

Package ‘scmSpillover’

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Type Package

Title Synthetic Control Method with Spillover Effects

Version 0.1.1

Description A general-purpose implementation of synthetic control methods that accounts for potential spillover effects between units. Based on the methodology of Cao and Dowd (2019) <[doi:10.48550/arXiv.1902.07343](https://doi.org/10.48550/arXiv.1902.07343)> ``Estimation and Inference for Synthetic Control Methods with Spillover Effects".

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`generate_test_data` *Generate simulated data for testing*

Description

Generate simulated data for testing

Usage

```
generate_test_data(
  n_units = 20,
  n_periods = 30,
  treatment_start = 20,
  effect_size = -5
)
```

Arguments

<code>n_units</code>	Number of units
<code>n_periods</code>	Number of time periods
<code>treatment_start</code>	When treatment begins
<code>effect_size</code>	Size of treatment effect

Value

A numeric matrix with dimensions (`n_periods` x `n_units`). The first column represents the treated unit, remaining columns are control units. Rows represent time periods.

Examples

```
# Generate test data with default parameters
data <- generate_test_data()
dim(data) # 30 periods x 20 units

# Generate smaller dataset
data <- generate_test_data(n_units = 10, n_periods = 20,
                           treatment_start = 15, effect_size = -3)
```

plot.scm_spillover *Plot method for scm_spillover objects*

Description

Plot method for scm_spillover objects

Usage

```
## S3 method for class 'scm_spillover'
plot(x, type = "effects", ...)
```

Arguments

x	An object of class scm_spillover
type	Type of plot: "effects" or "all"
...	Additional arguments passed to plotting functions

Value

A ggplot object (invisibly for type="all")

plot_all *Generate all plots for SCM analysis*

Description

Generate all plots for SCM analysis

Usage

```
plot_all(
  result,
  start_year = NULL,
  unit_name = "Treated Unit",
  outcome_label = "Outcome",
  treatment_label = "Treatment",
  show_ci = TRUE
)
```

Arguments

<code>result</code>	Output from <code>run_scm_spillover</code>
<code>start_year</code>	First year of treatment period
<code>unit_name</code>	Name of treated unit (default "Treated Unit")
<code>outcome_label</code>	Label for outcome variable (default "Outcome")
<code>treatment_label</code>	Label for treatment (default "Treatment")
<code>show_ci</code>	Logical, whether to show confidence bands (default TRUE)

Value

A list containing three ggplot objects (`p1`, `p2`, `p3`), returned invisibly. Called primarily for side effects (displaying plots).

Examples

```
## Not run:
data <- generate_test_data()
result <- run_scm_spillover(data, treatment_start = 20)
plots <- plot_all(result)

## End(Not run)
```

`plot_effects` *Plot treatment effects only*

Description

Plot treatment effects only

Usage

```
plot_effects(
  result,
  start_year = NULL,
  unit_name = "Treated Unit",
  outcome_label = "Outcome",
  show_ci = TRUE,
  show_vanilla = FALSE
)
```

Arguments

result	Output from run_scm_spillover
start_year	First year of treatment period
unit_name	Name of treated unit (default "Treated Unit")
outcome_label	Label for outcome variable (default "Outcome")
show_ci	Logical, whether to show confidence bands (default TRUE)
show_vanilla	Logical, whether to show vanilla SCM comparison (default FALSE)

Value

A ggplot object that can be further customized or saved.

Examples

```
data <- generate_test_data()
result <- run_scm_spillover(data, treatment_start = 20, verbose = FALSE)
p <- plot_effects(result)
```

print.scm_spillover *Print method for scm_spillover objects*

Description

Print method for scm_spillover objects

Usage

```
## S3 method for class 'scm_spillover'
print(x, ...)
```

Arguments

x	An object of class scm_spillover
...	Additional arguments (not used)

