALSE,
    fixed = FALSE
)
),
color_by = data_extract_spec(
  dataname = "ADSL",
  select = select_spec(
    label = "Select variable:",
    choices = variable_choices(
      data[["ADSL"]],
      c("AGE", "BMRKR1", "BMRKR2", "RACE", "REGION1")
    ),
    selected = NULL,
    multiple = FALSE,
    fixed = FALSE
)
),
size_by = data_extract_spec(
  dataname = "ADSL",
  select = select_spec(
    label = "Select variable:",
    choices = variable_choices(data[["ADSL"]], c("AGE", "BMRKR1")),
    selected = "AGE",
    multiple = FALSE,
    fixed = FALSE
)
),
row_facet = data_extract_spec(
  dataname = "ADSL",
  select = select_spec(
    label = "Select variable:",
    choices = variable_choices(data[["ADSL"]], c("BMRKR2", "RACE", "REGION1")),
    selected = NULL,
    multiple = FALSE,
    fixed = FALSE
)
),
col_facet = data_extract_spec(
  dataname = "ADSL",

```

```

    select = select_spec(
      label = "Select variable:",
      choices = variable_choices(data[["ADSL"]], c("BMRKR2", "RACE", "REGION1")),
      selected = NULL,
      multiple = FALSE,
      fixed = FALSE
    )
  ),
  ggplot2_args = ggplot2_args(
    labs = list(subtitle = "Plot generated by Scatterplot Module")
  )
)
)
)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_g_scatterplotmatrix*teal module: Scatterplot matrix***Description**

Generates a scatterplot matrix from selected variables from datasets. Each plot within the matrix represents the relationship between two variables, providing the overview of correlations and distributions across selected data.

Usage

```

tm_g_scatterplotmatrix(
  label = "Scatterplot Matrix",
  variables,
  plot_height = c(600, 200, 2000),
  plot_width = NULL,
  pre_output = NULL,
  post_output = NULL
)

```

Arguments

- | | |
|------------------|---|
| label | (character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details. |
| variables | (<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Specifies plotting variables from an incoming dataset with filtering and selecting. In case of <code>data_extract_spec</code> use <code>select_spec(..., ordered = TRUE)</code> if plot elements should be rendered according to selection order. |

<code>plot_height</code>	(<code>numeric</code>) optional, specifies the plot height as a three-element vector of value, <code>min</code> , and <code>max</code> intended for use with a slider UI element.
<code>plot_width</code>	(<code>numeric</code>) optional, specifies the plot width as a three-element vector of value, <code>min</code> , and <code>max</code> for a slider encoding the plot width.
<code>pre_output</code>	(<code>shiny.tag</code>) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
<code>post_output</code>	(<code>shiny.tag</code>) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.

Value

Object of class `teal_module` to be used in `teal` applications.

Note

For more examples, please see the vignette "Using scatterplot matrix" via `vignette("using-scatterplot-matrix", package = "teal.modules.general")`.

