

# Package ‘shinyCohortBuilder’

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**Type** Package

**Title** Modular Cohort-Building Framework for Analytical Dashboards

**Version** 0.4.0

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**Description** You can easily add advanced cohort-building component to your analytical dashboard or simple 'Shiny' app. Then you can instantly start building cohorts using multiple filters of different types, filtering datasets, and filtering steps. Filters can be complex and data-specific, and together with multiple filtering steps you can use complex filtering rules. The cohort-building sidebar panel allows you to easily work with filters, add and remove filtering steps. It helps you with handling missing values during filtering, and provides instant filtering feedback with filter feedback plots. The GUI panel is not only compatible with native shiny bookmarking, but also provides reproducible R code.

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**Depends** R (>= 3.5.0)

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shinyCohortBuilder-package  
*GUI layer for cohortBuilder package*

---

### Description

GUI layer for cohortBuilder package

---

`.cb_input`                      *Create input controller insensitive to server updates*

---

### Description

Input controllers created with `'cb_input'` are sending its value to server only when user changes it's value directly in browser. That means all the `'update*'` functions have only visible effect on application output.

The method should be used for each filter input controller and precise which filter value should be updated when the input selection is changes.

### Usage

```
.cb_input(ui, data_param, ..., priority = NULL)
```

### Arguments

<code>ui</code>	UI defining input controllers.
<code>data_param</code>	Name of the parameter that should be updated in filter whenever user change the input value.
<code>...</code>	Extra attributes passed to the input div container.
<code>priority</code>	Set to <code>'event'</code> to force sending value.

### Value

A `'shiny.tag'` object defining html structure of filter input container.

### Examples

```
if (interactive()) {  
  library(shiny)  
  library(shinyCohortBuilder)  
  
  shiny::addResourcePath(  
    "shinyCohortBuilder",  
    system.file("www", package = "shinyCohortBuilder")  
  )  
}
```

```

ui <- fluidPage(
  tags$head(
    shiny::tags$script(type = "text/javascript", src = file.path("shinyCohortBuilder", "scb.js"))
  ),
  actionButton("update", "Update with random value"),
  div(
    class = "cb_container",
    `data-ns_prefix` = "",
    div(
      class = "cb_step",
      `data-step_id` = "1",
      div(
        class = "cb_filter",
        `data-filter_id` = "filid",
        .cb_input(
          numericInput("val", "Value", value = 1),
          data_param = "range"
        )
      )
    )
  )
)

server <- function(input, output, session) {
  observeEvent(input$action, {
    # print should be avoided when value is changed due to update
    print(input$action)
  })
  observeEvent(input$update, {
    updateNumericInput(session, "val", value = rnorm(1))
  })
}

shinyApp(ui, server)
}

```

---

*.render\_filter*
*Define filter related output in filtering panel*


---

### **Description**

The method exported only for custom extensions use.

### **Usage**

```
.render_filter(filter, step_id, cohort, ns)
```

**Arguments**

<code>filter</code>	Filter object.
<code>step_id</code>	Id of the step.
<code>cohort</code>	Cohort object.
<code>ns</code>	Namespace function.

**Value**

A ‘shiny.tag’ class ‘div’ object defining html structure of filter input panel.

**Examples**

```

if (interactive()) {
  library(magrittr)
  library(shiny)
  library(cohortBuilder)
  library(shinyCohortBuilder)

  ui <- fluidPage(
    actionButton("add_filter", "Add Filter"),
    div(id = "filter_container")
  )

  server <- function(input, output, session) {
    add_gui_filter_layer <- function(public, private, ...) {
      private$steps[["1"]]$filters$copies$gui <- .gui_filter(
        private$steps[["1"]]$filters$copies
      )
    }
    add_hook("post_cohort_hook", add_gui_filter_layer)
    coh <- cohort(
      set_source(as.tbllist(librarian)),
      filter(
        "range", id = "copies", name = "Copies", dataset = "books",
        variable = "copies", range = c(5, 12)
      )
    ) %>% run()
    coh$attributes$session <- session
    coh$attributes$feedback <- TRUE

    observeEvent(input$add_filter, {
      insertUI(
        "#filter_container",
        ui = .render_filter(
          coh$get_filter("1", "copies"),
          step_id = "1",
          cohort = coh,
          ns = function(x) x
        )
      ), ignoreInit = TRUE, once = TRUE)
    }
  }

```

```

  shinyApp(ui, server)
}

```

---

.save_observer	<i>Save observer to user session</i>
----------------	--------------------------------------

---

### Description

The method used to store created observers (used to implement extra filter logic). The saved observers are then destroyed when filtering step is removed which prevents duplicated execution of accumulated observers.