Value

Invisibly returns the input object

qplot_all	<i>Quick plot all three main plots in sequence</i>
-----------	--

Description

Quick plot all three main plots in sequence

Usage

```
qplot_all(
  result,
  start_year = NULL,
  unit_name = "Treated Unit",
  outcome_label = "Outcome",
  treatment_label = "Treatment",
  show_ci = TRUE,
  pause = TRUE
)
```

Arguments

result	Output from run_scm_spillover
start_year	First year of treatment period
unit_name	Name of treated unit (default "Treated Unit")
outcome_label	Label for outcome variable (default "Outcome")
treatment_label	Label for treatment (default "Treatment")
show_ci	Logical, whether to show confidence bands (default TRUE)
pause	Logical, whether to pause between plots (default TRUE)

Value

List of ggplot objects (invisibly)

qplot_ci	<i>Quick plot with confidence intervals only</i>
----------	--

Description

Quick plot with confidence intervals only

Usage

```
qplot_ci(  
  result,  
  start_year = NULL,  
  unit_name = "Treated Unit",  
  outcome_label = "Outcome"  
)
```

Arguments

result	Output from run_scm_spillover
start_year	First year of treatment period
unit_name	Name of treated unit (default "Treated Unit")
outcome_label	Label for outcome variable (default "Outcome")

Value

ggplot object (invisibly)

qplot_compare

Quick plot for method comparison (Spillover vs Vanilla)

Description

Quick plot for method comparison (Spillover vs Vanilla)

Usage

```
qplot_compare(  
  result,  
  start_year = NULL,  
  unit_name = "Treated Unit",  
  outcome_label = "Outcome"  
)
```

Arguments

result	Output from run_scm_spillover
start_year	First year of treatment period
unit_name	Name of treated unit (default "Treated Unit")
outcome_label	Label for outcome variable (default "Outcome")

Value

ggplot object (invisibly)

qplot_effects*Quick plot for treatment effects with confidence intervals***Description**

Quick plot for treatment effects with confidence intervals

Usage

```
qplot_effects(
  result,
  start_year = NULL,
  unit_name = "Treated Unit",
  outcome_label = "Outcome",
  show_ci = TRUE,
  show_vanilla = FALSE
)
```

Arguments

<code>result</code>	Output from <code>run_scm_spillover</code>
<code>start_year</code>	First year of treatment period
<code>unit_name</code>	Name of treated unit (default "Treated Unit")
<code>outcome_label</code>	Label for outcome variable (default "Outcome")
<code>show_ci</code>	Logical, whether to show confidence bands (default TRUE)
<code>show_vanilla</code>	Logical, whether to show vanilla SCM comparison (default FALSE)

Value

`ggplot` object (invisibly)

qplot_point*Quick plot without confidence intervals***Description**

Quick plot without confidence intervals

Usage

```
qplot_point(
  result,
  start_year = NULL,
  unit_name = "Treated Unit",
  outcome_label = "Outcome"
)
```

Arguments

result	Output from run_scm_spillover
start_year	First year of treatment period
unit_name	Name of treated unit (default "Treated Unit")
outcome_label	Label for outcome variable (default "Outcome")

Value

ggplot object (invisibly)

qplot_series*Quick plot for actual vs synthetic control time series*

Description

Quick plot for actual vs synthetic control time series

Usage

```
qplot_series(  
  result,  
  start_year = NULL,  
  unit_name = "Treated Unit",  
  outcome_label = "Outcome",  
  treatment_label = "Treatment"  
)
```

Arguments

result	Output from run_scm_spillover
start_year	First year of treatment period
unit_name	Name of treated unit (default "Treated Unit")
outcome_label	Label for outcome variable (default "Outcome")
treatment_label	Label for treatment (default "Treatment")

Value

ggplot object (invisibly)

`run_scm_spillover` *Run Complete SCM Analysis with Spillover Effects*

Description

Run Complete SCM Analysis with Spillover Effects

Usage

```
run_scm_spillover(
  data,
  treatment_start,
  treated_unit = 1,
  affected_units = NULL,
  verbose = TRUE
)
```