Examples

```
# general data example
data <- teal_data()
data <- within(data, {
  countries <- data.frame(
    id = c("DE", "FR", "IT", "ES", "PT", "GR", "NL", "BE", "LU", "AT"),
    government = factor(
      c(2, 2, 2, 1, 2, 2, 1, 1, 1, 2),
      labels = c("Monarchy", "Republic")
    ),
    language_family = factor(
      c(1, 3, 3, 3, 2, 1, 1, 3, 1),
      labels = c("Germanic", "Hellenic", "Romance")
    ),
    population = c(83, 67, 60, 47, 10, 11, 17, 11, 0.6, 9),
    area = c(357, 551, 301, 505, 92, 132, 41, 30, 2.6, 83),
    gdp = c(3.4, 2.7, 2.1, 1.4, 0.3, 0.2, 0.7, 0.5, 0.1, 0.4),
    debt = c(2.1, 2.3, 2.4, 2.6, 2.3, 2.4, 2.3, 2.4, 2.4)
  )
  sales <- data.frame(
    id = 1:50,
    country_id = sample(
      c("DE", "FR", "IT", "ES", "PT", "GR", "NL", "BE", "LU", "AT"),
      size = 50,
      replace = TRUE
    ),
    year = sort(sample(2010:2020, 50, replace = TRUE)),
    venue = sample(c("small", "medium", "large", "online"), 50, replace = TRUE),
    cancelled = sample(c(TRUE, FALSE), 50, replace = TRUE),
    quantity = rnorm(50, 100, 20),
  )
})
```

```
costs = rnorm(50, 80, 20),
profit = rnorm(50, 20, 10)
)
})
datanames(data) <- c("countries", "sales")
join_keys(data) <- join_keys(
  join_key("countries", "countries", "id"),
  join_key("sales", "sales", "id"),
  join_key("countries", "sales", c("id" = "country_id")))
)

app <- init(
  data = data,
  modules = modules(
    tm_g_scatterplotmatrix(
      label = "Scatterplot matrix",
      variables = list(
        data_extract_spec(
          dataname = "countries",
          select = select_spec(
            label = "Select variables:",
            choices = variable_choices(data[["countries"]]),
            selected = c("area", "gdp", "debt"),
            multiple = TRUE,
            ordered = TRUE,
            fixed = FALSE
          )
        ),
        data_extract_spec(
          dataname = "sales",
          filter = filter_spec(
            label = "Select variable:",
            vars = "country_id",
            choices = value_choices(data[["sales"]], "country_id"),
            selected = c("DE", "FR", "IT", "ES", "PT", "GR", "NL", "BE", "LU", "AT"),
            multiple = TRUE
          ),
          select = select_spec(
            label = "Select variables:",
            choices = variable_choices(data[["sales"]], c("quantity", "costs", "profit")),
            selected = c("quantity", "costs", "profit"),
            multiple = TRUE,
            ordered = TRUE,
            fixed = FALSE
          )
        )
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}
```

```

# CDISC data example
data <- teal_data()
data <- within(data, {
  ADSL <- rADSL
  ADRS <- rADRS
})
datanames(data) <- c("ADSL", "ADRS")
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(
    tm_g_scatterplotmatrix(
      label = "Scatterplot matrix",
      variables = list(
        data_extract_spec(
          dataname = "ADSL",
          select = select_spec(
            label = "Select variables:",
            choices = variable_choices(data[["ADSL"]]),
            selected = c("AGE", "RACE", "SEX"),
            multiple = TRUE,
            ordered = TRUE,
            fixed = FALSE
          )
        ),
        data_extract_spec(
          dataname = "ADRS",
          filter = filter_spec(
            label = "Select endpoints:",
            vars = c("PARAMCD", "AVISIT"),
            choices = value_choices(data[["ADRS"]], c("PARAMCD", "AVISIT"), c("PARAM", "AVISIT")),
            selected = "INVET - END OF INDUCTION",
            multiple = TRUE
          ),
          select = select_spec(
            label = "Select variables:",
            choices = variable_choices(data[["ADRS"]]),
            selected = c("AGE", "AVAL", "ADY"),
            multiple = TRUE,
            ordered = TRUE,
            fixed = FALSE
          )
        )
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_missing_data teal module: Missing data analysis

Description

This module analyzes missing data in `data.frames` to help users explore missing observations and gain insights into the completeness of their data. It is useful for clinical data analysis within the context of CDISC standards and adaptable for general data analysis purposes.

Usage

```
tm_missing_data(  
  label = "Missing data",  
  plot_height = c(600, 400, 5000),  
  plot_width = NULL,  
  parent_dataname = "ADSL",  
  ggtheme = c("classic", "gray", "bw", "linedraw", "light", "dark", "minimal", "void"),  
  ggplot2_args = list(`Combinations Hist` = teal.widgets::ggplot2_args(labs =  
    list(caption = NULL)), `Combinations Main` = teal.widgets::ggplot2_args(labs =  
    list(title = NULL))),  
  pre_output = NULL,  
  post_output = NULL  
)
```

Arguments

<code>label</code>	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
<code>plot_height</code>	(numeric) optional, specifies the plot height as a three-element vector of value, min, and max intended for use with a slider UI element.
<code>plot_width</code>	(numeric) optional, specifies the plot width as a three-element vector of value, min, and max for a slider encoding the plot width.
<code>parent_dataname</code>	(character(1)) Specifies the parent dataset name. Default is ADSL for CDISC data. If provided and exists, enables additional analysis "by subject". For non-CDISC data, this parameter can be ignored.
<code>ggtheme</code>	(character) optional, specifies the default ggplot2 theme for plots. Defaults to <code>classic</code> .
<code>ggplot2_args</code>	(<code>ggplot2_args</code>) optional, object created by <code>teal.widgets::ggplot2_args()</code> with settings for all the plots or named list of <code>ggplot2_args</code> objects for plot-specific settings. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. List names should match the following: <code>c("default", "Summary Obs", "Summary Patients", "Combinations Main", "Combinations Hist", "By Subject")</code> . For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code> .

<code>pre_output</code>	(shiny.tag) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
<code>post_output</code>	(shiny.tag) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.

Value

Object of class `teal_module` to be used in `teal` applications.

Examples

