Package 'causalverse'

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Title Causality in Clarity

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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),
stats (>= 4.2.3),
PanelMatch (>= 2.0.1),
doParallel (>= 1.0.17),
fastDummies (>= 1.7.3),
magrittr (>= 2.0.3),

did (>= 2.1.2),
stats (>= 4.2.3),
foreach (>= 1.5.2)

VignetteBuilder knitr

R topics documented:

`ama_export_fig` *Function to export a figure with custom settings*

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 It exports both an archived version with the current date and a current version without a date.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 *Function to export a table with AMA style*

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 *Custom Label Formatting for ggplot2: American Marketing Association Style*

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 *Custom Color Scale for ggplot2: American Marketing Association Style*

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)
```

`balance_assessment` *Assess balance between treated and control groups*

Description

This function performs a balance assessment between treated and control groups using Seemingly Unrelated Regression (SUR) and Hotelling's T-squared test.

Usage

```
balance_assessment(data, treatment_col, ...)
```

Arguments

`data`	A dataframe containing the data to be assessed.
`treatment_col`	The name of the column that contains the treatment indicator (0 for control, 1 for treated).
`...`	Names of the dependent variables.

Value

A list with two elements: 'SUR' (results of the SUR) and 'Hotelling' (results of the Hotelling's T-squared test).

Examples

```
## Not run:
set.seed(123)
data = mtcars %>%
  dplyr::select(mpg, cyl, disp, hp, wt) %>%
  dplyr::rowwise() %>%
  dplyr::mutate(treatment = sample(c(0,1), 1, replace = TRUE)) %>%
  dplyr::ungroup()

results <- balance_assessment(data, "treatment", "mpg", "cyl")
print(results$SUR)
print(results$Hotelling)

## End(Not run)
```

`balance_scatter_custom` *Custom function to visualize the balance between treatment and control groups*

Description

Custom function to visualize the balance between treatment and control groups

Usage

```
balance_scatter_custom(
  matched_set_list,
  set.names = NULL,
  show.legend = TRUE,
  legend.title = "Type",
  legend.position = "right",
  xlim = c(0, 0.8),
  ylim = c(0, 0.8),
  main = "Standardized Mean Difference of Covariates",
  pchs = NULL,
  dot.size = NULL,
  covariates,
  data,
  x.axis.label = "Before Refinement",
  y.axis.label = "After Refinement",
  theme_use = causalverse::ama_theme(),
  ...
)
```

Arguments

`matched_set_list`	List of matched sets
`set.names`	Vector of names for matched sets. Defaults to NULL.
`show.legend`	Boolean to determine if legend should be shown. Defaults to TRUE.
`legend.title`	Legend title. Defaults to "Type".
`legend.position`	Position of legend. Defaults to "right".
`xlim`	Vector defining x-axis limits. Defaults to c(0, 0.8).
`ylim`	Vector defining y-axis limits. Defaults to c(0, 0.8).
`main`	Main title for the plot. Defaults to "Standardized Mean Difference of Covariates".
`pchs`	Plot characters. Defaults to NULL.
`dot.size`	Size of dots in the scatter plot. Defaults to NULL.
`covariates`	Covariates for calculating balance.
`data`	Dataset for balance calculation.

x.axis.label	x-axis label. Defaults to "Before Refinement".
y.axis.label	y-axis label. Defaults to "After Refinement".
theme_use	Custom theme that follows ggplots2. Defaults to causalverse::ama_theme().
...	Additional arguments passed to the labs() function

Value

ggplot object

Examples

```
## Not run:
library(PanelMatch)
# Maha 4-year lag, up to 5 matches
PM.results.maha.4lag.5m <- PanelMatch::PanelMatch(
   lag = 4,
   time.id = "year",
   unit.id = "wbcode2",
   treatment = "dem",
   refinement.method = "mahalanobis",
   data = PanelMatch::dem,
   match.missing = TRUE,
   covs.formula = ~ I(lag(tradewb, 1:4)) + I(lag(y, 1:4)),
   size.match = 5,
   qoi = "att",
   outcome.var = "y",
   lead = 0:4,
   forbid.treatment.reversal = FALSE,
   use.diagonal.variance.matrix = TRUE
)

