

Package ‘causalverse’

August 28, 2023

Type Package

Date 2023-08-16

Title Causality in Clarity

Version 0.0.0.9000

Maintainer The package maintainer <mikenguyen.contact@gmail.com>

Description CausalVerse: An R toolkit expediting causal research & analysis. Streamlines complex methodologies, empowering users to unveil causal relationships with precision. Your go-to for insightful causality exploration..

License GPL-3 | file LICENSE

Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.2.3

Suggests knitr,
rmarkdown,
testthat (>= 3.0.0)

Config/testthat/edition 3

Imports ggplot2 (>= 3.4.2),
ggthemes (>= 4.2.4),
tidyverse (>= 2.0.0),
lubridate (>= 1.9.2),
rio (>= 0.5.29),
xtable (>= 1.8.4),
dplyr (>= 1.1.1),
tidyr (>= 1.3.0),
scales (>= 1.2.1),
gridExtra (>= 2.3),
systemfit (>= 1.1.30),
Hotelling (>= 1.0.8),
MatchIt (>= 4.5.4),
rlang (>= 1.1.1),
fixest (>= 0.11.1),
stats (>= 4.2.3),
PanelMatch (>= 2.0.1)

VignetteBuilder knitr

R topics documented:

| | |
|---------------------------|----|
| ama_export_fig | 2 |
| ama_export_tab | 3 |
| ama_labs | 3 |
| ama_scale_color | 4 |
| ama_scale_fill | 5 |
| ama_theme | 6 |
| plot_coef_par_trends | 7 |
| plot_density_by_treatment | 8 |
| plot_par_trends | 9 |
| plot_treat_time | 11 |

| | |
|--------------|-----------|
| Index | 13 |
|--------------|-----------|

| | |
|----------------|---|
| ama_export_fig | <i>Function to export a figure with custom settings</i> |
|----------------|---|

Description

This function exports a ggplot2 figure to a given path. It exports both an archived version with the current date and a current version without a date. The function supports exporting to PDF and JPG formats.

Usage

```
ama_export_fig(figure, filename, filepath, width = 7, height = 7)
```

Arguments

| | |
|----------|---|
| figure | A ggplot2 object. |
| filename | A character string specifying the filename without the extension. |
| filepath | A character string specifying the directory to save the file. |
| width | The width of the image in inches (default is 7 inches). |
| height | The height of the image in inches (default is 7 inches). |

Examples

```
## Not run:
test_plot <- ggplot(mpg, aes(x=displ, y=hwy)) + geom_point() # Create a ggplot2 plot
filename <- "sample_plot" # Define a filename
filepath <- tempdir() # Define a path using a temporary directory
ama_export_fig(test_plot, filename, filepath) # Call the ama_export_fig function

## End(Not run)
```

| | |
|----------------|--|
| ama_export_tab | <i>Function to export a table with AMA style</i> |
|----------------|--|

Description

This function exports the provided table in both Excel (.xlsx) and LaTeX (.tex) formats. The table is archived with the current date in the filename for the Excel version, while the LaTeX version is saved with just the specified filename.

Usage

```
ama_export_tab(table, filename, filepath, caption = NULL)
```

Arguments

| | |
|----------|---|
| table | A data frame or matrix. |
| filename | A character string specifying the filename without the extension. |
| filepath | A character string specifying the directory to save the file. |
| caption | A character string specifying the caption for the table. |

Examples

```
## Not run:
data(mtcars) # Load the mtcars dataset
ama_export_tab(mtcars[1:5, 1:5], "sample_table", tempdir(), "Sample Caption for mtcars")

## End(Not run)
```

| | |
|----------|--|
| ama_labs | <i>Custom Label Formatting for ggplot2: American Marketing Association Style</i> |
|----------|--|

Description

This function provides custom label formatting for ggplot2 based on the guidelines set by the American Marketing Association.