```
library(teal.widgets)

# module specification used in apps below
tm_missing_data_module <- tm_missing_data(
  ggplot2_args = list(
    "Combinations Hist" = ggplot2_args(
      labs = list(subtitle = "Plot produced by Missing Data Module", caption = NULL)
    ),
    "Combinations Main" = ggplot2_args(labs = list(title = NULL))
  )
)

# general example data
data <- teal_data()
data <- within(data, {
  require(nestcolor)

  add_nas <- function(x) {
    x[sample(seq_along(x), floor(length(x) * runif(1, .05, .17)))] <- NA
    x
  }

  iris <- iris
  mtcars <- mtcars

  iris[] <- lapply(iris, add_nas)
  mtcars[] <- lapply(mtcars, add_nas)
  mtcars[["cyl"]] <- as.factor(mtcars[["cyl"]])
  mtcars[["gear"]] <- as.factor(mtcars[["gear"]])
})
data$names(data) <- c("iris", "mtcars")

app <- init(
  data = data,
  modules = modules(tm_missing_data_module)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}
```

```

# CDISC example data
data <- teal_data()
data <- within(data, {
  require(nestcolor)
  ADSL <- rADSL
  ADRS <- rADRS
})
datanames(data) <- c("ADSL", "ADRS")
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(tm_missing_data_module)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_outliers*teal module: Outliers analysis***Description**

Module to analyze and identify outliers using different methods such as IQR, Z-score, and Percentiles, and offers visualizations including box plots, density plots, and cumulative distribution plots to help interpret the outliers.

Usage

```

tm_outliers(
  label = "Outliers Module",
  outlier_var,
  categorical_var = NULL,
  ggtheme = c("gray", "bw", "linedraw", "light", "dark", "minimal", "classic", "void"),
  ggplot2_args = teal.widgets::ggplot2_args(),
  plot_height = c(600, 200, 2000),
  plot_width = NULL,
  pre_output = NULL,
  post_output = NULL
)

```

Arguments

<code>label</code>	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
<code>outlier_var</code>	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Specifies variable(s) to be analyzed for outliers.

categorical_var	(<i>data_extract_spec</i> or list of multiple <i>data_extract_spec</i>) optional, specifies the categorical variable(s) to split the selected outlier variables on.
ggtheme	(character) optional, <i>ggplot2</i> theme to be used by default. Defaults to "gray".
ggplot2_args	(<i>ggplot2_args</i>) optional, object created by <code>teal.widgets::ggplot2_args()</code> with settings for all the plots or named list of <i>ggplot2_args</i> objects for plot-specific settings. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. List names should match the following: <code>c("default", "Boxplot", "Density Plot", "Cumulative Distribution Plot")</code> . For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code> .
plot_height	(numeric) optional, specifies the plot height as a three-element vector of value, <i>min</i> , and <i>max</i> intended for use with a slider UI element.
plot_width	(numeric) optional, specifies the plot width as a three-element vector of value, <i>min</i> , and <i>max</i> for a slider encoding the plot width.
pre_output	(shiny.tag) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
post_output	(shiny.tag) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.

Value

Object of class `teal_module` to be used in `teal` applications.

Examples

```
library(teal.widgets)

# general data example
data <- teal_data()
data <- within(data, {
  C02 <- C02
  C02[["primary_key"]] <- seq_len(nrow(C02))
})
datanames(data) <- "C02"
join_keys(data) <- join_keys(join_key("C02", "C02", "primary_key"))

vars <- choices_selected(variable_choices(data[["C02"]], c("Plant", "Type", "Treatment")))

app <- init(
  data = data,
  modules = modules(
    tm_outliers(
      outlier_var = list(
        data_extract_spec(
          dataname = "C02",