# Maha 4-year lag, up to 10 matches
PM.results.maha.4lag.10m <- PanelMatch::PanelMatch(
   lag = 4,
   time.id = "year",
   unit.id = "wbcode2",
   treatment = "dem",
   refinement.method = "mahalanobis",
   data = PanelMatch::dem,
   match.missing = TRUE,
   covs.formula = ~ I(lag(tradewb, 1:4)) + I(lag(y, 1:4)),
   size.match = 10,
   qoi = "att",
   outcome.var = "y",
   lead = 0:4,
   forbid.treatment.reversal = FALSE,
   use.diagonal.variance.matrix = TRUE
)

# Using the function
balance_scatter_custom(
   matched_set_list = list(PM.results.maha.4lag.5m$att, PM.results.maha.4lag.10m$att),
   set.names = c("Maha 4 Lag 5 Matches", "Maha 4 Lag 10 Matches"),
   data = dem,
   covariates = c("y", "tradewb")
)
```

```
## End(Not run)
```

`nice_tab` *Nice Tabulation Function*

Description

Create a custom function that takes a data frame and a number of decimal places as input, rounds all numeric columns in the data frame to the specified number of decimal places, and returns the modified data frame.

Usage

```
nice_tab(data, digit_decimal = 2)
```

Arguments

`data`	A data frame.
`digit_decimal`	A number of decimal places.

Value

A data frame with all numeric columns rounded to the specified number of decimal places.

`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 "iplot".
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

```
## Not run:
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
)

## End(Not run)
```

`plot_covariate_balance_pretrend` *Plot Covariate Balance Over Pre-Treatment Period*

Description

This function visualizes the covariate balance over the pre-treatment period. It's particularly designed for outputs from methods like PanelMatch.

Usage

```
plot_covariate_balance_pretrend(
  balance_data,
  y_limits = c(-1, 1),
  theme_use = causalverse::ama_theme(),
  xlab = "Time to Treatment",
  ylab = "Balance (in SD unit)",
  main_title = "Covariate Balance Over Pre-Treatment Period",
  legend_title = "Covariate",
  show_legend = TRUE,
  ...
)
```

Arguments

`balance_data`	A matrix containing the covariate balance data over the pre-treatment period.
`y_limits`	A numeric vector of length 2 defining the y-axis limits.
`theme_use`	A ggplot2 theme. By default, it uses `causalverse::ama_theme()`.
`xlab`	A string indicating the label for the x-axis.
`ylab`	A string indicating the label for the y-axis.
`main_title`	A string for the main title of the plot.
`legend_title`	A string for the legend title.
`show_legend`	A logical; if TRUE, the legend is displayed, otherwise, it's hidden.
`...`	Additional arguments passed to the ggplot labs.

Value

A ggplot2 object.

Examples

```
## Not run:
  balance_data_sample <- matrix(rnorm(20), nrow = 5)
  plot_covariate_balance_pretrend(balance_data_sample)

## End(Not run)
```

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_PanelEstimate` *Plot Estimated Effects of Treatment Over Time*

Description

This function takes an object (result of PanelEstimate or similar) and plots its estimates over time.

Usage

```
plot_PanelEstimate(
  pe.object,
  ylab = "Estimated Effect of Treatment",
  xlab = "Time Since Treatment",
  main = "Estimated Effects of Treatment Over Time",
  ylim = NULL,
  theme_use = causalverse::ama_theme(),
  ...
)
```

Arguments

`pe.object`	The object with the estimation results.
`ylab`	The y-axis label.
`xlab`	The x-axis label.
`main`	The main title for the plot.
`ylim`	The limits for the y-axis.
`theme_use`	The theme to use for the plot. Defaults to causalverse::ama_theme().
`...`	Additional parameters to pass to labs() function.

Value

A ggplot object with the desired plot.

Examples

```
## Not run:
PM.results.ps.weight <- PanelMatch(lag = 4, time.id = "year", unit.id = "wbcode2",
                                   treatment = "dem", refinement.method = "ps.weight",
                                data = dem, match.missing = FALSE, listwise.delete = TRUE,
                                  covs.formula = ~ I(lag(tradewb, 1:4)) + I(lag(y, 1:4)),
                                   size.match = 5, qoi = "att", outcome.var = "y",
                                   lead = 0:4, forbid.treatment.reversal = FALSE)