Usage

```
ama_labs(
  title = NULL,
  subtitle = NULL,
  caption = NULL,
  x = NULL,
  y = NULL,
  fill = NULL,
  color = NULL,
  ...
)
```

Arguments

| | |
|----------|---|
| title | Plot title. |
| subtitle | Plot subtitle. |
| caption | Plot caption. |
| x | X-axis label. |
| y | Y-axis label. |
| fill | Fill legend title. |
| color | Color legend title. |
| ... | Additional arguments to be passed to <code>ggplot2::labs()</code> . |

Value

Modified labels for a `ggplot2` plot.

Examples

```
## Not run:
library(ggplot2)
ggplot(mtcars, aes(mpg, wt)) + geom_point() +
  ama_labs(title = "Sample Plot") +
  ama_theme()

## End(Not run)
```

| | |
|-----------------|---|
| ama_scale_color | <i>Custom Color Scale for ggplot2: American Marketing Association Style</i> |
|-----------------|---|

Description

This function provides a custom color scale for `ggplot2` plots based on the guidelines set by the American Marketing Association.

Usage

```
ama_scale_color(
  use_color = FALSE,
  palette_name = "OkabeIto",
  grayscale_limits = c(0.2, 0.8)
)
```

Arguments

| | |
|------------------|---|
| use_color | Logical. If TRUE, uses color, otherwise uses grayscale. |
| palette_name | Character. Name of the color palette to use. |
| grayscale_limits | Numeric vector. Limits for the grayscale gradient. |

Value

A color scale for a ggplot2 plot.

Examples

```
## Not run:
library(ggplot2)
ggplot(mtcars, aes(mpg, wt, color = gear)) + geom_point(size = 4) + ama_scale_color()

## End(Not run)
```

ama_scale_fill*Custom Fill Scale for ggplot2: American Marketing Association Style*

Description

This function provides a custom fill scale for ggplot2 plots based on the guidelines set by the American Marketing Association.

Usage

```
ama_scale_fill(
  use_color = FALSE,
  palette_name = "OkabeIto",
  grayscale_limits = c(0.2, 0.8)
)
```

Arguments

use_color Logical. If TRUE, uses color, otherwise uses grayscale.

palette_name Character. Name of the color palette to use.

grayscale_limits Numeric vector. Limits for the grayscale gradient.

Value

A fill scale for a ggplot2 plot.

Examples

```
## Not run:
library(ggplot2)
ggplot(mtcars, aes(mpg, wt, fill = gear)) +
  geom_point(shape = 21, size = 4) +
  ama_scale_fill()

## End(Not run)
```

ama_theme

*Custom Theme for ggplot2: American Marketing Association Style***Description**

This function provides a custom theme for ggplot2 following the guidelines set by the American Marketing Association.

Usage

```
ama_theme(
  base_size = 16,
  base_family = "sans",
  title_size = ggplot2::rel(1.2),
  axis_title_size = ggplot2::rel(1.2),
  legend_title_size = ggplot2::rel(0.6),
  legend_text_size = ggplot2::rel(0.6),
  axis_text_size = ggplot2::rel(1),
  ...
)
```

Arguments

| | |
|-------------------|--|
| base_size | Base font size. |
| base_family | Font family. Use "sans" for Arial and "serif" for Times New Roman. |
| title_size | Title font size as a relative value. |
| axis_title_size | Axis title font size as a relative value. |
| legend_title_size | Legend title font size as a relative value. |
| legend_text_size | Legend text font size as a relative value. |
| axis_text_size | Axis text font size as a relative value. |
| ... | Additional theme elements to be passed to ggplot2::theme(). |

Value

A ggplot2 theme.

Examples

```
## Not run:
library(ggplot2)
# Using Arial font
ggplot(mtcars, aes(mpg, wt)) + geom_point() + ama_theme()
# Using Times New Roman font
ggplot(mtcars, aes(mpg, wt)) + geom_point() + ama_theme(base_family = "serif")

## End(Not run)
```

plot_coef_par_trends *Plot Coefficients of Parallel Trends*

Description

This function generates coefplots or iplots based on fixest outputs, allowing the user to visualize interaction coefficients with ease.