### Usage

```
.save_observer(observer, id, session)
```

### Arguments

observer	An 'observe' or 'observeEvent' to be saved.
id	Id of the observer. Preferably prefixed with step_id. The saved observer is saved as 'session\$userData\$observers[['<id>-observer']]' object.
session	Shiny session object.

### Value

No return value, used for side effect which is saving the observer to 'session\$userData' object.

### Examples

```

if (interactive()) {
  library(shiny)
  library(shinyCohortBuilder)

  ui <- fluidPage(
    numericInput("power", "Power", min = 0, max = 10, value = 1, step = 1),
    numericInput("value", "Value", min = 0, max = 100, value = 2, step = 0.1),
    actionButton("add", "Observe the selected power"),
    actionButton("rm", "Stop observing the selected power")
  )

  server <- function(input, output, session) {
    observeEvent(input$add, {
      .save_observer(
        observeEvent(input$value, {
          print(input$value ^ input$power)
        }),
        as.character(input$power),
        session = session
      )
    })
  }
}

```

```
    )
  }, ignoreInit = TRUE)

  observeEvent(input$rm, {
    id <- paste0(input$power, "-observer")
    session$userData$observers[[id]]$destroy()
    session$userData$observers[[id]] <- NULL
  }, ignoreInit = TRUE)
}

shinyApp(ui, server)
}
```

---

`.sendOutput`*Send output rendering*

---

### Description

Functional approach to rendering output. Equivalent of `'output[[name]] <- rendering'`.

### Usage

```
.sendOutput(name, rendering, session = shiny::getDefaultReactiveDomain())
```

### Arguments

<code>name</code>	Name of the output to be rendered
<code>rendering</code>	Rendering expression to be sent.
<code>session</code>	Shiny session object.

### Value

No return value, used for side effect which is assigning rendering to the output object.

### Examples

```
if (interactive()) {
  library(shiny)
  library(shinyCohortBuilder)

  rendering <- function(x_max) {
    renderPlot({
      x <- seq(0, x_max, by = 0.01)
      plot(x, sin(x), type = "l")
    })
  }

  ui <- fluidPage(
    numericInput("xmax", "X Axis Limit", min = 0, max = 10, value = pi),
```

```
    plotOutput("out")
  )

  server <- function(input, output, session) {
    observeEvent(input$xmax, {
      .sendOutput("out", rendering(input$xmax))
    })
  }

  shinyApp(ui, server)
}
```

---

autofilter

*Generate filters definition based on the Source data*

---

## Description

The method should analyze source data structure, generate proper filters based on the data (e.g. column types) and attach them to source.

## Usage

```
autofilter(source, attach_as = c("step", "meta"), ...)

## Default S3 method:
autofilter(source, ...)

## S3 method for class 'tblist'
autofilter(source, attach_as = c("step", "meta"), ...)
```

## Arguments

source	Source object.
attach_as	Choose whether the filters should be attached as a new step, or list of available filters (used in filtering panel when 'new_step = "configure"'). By default in step.
...	Extra arguments passed to a specific method.

## Value

Source object having step configuration attached.

## See Also

[source-gui-layer](#)

## Examples

```
library(magrittr)
library(cohortBuilder)
library(shinyCohortBuilder)

iris_source <- set_source(tblist(iris = iris)) %>%
  autofilter()
iris_cohort <- cohort(iris_source)
sum_up(iris_cohort)

if (interactive()) {
  library(shiny)

  ui <- fluidPage(
    cb_ui("mycoh")
  )

  server <- function(input, output, session) {
    cb_server("mycoh", cohort = iris_cohort)
  }

  shinyApp(ui, server)
}
```

---

available-filters-choices

*Generate available filters choices based on the Source data*

---

## Description

The method should return the available choices for virtualSelect input.

## Usage

```
.available_filters_choices(source, cohort, ...)
```

## Default S3 method:

```
.available_filters_choices(source, cohort, ...)
```

## S3 method for class 'tblist'

```
.available_filters_choices(source, cohort, ...)
```

## Arguments

source	Source object.
cohort	cohortBuilder cohort object
...	Extra arguments passed to a specific method.

**Value**

‘shinyWidgets::prepare\_choices’ output value.

**Examples**

```

if (interactive()) {
  library(magrittr)
  library(shiny)
  library(cohortBuilder)
  library(shinyCohortBuilder)
  library(shinyWidgets)

  coh <- cohort(
    set_source(as.tbllist(librarian), available_filters = list(
      filter(
        "range", id = "copies", name = "Copies", dataset = "books",
        variable = "copies", range = c(5, 12)
      ),
      filter(
        "date_range", id = "registered", name = "Registered", dataset = "borrowers",
        variable = "registered", range = c(as.Date("2010-01-01"), Inf)
      )
    ))
  ) %>% run()
  filter_choices <- .available_filters_choices(coh$get_source(), coh)

  ui <- fluidPage(
    virtualSelectInput("filters", "Filters", choices = filter_choices, html = TRUE)
  )

  server <- function(input, output, session) {

  }

  shinyApp(ui, server)
}

```

---

cb\_changed

*Track changes of cohort data in Shiny*


---

**Description**

The function returns Shiny input object related to selected cohort that is triggered whenever cohort data filters were applied to it within filtering panel.