Arguments

<code>data</code>	Matrix or data frame (time x units)
<code>treatment_start</code>	Integer, first treatment period (row number)
<code>treated_unit</code>	Integer, column index of treated unit (default = 1)
<code>affected_units</code>	Vector of column indices for all affected units (including treated)
<code>verbose</code>	Logical, print progress

Value

List with all results

`safe_solve` *Safe matrix inversion with regularization*

Description

Safely inverts a matrix using regularization if needed

Usage

```
safe_solve(M, tol = 1e-10)
```

Arguments

<code>M</code>	Matrix to invert
<code>tol</code>	Tolerance for condition number check

Value

Inverted matrix

save_all_plots	<i>Save all plots to files</i>
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Description

Save all plots to files

Usage

```
save_all_plots(  
  result,  
  prefix = "scm",  
  path = NULL,  
  width = 10,  
  height = 6,  
  dpi = 300,  
  ...  
)
```

Arguments

result	Output from run_scm_spillover
prefix	File name prefix (default "scm")
path	Directory to save plots (required, no default)
width	Plot width in inches (default 10)
height	Plot height in inches (default 6)
dpi	Resolution (default 300)
...	Additional arguments passed to plotting functions

Value

Character vector of saved file paths, returned invisibly.

Examples

```
## Not run:  
data <- generate_test_data()  
result <- run_scm_spillover(data, treatment_start = 20)  
save_all_plots(result, path = tempdir())  
  
## End(Not run)
```

scm*Basic SCM function for single unit***Description**

Computes synthetic control weights for a single treated unit

Usage

```
scm(Y, lambda = 1e-06)
```

Arguments

<i>Y</i>	N x T matrix where first row is treated unit
<i>lambda</i>	Regularization parameter for numerical stability

Value

List containing intercept (a) and weights (b)

scm_batch*Batch SCM for all units***Description**

Computes synthetic control weights for all units

Usage

```
scm_batch(Y, verbose = FALSE)
```

Arguments

<i>Y</i>	N x T matrix of outcomes
<i>verbose</i>	Print progress messages

Value

List with intercepts (a) and weight matrix (B)

sp_andrews_spillover *Test for spillover effects*

Description

Tests whether spillover effects are significant

Usage

```
sp_andrews_spillover(Y0, Y1, A, alpha_sig = 0.05)
```

Arguments

Y0	N x T pre-treatment matrix
Y1	N x 1 post-treatment vector
A	N x k spillover structure matrix
alpha_sig	Significance level

Value

List with test results

sp_andrews_te *Andrews test for treatment effects*

Description

Tests significance of treatment effects

Usage

```
sp_andrews_te(Y0, Y1, A, C = NULL, alpha_sig = 0.05)
```

Arguments

Y0	N x T pre-treatment matrix
Y1	N x 1 post-treatment vector (single period)
A	N x k spillover structure matrix
C	Constraint matrix (default tests first unit)
alpha_sig	Significance level

Value

List with estimates, p-values, and confidence intervals

sp_estimation	<i>Spillover-adjusted SCM estimation</i>
---------------	--

Description

Estimates treatment effects accounting for spillovers

Usage

```
sp_estimation(Y_pre, Y_post, A, verbose = FALSE)
```

Arguments

Y_pre	N x T pre-treatment matrix
Y_post	N x S post-treatment matrix
A	N x k spillover structure matrix
verbose	Print progress messages

Value

List containing treatment effects and other results

summary.scm_spillover	<i>Summary method for scm_spillover objects</i>
-----------------------	---

Description

Summary method for scm_spillover objects

Usage

```
## S3 method for class 'scm_spillover'
summary(object, ...)
```

Arguments

object	An object of class scm_spillover
...	Additional arguments (not used)

Value

Invisibly returns a data frame with detailed results

vanilla_scm	<i>Standard SCM without spillovers</i>
-------------	--

Description

Computes standard synthetic control for comparison

Usage

```
vanilla_scm(Y_pre, Y_post)
```

Arguments

Y_pre	N x T pre-treatment matrix
Y_post	N x S post-treatment matrix

Value

Synthetic control values for all periods

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