```

```
select = select_spec(
  label = "Select variable:",
  choices = variable_choices(data[["CO2"]], c("conc", "uptake")),
  selected = "uptake",
  multiple = FALSE,
  fixed = FALSE
)
)
),
categorical_var = list(
  data_extract_spec(
    dataname = "CO2",
    filter = filter_spec(
      vars = vars,
      choices = value_choices(data[["CO2"]], vars$selected),
      selected = value_choices(data[["CO2"]], vars$selected),
      multiple = TRUE
    )
  )
),
ggplot2_args = list(
  ggplot2_args(
    labs = list(subtitle = "Plot generated by Outliers Module")
  )
)
)
)
)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

# CDISC data example
data <- teal_data()
data <- within(data, {
  ADSL <- rADSL
})
datanames(data) <- "ADSL"
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

fact_vars_adsl <- names(Filter(isTRUE, sapply(data[["ADSL"]], is.factor)))
vars <- choices_selected(variable_choices(data[["ADSL"]], fact_vars_adsl))

app <- init(
  data = data,
  modules = modules(
    tm_outliers(
      outlier_var = list(
        data_extract_spec(
          dataname = "ADSL",
          select = select_spec(
            label = "Select variable:",
            choices = variable_choices(data[["ADSL"]], c("AGE", "BMRKR1"))
          )
        )
      )
    )
  )
)
```

```

selected = "AGE",
multiple = FALSE,
fixed = FALSE
)
)
),
categorical_var = list(
  data_extract_spec(
    dataname = "ADSL",
    filter = filter_spec(
      vars = vars,
      choices = value_choices(data[["ADSL"]], vars$selected),
      selected = value_choices(data[["ADSL"]], vars$selected),
      multiple = TRUE
    )
  )
),
ggplot2_args = list(
  ggplot2_args(
    labs = list(subtitle = "Plot generated by Outliers Module")
  )
)
)
)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

tm_t_crosstable *teal module: Cross-table*

Description

Generates a simple cross-table of two variables from a dataset with custom options for showing percentages and sub-totals.

Usage

```

tm_t_crosstable(
  label = "Cross Table",
  x,
  y,
  show_percentage = TRUE,
  show_total = TRUE,
  pre_output = NULL,
  post_output = NULL,
  basic_table_args = teal.widgets::basic_table_args()
)

```

Arguments

label	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
x	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Object with all available choices with pre-selected option for variable X - row values. In case of <code>data_extract_spec</code> use <code>select_spec(..., ordered = TRUE)</code> if table elements should be rendered according to selection order.
y	(<code>data_extract_spec</code> or list of multiple <code>data_extract_spec</code>) Object with all available choices with pre-selected option for variable Y - column values. <code>data_extract_spec</code> must not allow multiple selection in this case.
show_percentage	(logical(1)) Indicates whether to show percentages (relevant only when x is a factor). Defaults to TRUE.
show_total	(logical(1)) Indicates whether to show total column. Defaults to TRUE.
pre_output	(shiny.tag) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
post_output	(shiny.tag) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.
basic_table_args	(<code>basic_table_args</code>) object created by <code>teal.widgets::basic_table_args()</code> with settings for the module table. The argument is merged with options variable <code>teal.basic_table_args</code> and default module setup. For more details see the vignette: <code>vignette("custom-basic-table-arguments", package = "teal.widgets")</code>

Value

Object of class `teal_module` to be used in `teal` applications.

Note

For more examples, please see the vignette "Using cross table" via `vignette("using-cross-table", package = "teal.modules.general")`.

Examples

```
# general data example
library(teal.widgets)

data <- teal_data()
data <- within(data, {
  mtcars <- mtcars
  for (v in c("cyl", "vs", "am", "gear")) {
    mtcars[[v]] <- as.factor(mtcars[[v]])
  }
})
```

```

mtcars[["primary_key"]] <- seq_len(nrow(mtcars))
})
datanames(data) <- "mtcars"
join_keys(data) <- join_keys(join_key("mtcars", "mtcars", "primary_key"))

app <- init(
  data = data,
  modules = modules(
    tm_t_crosstable(
      label = "Cross Table",
      x = data_extract_spec(
        dataname = "mtcars",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["mtcars"]], c("cyl", "vs", "am", "gear")),
          selected = c("cyl", "gear"),
          multiple = TRUE,
          ordered = TRUE,
          fixed = FALSE
        )
      ),
      y = data_extract_spec(
        dataname = "mtcars",
        select = select_spec(
          label = "Select variable:",
          choices = variable_choices(data[["mtcars"]], c("cyl", "vs", "am", "gear")),
          selected = "vs",
          multiple = FALSE,
          fixed = FALSE
        )
      ),
      basic_table_args = basic_table_args(
        subtitles = "Table generated by Crosstable Module"
      )
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