**Usage**

```
cb_changed(session, cohort_id, step_id = NULL)
```

## Arguments

session	Shiny session object.
cohort_id	Id of the cohort.
step_id	Id of the step to check. When NULL (default) all the steps are checked for changes.

## Details

The function is meant to be used as a trigger for Shiny render functions and observers.

---

cb_ui	<i>Include filtering panel in Shiny</i>
-------	---

---

## Description

The function returns filtering panel placeholder, you may use in you custom Shiny application. Use in the UI part of your application.

## Usage

```
cb_ui(  
  id,  
  ...,  
  state = FALSE,  
  steps = TRUE,  
  code = TRUE,  
  attrition = TRUE,  
  new_step = c("clone", "configure"),  
  manage_step = FALSE  
)  
  
cb_server(  
  id,  
  cohort,  
  run_button = "none",  
  stats = c("pre", "post"),  
  feedback = FALSE,  
  enable_bookmarking = shiny::getShinyOption("bookmarkStore", default = "disable"),  
  show_help = TRUE,  
  ...  
)
```

**Arguments**

id	Id of the module used to render the panel.
...	Extra attributes passed to the panel div container.
state	Set to TRUE (default) to enable get/set state panel.
steps	Set to TRUE (default) if multiple steps should be available.
code	Set to TRUE (default) to enable reproducible code panel.
attrition	Set to TRUE (default) to enable attrition plot panel.
new_step	Choose which add step method should be used for creating new step. Possible options are: "clone" - copy filters from last step, "configure" - opening modal and allow to chose filters from available filters.
manage_step	When 'TRUE', enables feature, that allows to modify the latest step filters (add/remove them). Available list of filters used by the feature should be stores as 'source\$attributes\$available_filters' object.
cohort	Cohort object storing filtering steps configuration.
run_button	Should Run button be displayed? If so, the current step computations are run only when clicked. Three options are available "none" - no button, "local" - button displayed at each step panel, "global" - button visible in top filtering panel.
stats	Choose which statistics should be displayed for data (and some filters). Possible options are: "pre" - previous step stat, "post" - current step stats, 'c("pre", "post")' - for both and NULL for no stats.
feedback	Set to TRUE (default) if feedback plots should be displayed at each filter.
enable_bookmarking	Set to TRUE (default) if panel should be compatible with native shiny bookmarking.
show_help	Set to TRUE (default) to enable help buttons.

**Value**

Nested list of 'shiny.tag' objects - html structure of filtering panel module.

'shiny::moduleServer' output providing server logic for filtering panel module.

**Examples**

```
if (interactive()) {
  library(cohortBuilder)
  library(shiny)
  library(shinyCohortBuilder)

  librarian_source <- set_source(as.tblist(librarian))
  librarian_cohort <- cohort(
    librarian_source,
    filter(
      "discrete", id = "author", dataset = "books",
      variable = "author", value = "Dan Brown",
```

```

      active = FALSE
    ),
    filter(
      "range", id = "copies", dataset = "books",
      variable = "copies", range = c(5, 10),
      active = FALSE
    ),
    filter(
      "date_range", id = "registered", dataset = "borrowers",
      variable = "registered", range = c(as.Date("2010-01-01"), Inf),
      active = FALSE
    )
  )
)

ui <- fluidPage(
  sidebarLayout(
    sidebarPanel(
      cb_ui("librarian")
    ),
    mainPanel()
  )
)

server <- function(input, output, session) {
  cb_server("librarian", librarian_cohort)
}

shinyApp(ui, server)
}

if (interactive()) {
  # enabling latest step filters management
  library(cohortBuilder)
  library(shiny)
  library(shinyCohortBuilder)

  librarian_source <- set_source(as.tbllist(librarian))
  librarian_source$attributes$available_filters <- list(
    filter(
      "discrete", id = "author", dataset = "books",
      variable = "author", value = "Dan Brown",
      active = FALSE
    ),
    filter(
      "range", id = "copies", dataset = "books",
      variable = "copies", range = c(5, 10),
      active = FALSE
    ),
    filter(
      "date_range", id = "registered", dataset = "borrowers",
      variable = "registered", range = c(as.Date("2010-01-01"), Inf),
      active = FALSE
    ),
    filter(

