

# Package: boxly (via r-universe)

September 14, 2024

**Title** Interactive Box Plot

**Version** 0.1.1

**Description** Interactive box plot using 'plotly' for clinical trial analysis.

**License** GPL (>= 3)

**URL** <https://merck.github.io/boxly/>, <https://github.com/Merck/boxly>

**BugReports** <https://github.com/Merck/boxly/issues>

**Encoding** UTF-8

**LazyData** true

**Depends** R (>= 4.1.0)

**Imports** DT, brew, rlang, crosstalk, ggplot2, htmlwidgets, htmltools, metalite, plotly, uuid

**Suggests** covr, knitr, rmarkdown, testthat (>= 3.0.0)

**VignetteBuilder** knitr

**Config/testthat/edition** 3

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.2.3

**Repository** <https://merck.r-universe.dev>

**RemoteUrl** <https://github.com/merck/boxly>

**RemoteRef** HEAD

**RemoteSha** f585e7cc70d44645548d640706407fc298beb8b5

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|              |                                       |
|--------------|---------------------------------------|
| <b>boxly</b> | <i>Create an interactive box plot</i> |
|--------------|---------------------------------------|

**Description**

Create an interactive box plot

**Usage**

```
boxly(
  outdata,
  color = NULL,
  hover_summary_var = c("n", "min", "q1", "median", "mean", "q3", "max"),
  hover_outlier_label = c("Participant ID", "Parameter value"),
  x_label = "Visit",
  y_label = "Change",
  heading_select_list = "Lab parameter",
  heading_summary_table = "Number of Participants"
)
```

**Arguments**

|                                    |  |
|------------------------------------|--|
| <code>outdata</code>               | An <code>outdata</code> object created from <code>prepare_ae_forestly()</code> .                                     |
| <code>color</code>                 | Color for box plot.  |
| <code>hover_summary_var</code>     | A character vector of statistics to be displayed on hover label of box.  |
| <code>hover_outlier_label</code>   | A character vector of hover label for outlier. A label from an input data is used if NA for a variable is specified. |
| <code>x_label</code>               | x-axis label.  |
| <code>y_label</code>               | y-axis label.  |
| <code>heading_select_list</code>   | Select list menu label.  |
| <code>heading_summary_table</code> | Summary table label.   |

**Value**

Interactive box plot.

## Examples

```
# Only run this example in interactive R sessions
if (interactive()) {
  library(metalite)

  meta_boxly(
    boxly_adsl,
    boxly_adlb,
    population_term = "apat",
    observation_term = "wk12"
  ) |>
  prepare_boxly() |>
  boxly()
}
```

---

boxly\_adeg

*An example ADEG dataset*

---

## Description

Definition of each variable can be found in <https://github.com/phuse-org/phuse-scripts/tree/master/data/adam/cdisc>

## Usage

boxly\_adeg

## Format

A data frame with 32139 and 35 variables:

## Source

<https://github.com/phuse-org/phuse-scripts/tree/master/data/adam/cdisc>

---

boxly\_adlb

*An example ADLB dataset*

---

## Description

Definition of each variable can be found in <https://github.com/phuse-org/phuse-scripts/tree/master/data/adam/cdisc>

## Usage

boxly\_adlb

**Format**

A data frame with 24746 and 24 variables:

**Source**

<https://github.com/phuse-org/phuse-scripts/tree/master/data/adam/cdisc>

---

boxly\_adsl

*A Subject Level Demographic Dataset*

---

**Description**

A dataset containing the demographic information of a clinical trial following CDISC ADaM standard.

**Usage**

boxly\_adsl

**Format**

A data frame with 254 rows and 51 variables.

**Details**

Definition of each variable can be found in <https://github.com/phuse-org/phuse-scripts/tree/master/data/adam/cdisc>

**Source**

<https://github.com/phuse-org/phuse-scripts/tree/master/data/adam/cdisc>

---

boxly\_advs

*An example ADVS dataset*

---

**Description**

Definition of each variable can be found in <https://github.com/phuse-org/phuse-scripts/tree/master/data/adam/cdisc>

**Usage**

boxly\_advs

**Format**

A data frame with 32139 and 34 variables:

**Source**

<https://github.com/phuse-org/phuse-scripts/tree/master/data/adam/cdisc>

---

meta\_boxly

*Create an example metadata object*

---

**Description**

Create an example metadata object

**Usage**

```
meta_boxly(  
  dataset_adsl,  
  dataset_param,  
  population_term,  
  population_subset = SAFFL == "Y",  
  observation_term,  
  observation_subset = SAFFL == "Y",  
  parameters = unique(dataset_param$PARAMCD)  
)
```

**Arguments**

dataset\_adsl     ADSL source dataset.  
dataset\_param    Observation level source dataset for boxplot.  
population\_term                A character value of population term name.  
population\_subset              An unquoted condition for selecting the populations from ADSL dataset.  
observation\_term              A character value of observation term name.  
observation\_subset            An unquoted condition for selecting the observations from dataset\_param dataset.  
parameters                  A character vector of parameters defined in dataset\_param\$PARAMCD

**Value**

A metalite object.

**Examples**

```
meta_boxly(  
  boxly_adsl,  
  boxly_adlb,  
  population_term = "apat",  
  observation_term = "wk12"  
)
```

---

|                            |  |
|----------------------------|--|
| <code>prepare_boxly</code> | <i>Prepare data for interactive box plot</i> |
|----------------------------|--|

---

## Description

Prepare data for interactive box plot

## Usage

```
prepare_boxly(
  meta,
  population = NULL,
  observation = NULL,
  analysis = NULL,
  hover_var_outlier = c("USUBJID", metalite::collect_adam_mapping(meta, analysis)$y)
)
```

## Arguments

|                                |   |
|--------------------------------|---|
| <code>meta</code>              | A metadata object created by metalite.  |
| <code>population</code>        | A character value of population term name. The term name is used as key to link information.  |
| <code>observation</code>       | A character value of observation term name. The term name is used as key to link information. |
| <code>analysis</code>          | A character value of analysis term name. The term name is used as key to link information.    |
| <code>hover_var_outlier</code> | A character vector of hover variables for outlier.  |

## Value

Metadata list with plotting dataset.

Metadata list with plotting dataset

## Examples

```
library(metalite)

meta <- meta_boxly(
  boxly_ads1,
  boxly_ad1b,
  population_term = "apat",
  observation_term = "wk12"
)
prepare_boxly(meta)
```

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