

# Package ‘safetyCharts’

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**Title** Charts for Monitoring Clinical Trial Safety

**Version** 0.2.0

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**Description** Contains chart code for monitoring clinical trial safety. Charts can be used as standalone output, but are also designed for use with the 'safetyGraphics' package, which makes it easy to load data and customize the charts using an interactive web-based interface created with Shiny.

**URL** <https://github.com/SafetyGraphics/safetyCharts>

**BugReports** <https://github.com/SafetyGraphics/safetyCharts/issues>

**License** MIT + file LICENSE

**Encoding** UTF-8

**RoxygenNote** 7.1.1

**Imports** dplyr, DT, Tplyr, ggplot2, rlang, shiny, knitr, RColorBrewer, stringr, forcats, Tendiril, kableExtra, huxtable, pharmaRTF

**Suggests** testthat, shinytest, safetyData, safetyGraphics, yaml

**Depends** R (>= 4.0)

**NeedsCompilation** no

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**Repository** CRAN

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demogRTF\_server      *Demographics Table RTF - UI*

**Description**

Demographics Table RTF - UI

**Usage**

demogRTF\_server(input, output, session, params)

**Arguments**

input	module input
output	module output
session	module session
params	parameters object with data and settings options.

**Value**

returns shiny module Server function

demogRTF\_table      *create demographics RTF table*

**Description**

create demographics RTF table

**Usage**

demogRTF\_table(data, settings)

**Arguments**

data	demographics data frame with columns specified in settings object
settings	list with parameters specifying the column names for: <ul style="list-style-type: none"><li>• sex (settings\$sex_col),</li><li>• race (settings\$race_col)</li><li>• age (settings\$age_Col)</li></ul>

**Value**

rtf doc object

**Examples**

```
settings <- list(treatment_col = "ARM", sex_col = "SEX", race_col = "RACE", age_col = "AGE")
demogRTF_table(safetyData::sdtm_dm, settings)
```

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demogRTF_ui	<i>Demographics Table RTF - UI</i>
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**Description**

Demographics Table RTF - UI

**Usage**

```
demogRTF_ui(id)
```

**Arguments**

id	module id
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**Value**

returns shiny module UI

---

init\_aeExplorer      *Initialize Settings for Adverse Event Explorer widget*

---

**Description**

Initialize Settings for Adverse Event Explorer widget

**Usage**

```
init_aeExplorer(data, settings)
```

**Arguments**

data	labs data structured as one record per person per visit per measurement. See details for column requirements.
settings	named list of settings

**Value**

returns list with data and settings

---

init\_paneledOutlierExplorer  
*Initialize Settings for Paneled Outlier Explorer widget*

---

**Description**

Initialize Settings for Paneled Outlier Explorer widget

**Usage**

```
init_paneledOutlierExplorer(data, settings)
```

**Arguments**

data	labs data structured as one record per person per visit per measurement. See details for column requirements.
settings	named list of settings

**Value**

returns list with data and settings

---

`init_safetyOutlierExplorer`

*Initialize Settings for Safety Outlier Explorer widget*

---

**Description**

Initialize Settings for Safety Outlier Explorer widget

**Usage**

`init_safetyOutlierExplorer(data, settings)`

**Arguments**

<code>data</code>	labs data structured as one record per person per visit per measurement. See details for column requirements.
<code>settings</code>	named list of settings

**Value**

returns list with data and settings

---

`init_safetyResultsOverTime`

*Initialize Settings for Safety Results Over Time widget*

---

**Description**

Initialize Settings for Safety Results Over Time widget

**Usage**

`init_safetyResultsOverTime(data, settings)`

**Arguments**

<code>data</code>	labs data structured as one record per person per visit per measurement. See details for column requirements.
<code>settings</code>	named list of settings

**Value**

returns list with data and settings

`init_safetyShiftPlot` *Initialize Settings for Safety Shift Plot widget*

---

**Description**

Initialize Settings for Safety Shift Plot widget

**Usage**

```
init_safetyShiftPlot(data, settings)
```

**Arguments**

<code>data</code>	labs data structured as one record per person per visit per measurement. See details for column requirements.
<code>settings</code>	named list of settings

**Value**

returns list with data and settings

---

`lab_distribution_server`

*lab distribution Module - Server*

---

**Description**

A simple server for a shiny module looking at lab histograms. Intended primarily for technical demos.

**Usage**

```
lab_distribution_server(input, output, session, params)
```

**Arguments**

<code>input</code>	module input
<code>output</code>	module output
<code>session</code>	module session
<code>params</code>	parameters object with data and settings options.