```

```

      "discrete", id = "program", dataset = "borrowers",
      variable = "program", value = NA,
      active = FALSE
    )
  )
  librarian_cohort <- cohort(
    librarian_source,
    filter(
      "range", id = "copies", dataset = "books",
      variable = "copies", range = c(5, 10),
      active = FALSE
    ),
    filter(
      "date_range", id = "registered", dataset = "borrowers",
      variable = "registered", range = c(as.Date("2010-01-01"), Inf),
      active = FALSE
    )
  )

  ui <- fluidPage(
    sidebarLayout(
      sidebarPanel(
        cb_ui("librarian", manage_step = TRUE)
      ),
      mainPanel()
    )
  )

  server <- function(input, output, session) {
    cb_server("librarian", librarian_cohort)
  }

  shinyApp(ui, server)
}

```

---

demo\_app

*Run demo application*

---

## Description

The demo presents available filters and toolbox features.

## Usage

```

demo_app(
  steps = TRUE,
  stats = c("pre", "post"),
  run_button = "none",
  feedback = TRUE,
  state = TRUE,

```

```

bootstrap = 5,
enable_bookmarking = TRUE,
code = TRUE,
attrition = TRUE,
show_help = TRUE,
new_step = c("clone", "configure"),
manage_step = FALSE,
...,
run_app = TRUE
)

```

## Arguments

steps	Set to TRUE (default) if multiple steps should be available.
stats	Choose which statistics should be displayed for data (and some filters). Possible options are: "pre" - previous step stat, "post" - current step stats, 'c("pre", "post")' - for both and NULL for no stats.
run_button	Should Run button be displayed? If so, the current step computations are run only when clicked. Three options are available "none" - no button, "local" - button displayed at each step panel, "global" - button visible in top filtering panel.
feedback	Set to TRUE (default) if feedback plots should be displayed at each filter.
state	Set to TRUE (default) to enable get/set state panel.
bootstrap	Bootstrap version to be used for filtering panel. See <a href="#">bs_theme</a> version argument.
enable_bookmarking	Set to TRUE (default) if panel should be compatible with native shiny bookmarking.
code	Set to TRUE (default) to enable reproducible code panel.
attrition	Set to TRUE (default) to enable attrition plot panel.
show_help	Set to TRUE (default) to enable help buttons.
new_step	Choose which add step method should be used for creating new step. Possible options are: "clone" - copy filters from last step, "configure" - opening modal and allow to chose filters from available filters.
manage_step	Set to TRUE to enable managing filters tool.
...	Extra parameters passed to selected cohort methods. Currently unused.
run_app	If 'TRUE' the application will run using <a href="#">runApp</a> , otherwise <a href="#">shinyApp</a> object is returned.

## Value

In case of 'run\_app=TRUE' no return value, used for side effect which is running a Shiny application. Otherwise [shinyApp](#) object.

**Examples**

```

if (interactive()) {
  library(shinyCohortBuilder)
  demo_app(steps = FALSE)
}
if (interactive()) {
  library(shinyCohortBuilder)
  demo_app(run_button = "local", state = FALSE)
}

```

---

filter-position	<i>Create target filter position in UI panel</i>
-----------------	--

---

**Description**

Create target filter position in UI panel

**Usage**

```

.filter_position(source, step_id, filter, ns, ...)

## Default S3 method:
.filter_position(source, step_id, filter, ns, ...)

## S3 method for class 'tblist'
.filter_position(source, step_id, filter, ns, ...)

```

**Arguments**

source	Source object.
step_id	Id of the step that filter belong to.
filter	Filter object.
ns	Namespace function.
...	Extra arguments passed to related method.

---

gui	<i>Run filtering panel locally</i>
-----	------------------------------------

---

**Description**

Run filtering panel locally

**Usage**

```
gui(
  cohort,
  steps = TRUE,
  stats = c("pre", "post"),
  run_button = "none",
  feedback = TRUE,
  state = TRUE,
  bootstrap = 5,
  enable_bookmarking = TRUE,
  code = TRUE,
  attrition = TRUE,
  show_help = TRUE,
  new_step = c("clone", "configure"),
  manage_step = FALSE
)
```

**Arguments**

cohort	Cohort object with configured filters.
steps	Set to TRUE (default) if multiple steps should be available.
stats	Choose which statistics should be displayed for data (and some filters). Possible options are: "pre" - previous step stat, "post" - current step stats, 'c("pre", "post")' - for both and NULL for no stats.
run_button	Should Run button be displayed? If so, the current step computations are run only when clicked. Three options are available "none" - no button, "local" - button displayed at each step panel, "global" - button visible in top filtering panel.
feedback	Set to TRUE (default) if feedback plots should be displayed at each filter.
state	Set to TRUE (default) to enable get/set state panel.
bootstrap	Bootstrap version to be used for filtering panel. See <a href="#">bs_theme</a> version argument.
enable_bookmarking	Set to TRUE (default) if panel should be compatible with native shiny bookmarking.
code	Set to TRUE (default) to enable reproducible code panel.
attrition	Set to TRUE (default) to enable attrition plot panel.
show_help	Set to TRUE (default) to enable help buttons.
new_step	Choose which add step method should be used for creating new step. Possible options are: "clone" - copy filters from last step, "configure" - opening modal and allow to chose filters from available filters.
manage_step	Set to TRUE to enable managing filters tool.