Usage

```
plot_coef_par_trends(
  data,
  dependent_vars,
  time_var,
  unit_treatment_status,
  unit_id_var,
  plot_type = "coefplot",
  combined_plot = TRUE,
  legend_position = "bottomleft",
  legend_title = "Legend Title",
  legend_args = list(),
  plot_args = list()
)
```

Arguments

| | |
|-----------------------|--|
| data | Data frame containing the data to be used in the model. |
| dependent_vars | Named list of dependent variables to model and their respective labels. |
| time_var | Name of the time variable in the data. |
| unit_treatment_status | Name of the treatment status variable. |
| unit_id_var | Name of the unit identification variable. |
| plot_type | Type of plot to generate. Either "coefplot" or "ipplot". |
| combined_plot | Logical indicating whether to combine plots for all dependent variables. |
| legend_position | Position of the legend on the plot. |
| legend_title | Title for the legend. |
| legend_args | List of additional arguments to customize the legend. |
| plot_args | List of additional arguments to customize the plot. |

Value

A plot visualizing interaction coefficients.

Examples

```
library(fixest)
data("base_did")

# Sample call to the function:
plot_coef_par_trends(
  data = base_did,
  dependent_vars = c(y = "Outcome 1", x1 = "Outcome 2"),
  time_var = "period",
  unit_treatment_status = "treat",
  unit_id_var = "id",
  plot_type = "coefplot",
  combined_plot = TRUE,
  plot_args = list(main = "Interaction coefficients Plot"),
  legend_title = "Metrics",
  legend_position = "bottomleft"
)

plot_coef_par_trends(
  data = base_did,
  dependent_vars = c(y = "Outcome 1", x1 = "Outcome 2"),
  time_var = "period",
  unit_treatment_status = "treat",
  unit_id_var = "id",
  plot_type = "coefplot",
  combined_plot = FALSE
)
```

```
plot_density_by_treatment
```

Plot Density by Treatment

Description

This function creates a list of ggplot density plots for specified variables by treatment groups.

Usage

```
plot_density_by_treatment(
  data,
  var_map,
  treatment_var,
  show_legend = TRUE,
  theme_use = ggplot2::theme_minimal(),
  ...
)
```

Arguments

| | |
|---------|--|
| data | A data frame containing the variables to plot and a treatment variable. |
| var_map | A named list mapping the column names in the data to display names for plotting. |

| | |
|---------------|--|
| treatment_var | A named vector where the name is the treatment column in the data and the value is the legend title. |
| show_legend | A logical value indicating whether to show the legend. Defaults to TRUE. |
| theme_use | ggplot2 theme. Defaults to ggplot2::theme_minimal(). |
| ... | Additional arguments to be passed to geom_density. |

Value

A list of ggplot objects for each variable in var_map.

Examples

```
## Not run:
data(mtcars)
data <- mtcars %>%
  dplyr::select(mpg, cyl) %>%
  dplyr::rowwise() %>%
  dplyr::mutate(treatment = sample(c(0,1), 1, replace = TRUE)) %>%
  dplyr::ungroup()

plots <- plot_density_by_treatment(
  data = data,
  var_map = list("mpg" = "Var 1",
                 "cyl" = "Var 2"),
  treatment_var = c("treatment" = "Treatment Name\nin Legend")
)

## End(Not run)
```

plot_par_trends

Plot Parallel Trends

Description

Plots parallel trends for given metrics.

Usage

```
plot_par_trends(
  data,
  metrics_and_names,
  treatment_status_var,
  time_var,
  conf_level = 0.95,
  non_negative = FALSE,
  display_CI = TRUE,
  output_format = "plot",
  smoothing_method = NULL,
  title_prefix = "Parallel Trends for",
  theme_use = causalverse::ama_theme()
)
```

Arguments

| | |
|-----------------------------------|---|
| <code>data</code> | A data frame containing the data to plot. |
| <code>metrics_and_names</code> | A named list of metrics to plot. |
| <code>treatment_status_var</code> | The variable indicating treatment status. |
| <code>time_var</code> | The variable indicating time. |
| <code>conf_level</code> | Confidence level for confidence intervals (default is 0.95). |
| <code>non_negative</code> | Logical; if TRUE, sets negative lower confidence bounds to 0. |
| <code>display_CI</code> | Logical; if TRUE, displays confidence intervals. |
| <code>output_format</code> | Format of the output; "plot" returns a list of ggplots, "data.frame" returns a data frame. |
| <code>smoothing_method</code> | Method to use for smoothing; NULL means no smoothing. |
| <code>title_prefix</code> | A character string specifying the prefix for the plot title (default is "Parallel Trends for"). |
| <code>theme_use</code> | Custom theme that follows ggplots2 |

Value

A list of ggplot objects or a data frame.

