

# Package ‘causalverse’

August 25, 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..

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**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)

**VignetteBuilder** knitr

## R topics documented:

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ama_export_fig	<i>Function to export a figure with AMA theme</i>
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### Description

This function applies a custom theme to a ggplot2 figure and exports it to a given path. It exports both an archived version with the current date and a current version without a date.

### Usage

```
ama_export_fig(figure, filename, filepath)
```

### 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.

### 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>
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---

**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 ggplot2::labs().

**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**

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

**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**

<code>data</code>	Data frame containing the data to be used in the model.
<code>dependent_vars</code>	Named list of dependent variables to model and their respective labels.
<code>time_var</code>	Name of the time variable in the data.
<code>unit_treatment_status</code>	Name of the treatment status variable.
<code>unit_id_var</code>	Name of the unit identification variable.
<code>plot_type</code>	Type of plot to generate. Either "coefplot" or "iplot".
<code>combined_plot</code>	Logical indicating whether to combine plots for all dependent variables.
<code>legend_position</code>	Position of the legend on the plot.
<code>legend_title</code>	Title for the legend.
<code>legend_args</code>	List of additional arguments to customize the legend.
<code>plot_args</code>	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,
  theme_use = causalverse::ama_theme(),
  ...
)

```

## Arguments

<code>data</code>	A data frame containing the variables to plot and a treatment variable.
<code>var_map</code>	A named list mapping the column names in the data to display names for plotting.
<code>treatment_var</code>	A named vector where the name is the treatment column in the data and the value is the legend title.
<code>theme_use</code>	ggplot2 theme. Defaults to <code>ggplot2::theme_minimal()</code> .
<code>...</code>	Additional arguments to be passed to <code>geom_density</code> .

## 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	<i>Plot Parallel Trends</i>
-----------------	-----------------------------

---

## 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

data	A data frame containing the data to plot.
metrics_and_names	A named list of metrics to plot.
treatment_status_var	The variable indicating treatment status.
time_var	The variable indicating time.
conf_level	Confidence level for confidence intervals (default is 0.95).
non_negative	Logical; if TRUE, sets negative lower confidence bounds to 0.
display_CI	Logical; if TRUE, displays confidence intervals.
output_format	Format of the output; "plot" returns a list of ggplots, "data.frame" returns a data frame.
smoothing_method	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)
```

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