**Value**

No return value, used for side effect which is running a Shiny application.

**Examples**

```

if (interactive()) {
  library(magrittr)
  library(cohortBuilder)
  library(shinyCohortBuilder)
  mtcars_source <- set_source(tblist(mtcars = mtcars))
  mtcars_cohort <- cohort(
    mtcars_source,
    filter("discrete", id = "am", dataset = "mtcars", variable = "am", value = 1)
  ) %>% run()
  gui(mtcars_cohort)
}

```

---

gui-filter-layer

*Return GUI layer methods for filter of specified type*


---

**Description**

For each filter type `‘.gui_filter’` method should return a list of the below objects:

- `input` - UI structure defining filter input controllers.
- `feedback` - List defining feedback plot output.
- `server` - Optional server-side expression attached to filter panel (e.g. filter specific observers).
- `update` - An expression used for updating filter panel based on its configuration.
- `post_stats` - TRUE if post statistics are displayed in filter controller (e.g. for discrete filter). If FALSE, some operations are skipped which results with better performance.
- `multi_input` - TRUE if multiple input controllers are used for providing filter value (e.g. range input where both `numericInput` and `sliderInput` are used). If FALSE, some operations are skipped which results with better performance.

If you want to learn more about creating filter layers see `‘vignette("gui-filter-layer")’`.

**Usage**

```

.gui_filter(filter, ...)

## S3 method for class 'discrete'
.gui_filter(filter, ...)

## S3 method for class 'range'
.gui_filter(filter, ...)

## S3 method for class 'date_range'
.gui_filter(filter, ...)

```

```
## S3 method for class 'datetime_range'
.gui_filter(filter, ...)

## S3 method for class 'discrete_text'
.gui_filter(filter, ...)

## S3 method for class 'multi_discrete'
.gui_filter(filter, ...)

## S3 method for class 'query'
.gui_filter(filter, ...)
```

### Arguments

filter	Filter object.
...	Extra arguments passed to a specific method.

### Value

List consisting filter metadata and methods that allow to perform filter based operations. See `'vignette("custom-filters")'`.

### See Also

[source-gui-layer](#)

### Examples

```
library(cohortBuilder)
librarian_source <- set_source(as.tbllist(librarian))
copies_filter <- filter(
  "range", id = "copies", name = "Copies", dataset = "books",
  variable = "copies", range = c(5, 12)
)
copies_filter_evaled <- copies_filter(librarian_source)
copies_filter_evaled$gui <- .gui_filter(copies_filter_evaled)

str(copies_filter_evaled$gui)
```

---

keep\_na\_input

*Generate NA's filter selection GUI input*

---

### Description

When used within filter's GUI input method, the component is responsible for updating 'keep\_na' filter parameter.

Use `'update_keep_na_input'` inside filter's GUI update method to update the output based on the filter state.

**Usage**

```
.keep_na_input(
  input_id,
  filter,
  cohort,
  msg_fun = function(x) glue::glue("Keep missing values ({x})")
)

.update_keep_na_input(
  session,
  input_id,
  filter,
  cohort,
  msg_fun = function(x) glue::glue("Keep missing values ({x})")
)
```

**Arguments**

input_id	Id of the keep na input.
filter	Filter object.
cohort	Cohort object.
msg_fun	Function taking number of missing values as an argument and returning missing values label.
session	Shiny session object.

**Value**

Nested list of 'shiny.tag' objects storing html structure of the input, or no value in case of usage 'update' method.

**Examples**

```
library(magrittr)
library(cohortBuilder)

librarian_source <- set_source(as.tbllist(librarian))
coh <- cohort(
  librarian_source,
  filter(
    "range", id = "copies", name = "Copies", dataset = "books",
    variable = "copies", range = c(5, 12)
  )
) %>% run()
.keep_na_input("keep_na", coh$get_filter("1", "copies"), coh)
```

---

pre\_post\_stats      *Generate structure of pre/post statistics*

---

### Description

The method exported only for custom extensions use.

‘pre\_post\_stats’ returns the statistics having html tags structure. ‘pre\_post\_stats\_text’ returns the same output but flatten to a single character object. The latter function works faster and supports vector arguments.