Examples

```
## Not run:
library(tidyverse)
data <- expand.grid(entity = 1:100, time = 1:10) %>%
  dplyr::arrange(entity, time) %>%
  dplyr::mutate(
    treatment = ifelse(entity <= 50, "Treated", "Control"),
    outcome1 = 0.5 * time + rnorm(n(), 0, 2) + ifelse(treatment == "Treated", 0, 0),
    outcome2 = 3 + 0.3 * time + rnorm(n(), 0, 1) + ifelse(treatment == "Treated", 0, 2)
  )
results <- plot_par_trends(
  data = data,
  metrics_and_names = list(outcome1 = "Outcome 1", outcome2 = "Outcome 2"),
  treatment_status_var = "treatment",
  time_var = list(time = "Time"),
  smoothing_method = "loess"
)
library(gridExtra)
gridExtra::grid.arrange(grobs = results, ncol = 1)

## End(Not run)
```

| | |
|-----------------|--|
| plot_treat_time | <i>Plot number of treated units over time or return a dataframe.</i> |
|-----------------|--|

Description

Plot number of treated units over time or return a dataframe.

Usage

```
plot_treat_time(
  data,
  time_var,
  unit_treat,
  outlier_method = "iqr",
  show_legend = FALSE,
  theme_use = causalverse::ama_theme(),
  legend_title = "Point Type",
  legend_labels = c("Regular", "Outlier"),
  regular_size = 3,
  outlier_size = 5,
  regular_color = "black",
  outlier_color = "red",
  regular_shape = 16,
  outlier_shape = 17,
  title = "Random Time Assignment",
  xlab = "Time",
  ylab = "Number of Treated Units",
  output = "plot",
  ...
)
```

Arguments

| | |
|----------------|--|
| data | Dataframe containing data. |
| time_var | Time variable for aggregating the number of treated units. |
| unit_treat | Variable indicating if the unit was treated in a specific time period. |
| outlier_method | Method for outlier detection ("iqr" or "z-score"). |
| show_legend | Logical indicating whether to show legend. |
| theme_use | ggplot2 theme to use. |
| legend_title | Title for legend. |
| legend_labels | Labels for regular and outlier points. |
| regular_size | Size of regular points. |
| outlier_size | Size of outlier points. |
| regular_color | Color of regular points. |
| outlier_color | Color of outlier points. |
| regular_shape | Shape of regular points. |
| outlier_shape | Shape of outlier points. |

| | |
|---------------------|--|
| <code>title</code> | Plot title. |
| <code>xlab</code> | X-axis label. |
| <code>ylab</code> | Y-axis label. |
| <code>output</code> | Type of output ("plot" or "dataframe"). |
| <code>...</code> | Additional arguments to pass to <code>ggplot2::labs</code> . |

Value

`ggplot2` object or dataframe.

Examples

```
# Example usage:
## Not run:
data <- data.frame(time = c(1,1,2,2,3,3), treat = c(0,1,1,1,0,0))
plot_treat_time(data, time_var = time, unit_treat = treat)
plot_treat_time(data, time_var = time, unit_treat = treat, output = "dataframe")

## End(Not run)
```

Index

`ama_export_fig`, [2](#)
`ama_export_tab`, [3](#)
`ama_labs`, [3](#)
`ama_scale_color`, [4](#)
`ama_scale_fill`, [5](#)
`ama_theme`, [6](#)

`plot_coef_par_trends`, [7](#)
`plot_density_by_treatment`, [8](#)
`plot_par_trends`, [9](#)
`plot_treat_time`, [11](#)