PE.results <- PanelEstimate(sets = PM.results.ps.weight,
                            data = dem,
                            se.method = "bootstrap",
                            number.iterations = 1000,
                            confidence.level = .95)

plot_PanelEstimate(PE.results)

## 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

`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.
`title_prefix`	A character string specifying the prefix for the plot title (default is "Parallel Trends for").
`theme_use`	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 *Plot number of treated units over time or return a dataframe.*

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

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)
```

---

plot_trends_across_group

Custom Faceted Line Plot with Optional Standard Error

Description

This function generates a faceted line plot for a given dataset, allowing the user to specify the x-axis, y-axis, grouping variable, and facet variable. Additionally, users can include standard errors and customize labels.

Usage

```
plot_trends_across_group(
  data,
  x_var,
  y_var,
  grouping_var,
  facet_var,
  se = NULL,
  include_legend = TRUE,
  title = "Dependent Variable across Years by Group and Industry",
  x_label = "Year",
  y_label = "Dependent Variable",
  theme = causalverse::ama_theme(),
  ...
)
```

Arguments

data	A data frame containing the data to be plotted.
x_var	A character string specifying the x-axis variable.
y_var	A character string specifying the y-axis variable.
grouping_var	A character string specifying the grouping variable.
facet_var	A character string specifying the facet variable.
se	A character string specifying the standard error variable, or NULL (default) if not provided.
include_legend	Logical. If TRUE, includes the legend, otherwise it does not.
title	Character string specifying the main plot title.
x_label	Character string specifying the x-axis label.
y_label	Character string specifying the y-axis label.
theme	A ggplot2 theme. Defaults to ama_theme.
...	Additional arguments passed to labs.

Value

A ggplot object.

Examples

```
## Not run:
# Create a small sample dataset
sample_data <- data.frame(
  year = rep(2001:2005, each = 2),
  dependent_variable = rnorm(10, mean = 50, sd = 10),
  group = rep(c("treated", "control"), times = 5),
  industry = rep(c("Tech", "Healthcare"), each = 5)
)

# Use the function
plot_trends_across_group(data = sample_data,
                         x_var = "year",
```

```
                        y_var = "dependent_variable",
                        grouping_var = "group",
                        facet_var = "industry",
                        title = "Sample Title")

## End(Not run)
```

`stack_data` *Stacked Data for Staggered DiD Analysis*

Description

`stack_data` processes datasets used in staggered Difference-in-Differences (DiD) designs. Staggered DiD designs arise when different units (e.g., firms, regions, countries) get treated at different time periods. This function creates cohorts based on the provided treatment period variable and stacks them together to create a comprehensive longitudinal format suitable for staggered DiD analyses.

Usage

```
stack_data(
  treated_period_var,
  time_var,
  pre_window,
  post_window,
  data,
  control_type = c("both", "never-treated", "not-yet-treated")
)
```

Arguments

`treated_period_var`	A character string indicating the column name of the treatment period variable.
`time_var`	A character string indicating the column name for time.
`pre_window`	An integer indicating the number of periods before the treatment to consider.
`post_window`	An integer indicating the number of periods after the treatment to consider.
`data`	A data frame containing the dataset to be processed.
`control_type`	A character string indicating which control type to use. One of "both", "never-treated", or "not-yet-treated".

Details

The function emphasizes the importance of having a control group, which should be represented by the value 10000 in the `treated_period_var` column of the provided dataset. The output data will be augmented with relative period dummy variables for ease of subsequent analysis.

Value

A data frame with the stacked data, augmented with relative period dummy variables, suitable for staggered DiD analysis.

Examples

```
## Not run:
  library(did)
  library(tidyverse)
  library(fixest)
  data(base_stagg)
 stacked_data <- stack_data("year_treated", "year", 3, 3, base_stagg, control_type = "both")
  feols_result <- feols(as.formula(paste0(
    "y ~ ",
    paste(paste0("`rel_period_", c(-3:-2, 0:3), "`"), collapse = " + "),
    " | id ^ df + year ^ df"
  )), data = stacked_data)
  print(feols_result)

## End(Not run)
```

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