### Usage

```
.pre_post_stats(  
  current,  
  previous,  
  name,  
  brackets = FALSE,  
  percent = FALSE,  
  stats = c("pre", "post")  
)
```

```
.pre_post_stats_text(  
  current,  
  previous,  
  name,  
  brackets = TRUE,  
  percent = FALSE,  
  stats = c("pre", "post")  
)
```

### Arguments

current	Current step statistic value.
previous	Previous step statistic value.
name	Name displayed nearby the statistics output.
brackets	If TRUE, statistics will be displayed in brackets.
percent	Should current/previous ration in percentages be displayed?
stats	Vector of "pre" and "post" defining which statistics should be returned. "pre" for previous, "post" for current and NULL for none.

### Value

A ‘shiny.tag’ class ‘span’ object defining html structure of data/value statistics, or character object.

## Examples

```
.pre_post_stats(5, 10, "books")
.pre_post_stats_text(5, 10, "books")
.pre_post_stats(5, 10, "books", brackets = TRUE)
.pre_post_stats_text(5, 10, "books", brackets = TRUE)
.pre_post_stats(5, 10, "books", percent = TRUE)
.pre_post_stats_text(5, 10, "books", percent = TRUE)
.pre_post_stats_text(5:6, 10:11, "books", percent = TRUE)
```

---

rendering-custom-attribution

*Method for generating custom attrition output*

---

## Description

When method is defined for selected source, the output is displayed in attrition modal tab.

## Usage

```
.custom_attrition(source, ...)
```

## Arguments

source	Source object.
...	Extra arguments passed to specific method.

## Details

Similar to [.step\\_attrition](#) the method should return list of 'render' and 'output' expressions.

## Value

List of two objects: 'render' and 'output' defining rendering and output placeholder for custom attrition plot feature.

## See Also

[source-gui-layer](#)

## Examples

```
if (interactive()) {
  library(magrittr)
  library(shiny)
  library(cohortBuilder)
  library(shinyCohortBuilder)
```

```

.custom_attrition.tbllist <- function(source, id, cohort, session, ...) {
  ns <- session$ns
  choices <- names(source$dtconn)

  list(
    render = shiny::renderPlot({
      cohort$show_attrition(dataset = session$input$attrition_input)
    }),
    output = shiny::tagList(
      shiny::h3("Step-wise Attrition Plot"),
      shiny::selectInput(ns("attrition_input"), "Choose dataset", choices),
      shiny::plotOutput(id)
    )
  )
}
coh <- cohort(
  set_source(as.tbllist(librarian)),
  step(
    filter(
      "range", id = "copies", dataset = "books",
      variable = "copies", range = c(5, 12)
    )
  ),
  step(
    filter(
      "range", id = "copies", dataset = "books",
      variable = "copies", range = c(6, 8)
    )
  )
) %>% run()

ui <- fluidPage(
  div(id = "attrition")
)

server <- function(input, output, session) {
  rendering <- .custom_attrition(
    coh$get_source(), id = "attr", cohort = coh, session = session, dataset = "books"
  )
  insertUI("#attrition", ui = rendering$output)
  output$attr <- rendering$render
}

shinyApp(ui, server)
}

```

**Description**

The method exported only for custom extensions use.

**Usage**

```
.render_filters(source, ...)

## Default S3 method:
.render_filters(source, cohort, step_id, ns, ...)

## S3 method for class 'tblist'
.render_filters(source, cohort, step_id, ns, ...)
```

**Arguments**

source	Source object.
...	Extra arguments passed to a specific method.
cohort	Cohort object.
step_id	Id of the step.
ns	Namespace function.

**Details**

Within the method you should define source data stats output (see [.update\\_data\\_stats](#)), and define a loop that renders filtering panel for each filter (using [.render\\_filter](#)).

**Value**

Nested list of 'shiny.tag' objects storing html structure of filter input panels.

**See Also**

[source-gui-layer](#)

**Examples**

```
if (interactive()) {
  library(magrittr)
  library(shiny)
  library(cohortBuilder)
  library(shinyCohortBuilder)

  ui <- fluidPage(
    actionButton("add_filter", "Add Filter"),
    div(id = "filter_container")
  )

  server <- function(input, output, session) {
    add_gui_filter_layer <- function(public, private, ...) {
```

```

    private$steps[["1"]]$filters$copies$gui <- .gui_filter(
      private$steps[["1"]]$filters$copies
    )
    private$steps[["1"]]$filters$registered$gui <- .gui_filter(
      private$steps[["1"]]$filters$registered
    )
  }
  add_hook("post_cohort_hook", add_gui_filter_layer)
  coh <- cohort(
    set_source(as.tblist(librarian)),
    filter(
      "range", id = "copies", name = "Copies", dataset = "books",
      variable = "copies", range = c(5, 12)
    ),
    filter(
      "date_range", id = "registered", name = "Registered", dataset = "borrowers",
      variable = "registered", range = c(as.Date("2010-01-01"), Inf)
    )
  ) %>% run()
  coh$attributes$session <- session
  coh$attributes$feedback <- TRUE

  observeEvent(input$add_filter, {
    insertUI(
      "#filter_container",
      ui = .render_filters(
        coh$get_source(),
        cohort = coh,
        step_id = "1",
        ns = function(x) x
      )
    ), ignoreInit = TRUE, once = TRUE)
  }

  shinyApp(ui, server)
}