**Value**

returns shiny module Server function

---

lab\_distribution\_ui    *Lab distribution Module - UI*

---

**Description**

A simple UI for a shiny module looking at lab histograms. Intended primarily for technical demos.

**Usage**

```
lab_distribution_ui(id)
```

**Arguments**

id	module id
----	-----------

**Value**

returns shiny module UI

---

safetyOutlierExplorer\_server  
*Safety Outlier Explorer Module - UI*

---

**Description**

Safety Outlier Explorer Module - UI

**Usage**

```
safetyOutlierExplorer_server(input, output, session, params)
```

**Arguments**

input	module input
output	module output
session	module session
params	parameters object with data and settings options.

**Value**

returns shiny module Server function

---

safetyOutlierExplorer\_ui

*Safety Outlier Explorer Module - UI*

---

### Description

Safety Outlier Explorer Module - UI

### Usage

```
safetyOutlierExplorer_ui(id)
```

### Arguments

id	module id
----	-----------

### Value

returns shiny module UI

---

safety\_outlier\_explorer

*Safety Outlier Explorer*

---

### Description

Safety Outlier Explorer

### Usage

```
safety_outlier_explorer(data, settings)
```

### Arguments

data	labs data structured as one record per person per visit per measurement. See details for column requirements.
settings	named list of settings with the parameters specified below.

### Details

The settings object provides details the columns in the data set.

- "id\_col" ID column
- "value\_col" Value column
- "measure\_col" Measure column
- "measure\_values" Measure values
- "studyday\_col" Study Day (numeric)



**Value**

returns a chart object

**Examples**

```
settings <- list(  
  id_col = "USUBJID",  
  measure_col = "LBTEST",  
  measure_values = c("Albumin", "Bilirubin", "Chloride"),  
  studyday_col = "VISITDY",  
  value_col = "LBORRES"  
)  
safety_outlier_explorer(safetyData::sdtm_lb, settings)
```

---

safety\_results\_over\_time

*Safety Results Over Time plot*

---

**Description**

Safety Results Over Time plot

**Usage**

```
safety_results_over_time(data, settings)
```

**Arguments**

data	labs data structured as one record per person per visit per measurement. See details for column requirements.
settings	named list of settings with the parameters specified below.

**Details**

The settings object provides details the columns in the data set.

- "value\_col" Value column
- "measure\_col" Measure column
- "measure\_values" Measure values
- "visit\_col" Study Visit
- "visitn\_col" Study Number
- "group\_col" Grouping column
- "violins" Show Violin plots?
- "boxplots" Show Box Plots?
- "axis" set to "log" to use a log transformed axis, linear otherwise
- "drop\_visit\_string" Drop visits that contain this string. e.g. "unscheduled"

**Value**

returns a chart object

**Examples**

```
library(dplyr)
lb <- safetyData::sdtm_lb
sub_ids <- unique(lb$USUBJID)[1:100]
lb <- lb %>% filter(USUBJID %in% sub_ids)
settings <- list(
  value_col = "LBORRES",
  measure_col = "LBTEST",
  measure_values = c("Chloride"),
  visit_col = "VISIT",
  visitn_col = "VISITNUM",
  axis = "log"
)
safety_results_over_time(lb, settings)

# remove unscheduled visits, add violin plot and 2nd panel
settings$drop_visit_string <- "unscheduled"
settings$violins <- TRUE
settings$measure_values <- c("Albumin")
safety_results_over_time(lb, settings)

# add grouping by treatment
dm_sub <- safetyData::sdtm_dm %>% select(USUBJID, ARM)
dm_lb <- dm_sub %>% left_join(lb)
settings$group_col <- "ARM"
safety_results_over_time(dm_lb, settings)
```

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tendrill\_chart

*Tendrill plot*


---

**Description**

Create a plot using the Tendril package

**Usage**

```
tendrill_chart(data, settings)
```

**Arguments**

data	list of data frames including dataframes named aes (adverse events) and dm (demographics)
settings	named list of domain-specific settings with the parameters specified below.

**Details**

The settings object provides details regarding the columns in the data sets.

- "settings\$dm\$id\_col" ID column
- "settings\$dm\$treatment\_col" Treatment column
- "settings\$dm\$treatment\_values-group1" Name of treatment 1
- "settings\$dm\$treatment\_values-group2" Name of treatment 2
- "settings\$aes\$id\_col" ID column)
- "settings\$aes\$bodsys\_col" Body System
- "settings\$aes\$stdy\_col" Study Day

**Value**

returns a chart object

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