# CDISC data example
library(teal.widgets)

data <- teal_data()
data <- within(data, {
  ADSL <- rADSL
})
datanames(data) <- "ADSL"
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,

```

```

modules = modules(
  tm_t_crosstable(
    label = "Cross Table",
    x = data_extract_spec(
      dataname = "ADSL",
      select = select_spec(
        label = "Select variable:",
        choices = variable_choices(data[["ADSL"]], subset = function(data) {
          idx <- !vapply(data, inherits, logical(1), c("Date", "POSIXct", "POSIXlt"))
          return(names(data)[idx])
        }),
        selected = "COUNTRY",
        multiple = TRUE,
        ordered = TRUE,
        fixed = FALSE
      )
    ),
    y = data_extract_spec(
      dataname = "ADSL",
      select = select_spec(
        label = "Select variable:",
        choices = variable_choices(data[["ADSL"]], subset = function(data) {
          idx <- vapply(data, is.factor, logical(1))
          return(names(data)[idx])
        }),
        selected = "SEX",
        multiple = FALSE,
        fixed = FALSE
      )
    ),
    basic_table_args = basic_table_args(
      subtitles = "Table generated by Crosstable Module"
    )
  )
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

```

`tm_variable_browser` teal module: *Variable browser*

Description

Module provides provides a detailed summary and visualization of variable distributions for `data.frame` objects, with interactive features to customize analysis.

Usage

```
tm_variable_browser(
  label = "Variable Browser",
  datasets_selected = character(0),
  parent_dataname = "ADSL",
  pre_output = NULL,
  post_output = NULL,
  ggplot2_args = teal.widgets::ggplot2_args()
)
```

Arguments

<code>label</code>	(character(1)) Label shown in the navigation item for the module or module group. For <code>modules()</code> defaults to "root". See Details.
<code>datasets_selected</code>	(character) vector of datasets which should be shown, in order. Names must correspond with datasets names. If vector of length zero (default) then all datasets are shown. Note: Only <code>data.frame</code> objects are compatible; using other types will cause an error.
<code>parent_dataname</code>	(character(1)) string specifying a parent dataset. If it exists in <code>datasets_selected</code> then an extra checkbox will be shown to allow users to not show variables in other datasets which exist in this dataname. This is typically used to remove ADSL columns in CDISC data. In non CDISC data this can be ignored. Defaults to "ADSL".
<code>pre_output</code>	(shiny.tag) optional, text or UI element to be displayed before the module's output, providing context or a title. with text placed before the output to put the output into context. For example a title.
<code>post_output</code>	(shiny.tag) optional, text or UI element to be displayed after the module's output, adding context or further instructions. Elements like <code>shiny::helpText()</code> are useful.
<code>ggplot2_args</code>	(<code>ggplot2_args</code>) object created by <code>teal.widgets::ggplot2_args()</code> with settings for the module plot. The argument is merged with options variable <code>teal.ggplot2_args</code> and default module setup. For more details see the vignette: <code>vignette("custom-ggplot2-arguments", package = "teal.widgets")</code>

Details

Numeric columns with fewer than 30 distinct values can be treated as either discrete or continuous with a checkbox allowing users to switch how they are treated(if < 6 unique values then the default is discrete, otherwise it is continuous).

Value

Object of class `teal_module` to be used in `teal` applications.

Examples

```
library(teal.widgets)

# Module specification used in apps below
tm_variable_browser_module <- tm_variable_browser(
  label = "Variable browser",
  ggplot2_args = ggplot2_args(
    labs = list(subtitle = "Plot generated by Variable Browser Module")
  )
)

# general data example
data <- teal_data()
data <- within(data, {
  iris <- iris
  mtcars <- mtcars
  women <- women
  faithful <- faithful
  CO2 <- CO2
})
datanames(data) <- c("iris", "mtcars", "women", "faithful", "CO2")

app <- init(
  data = data,
  modules = modules(tm_variable_browser_module)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}

# CDISC example data
data <- teal_data()
data <- within(data, {
  ADSL <- rADSL
  ADTTE <- rADTTE
})
datanames(data) <- c("ADSL", "ADTTE")
join_keys(data) <- default_cdisc_join_keys[datanames(data)]

app <- init(
  data = data,
  modules = modules(tm_variable_browser_module)
)
if (interactive()) {
  shinyApp(app$ui, app$server)
}
```

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