```

---

rendering-step-attribution

*Generate output of attrition plot*

---

### Description

The method should return list of two object:

- render - Rendering expression of attrition output.
- output - Output expression related to rendering (with id equal to 'id' parameter).

For example:

```
list(  
  render = shiny::renderPlot({  
    cohort$show_attrition()  
  }),  
  output = shiny::plotOutput(id)  
)
```

### Usage

```
.step_attrition(source, ...)  
  
## Default S3 method:  
.step_attrition(source, id, cohort, session, ...)  
  
## S3 method for class 'tblist'  
.step_attrition(source, id, cohort, session, ...)
```

### Arguments

source	Source object.
...	Extra arguments passed to specific method.
id	Id of attrition output.
cohort	Cohort object.
session	Shiny session object.

### Value

List of two objects: 'render' and 'output' defining rendering and output placeholder for step attrition plot feature.

### See Also

[source-gui-layer](#)

### Examples

```
if (interactive()) {  
  library(magrittr)  
  library(shiny)  
  library(cohortBuilder)  
  library(shinyCohortBuilder)  
  
  coh <- cohort(  
    set_source(as.tblist(librarian)),  
    step(  
      filter(  
        "range", id = "copies", dataset = "books",  
        variable = "copies", range = c(5, 12)  
      )  
    ),  
  ),  
}
```

```
    step(
      filter(
        "range", id = "copies", dataset = "books",
        variable = "copies", range = c(6, 8)
      )
    )
  ) %>% run()

  ui <- fluidPage(
    div(id = "attrition")
  )

  server <- function(input, output, session) {
    rendering <- .step_attrition(
      coh$get_source(), id = "attr", cohort = coh, session = session, dataset = "books"
    )
    insertUI("#attrition", ui = rendering$output)
    output$attr <- rendering$render
  }

  shinyApp(ui, server)
}
```

---

scb\_chart\_palette

*Default color palette used for filter feedback plots*

---

## Description

It's a list of the following elements:

## Usage

```
scb_chart_palette
```

## Format

An object of class `list` of length 3.

## Details

- `discrete` - Discrete filter plot colors.
- `histogram` - Range and date range histogram color.
- `no_data` - Color used to mark missing variables on feedback plots.

The palette is used as default `scb_chart_palette` option, that can be overwritten with custom palettes.

---

scb_icons	<i>Default filtering panel icons</i>
-----------	--------------------------------------

---

**Description**

Icons can be overwritten with using `sbc_icons` option.

**Usage**

`scb_icons`

**Format**

An object of class `list` of length 16.

---

scb_labels	<i>Default filtering panel labels</i>
------------	---------------------------------------

---

**Description**

Labels can be overwritten with using `sbc_labels` option.

**Usage**

`scb_labels`

**Format**

An object of class `list` of length 16.

---

source-gui-layer	<i>Source compatibility methods.</i>
------------------	--------------------------------------

---

**Description**

List of methods that allow compatibility of different source types. Most of the methods should be defined in order to make new source layer functioning. See 'Details' section for more information.

## Details

The package is designed to make the functionality work with multiple data sources. Data source can be based for example on list of tables, connection to database schema or API service that allows to access and operate on data. In order to make new source type layer functioning, the following list of methods should be defined:

- `.render_filters`
- `.update_data_stats`
- `.step_attrition`
- `.custom_attrition`
- `.available_filter_choices`
- `autofilter`

Except from the above methods, you may extend the existing or new source with providing custom gui filtering methods. See [gui-filter-layer](#). In order to see more details about how to implement custom source check `'vignette("custom-gui-layer")'`.

## Value

Various type outputs dependent on the selected method. See each method documentation for details.

---

<code>trigger-action</code>	<i>Trigger filtering panel action</i>
-----------------------------	---------------------------------------

---

## Description

The two functions that allow to trigger a specific filtering panel action directly from Shiny server (`.trigger_action`) or application browser (`.trigger_action_js`) attached to a specific JS event, e.g. `onclick`.

Check Details section to see possible options.

## Usage

```
.trigger_action(session, action, params = NULL)
```

```
.trigger_action_js(action, params = list(), ns = function(id) id)
```

## Arguments

<code>session</code>	Shiny session object.
<code>action</code>	Id of the action.
<code>params</code>	List of parameters passed to specific action method.
<code>ns</code>	Namespace function (if used within Shiny modal).

## Details

The list of possible actions:

- `update_filter` - Calls `'shinyCohortBuilder:::gui_update_filter'` that triggers filter arguments update.
- `add_step` - Calls `'shinyCohortBuilder:::gui_add_step'` that triggers adding a new filtering step (based on configuration of the previous one).
- `rm_step` - Calls `'shinyCohortBuilder:::gui_rm_step'` used to remove a selected filtering step.,
- `clear_step` - Calls `'shinyCohortBuilder:::gui_clear_step'` used to clear filters configuration in selected step.
- `update_step` - Calls `'shinyCohortBuilder:::gui_update_step'` used to update filters and feedback plots for the specific filter GUI panel.
- `update_data_stats` - Calls `'shinyCohortBuilder:::gui_update_data_stats'` that is called to update data statistics.
- `show_repro_code` - Calls `'shinyCohortBuilder:::gui_show_repro_code'` that is used to show reproducible code.
- `run_step` - Calls `'shinyCohortBuilder:::gui_run_step'` used to trigger specific step data calculation.
- `show_state` - Calls `'shinyCohortBuilder:::gui_show_state'` that is used to show filtering panel state json.
- `input_state` - Calls `'shinyCohortBuilder:::gui_input_state'` that is used to generate modal in which filtering panel state can be provided (as json).
- `restore_state` - Calls `'shinyCohortBuilder:::gui_restore_state'` used for restoring filtering panel state based on provided json.
- `show_attrition` - Calls `'shinyCohortBuilder:::gui_show_attrition'` a method used to show attrition data plot(s).

Both `'trigger_action'` and `'trigger_action_js'` methods are exported for advanced use only.

## Value

No return value (`'trigger_action'` - sends message to the browser) or character string storing JS code for sending input value to Shiny server (`'trigger_action_js'`).

## Examples

```
if (interactive()) {
  library(shiny)
  library(shinyCohortBuilder)

  shiny::addResourcePath(
    "shinyCohortBuilder",
    system.file("www", package = "shinyCohortBuilder")
  )
  ui <- fluidPage(
    tags$head(
```

```

shiny::tags$script(type = "text/javascript", src = file.path("shinyCohortBuilder", "scb.js"))
),
tags$button(
  "Trigger action from UI", class = "btn btn-default",
  onclick = .trigger_action_js("uiaction", params = list(a = 1))
),
actionButton("send", "Trigger action from server")
)

server <- function(input, output, session) {
  observeEvent(input$send, {
    .trigger_action(session, "serveraction", params = list(a = 2))
  })
  observeEvent(input$action, {
    print(input$action)
  })
}

shinyApp(ui, server)
}

```

---

 updating-data-statistics

*Render source data related statistics*


---

### Description

The function should assign rendering that displays data source statistics to the valid output. By default, the output is placed within `.render_filters` method.

### Usage

```

.update_data_stats(source, ...)

## Default S3 method:
.update_data_stats(source, step_id, cohort, session, ...)

## S3 method for class 'tblist'
.update_data_stats(source, step_id, cohort, session, ...)

```

### Arguments

source	Source object.
...	Extra arguments passed to a specific method.
step_id	Id if filtering step.
cohort	Cohort object.
session	Shiny session object.

**Details**

When rendering the output, a good practice is to use cached data statistics available with `'cohort$get_cache(step_id)'`. This way, you omit running additional computations which results with performance improvement.

**Value**

No return value, used for side effect which assigning Cohort data statistics to the `'output'` object.

**See Also**

[source-gui-layer](#)

**Examples**

```
if (interactive()) {
  library(magrittr)
  library(shiny)
  library(cohortBuilder)
  library(shinyCohortBuilder)

  ui <- fluidPage(
    sliderInput("step_two_max", "Max step two copies", min = 6, max = 12, value = 8),
    uiOutput("2-stats_books")
  )

  server <- function(input, output, session) {
    coh <- cohort(
      set_source(as.tblist(librarian)),
      step(
        filter(
          "range", id = "copies", dataset = "books",
          variable = "copies", range = c(5, 12)
        )
      ),
      step(
        filter(
          "range", id = "copies", dataset = "books",
          variable = "copies", range = c(6, 8)
        )
      )
    ) %>% run()
    coh$attributes$stats <- c("pre", "post")
    observeEvent(input$step_two_max, {
      coh$update_filter("copies", step_id = 2, range = c(6, input$step_two_max))
      run(coh, min_step_id = "2")
      .update_data_stats(coh$get_source(), step_id = "2", cohort = coh, session = session)
    })
  }

  shinyApp(ui, server)
}
```



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