

# Package ‘MRStdLCRT’

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**Type** Package

**Title** Model-Robust Standardization for Longitudinal Cluster-Randomized Trials

**Version** 0.1.0

**Description** Provides estimation and leave-one-cluster-out jackknife standard errors for four longitudinal cluster-randomized trial estimands: horizontal individual average treatment effect (h-iATE), horizontal cluster average treatment effect (h-cATE), vertical individual average treatment effect (v-iATE), and vertical cluster-period average treatment effect (v-cATE), using unadjusted and augmented (model-robust standardization) estimators. The working model may be fit using linear mixed models for continuous outcomes or generalized estimating equations and generalized linear mixed models for binary outcomes. Period inclusion for aggregation is determined automatically: only periods with both treated and control clusters are included in the construction of the marginal means and treatment effect contrasts. See Fang et al. (2025) <[doi:10.48550/arXiv.2507.17190](https://doi.org/10.48550/arXiv.2507.17190)>.

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**Encoding** UTF-8

**LazyData** true

**Depends** R (>= 4.1.0)

**Imports** reformulas, dplyr (>= 1.1.0), tidyr (>= 1.3.0), rlang (>= 1.1.0), tidyselect, gee, lme4 (>= 1.1-30), ggplot2 (>= 3.4.0), stats, utils

**Suggests** testthat (>= 3.0.0)

**Config/testthat/edition** 3

**RoxxygenNote** 7.3.3

**NeedsCompilation** no

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**Repository** CRAN

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### *mrstdlcrt\_fit*

*Fit model-robust standardization for longitudinal CRTs*

---

#### Description

Fit model-robust standardization for longitudinal CRTs

#### Usage

```
mrstdlcrt_fit(
  data,
  formula,
  cluster_id = "cluster",
  period = "period",
  trt = "trt",
  method = c("gee", "lmer", "glmer"),
  family = c("gaussian", "binomial"),
  corstr = "independence",
  scale = c("RD", "RR", "OR")
)
```

#### Arguments

|                   |  |
|-------------------|--|
| <i>data</i>       | data.frame with outcome, treatment, period, cluster, covariates. |
| <i>formula</i>    | model formula; may include interactions and random effects.      |
| <i>cluster_id</i> | cluster id column name.  |
| <i>period</i>     | period column name.  |
| <i>trt</i>        | treatment column name (0/1).                                     |
| <i>method</i>     | "gee", "lmer", "glmer".  |
| <i>family</i>     | "gaussian", "binomial".  |
| <i>corstr</i>     | gee correlation.   |
| <i>scale</i>      | For binomial: "RD", "RR", "OR" (RR/OR are on log scale).         |

**Value**

Object of class "mrs".

---

**plot.mrs**

*Plot method for mrs objects*

---

**Description**

Plot method for mrs objects

**Usage**

```
## S3 method for class 'mrs'  
plot(x, level = 0.95, estimand = NULL, point_size = 2.8, ...)
```

**Arguments**

|            |                              |
|------------|------------------------------|
| x          | An object of class "mrs".    |
| level      | Confidence level.            |
| estimand   | Subset of estimands to plot. |
| point_size | Point size.                  |
| ...        | Unused.                      |

**Value**

ggplot object invisibly.

---

**print.mrs**

*Print method for mrs objects*

---

**Description**

Print method for mrs objects

**Usage**

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

**Arguments**

|     |                           |
|-----|---------------------------|
| x   | An object of class "mrs". |
| ... | Unused.                   |

**Value**

x invisibly.

`summary.mrs`      *Summarize an mrs fit*

### Description

Summarize an mrs fit

### Usage

```
## S3 method for class 'mrs'
summary(
  object,
  level = 0.95,
  estimand = NULL,
  digits = 6,
  show_counts = TRUE,
  ...
)
```

### Arguments

|                          |                               |
|--------------------------|-------------------------------|
| <code>object</code>      | An object of class "mrs".     |
| <code>level</code>       | Confidence level.             |
| <code>estimand</code>    | Optional subset of estimands. |
| <code>digits</code>      | Digits to print.              |
| <code>show_counts</code> | Print counts tables.          |
| ...                      | Unused.                       |

### Value

Invisibly returns a list of printed tables and metadata.

`sw_b`      *Example stepped wedge CRT dataset with binary outcome*

### Description

A toy dataset with cluster, period, and individual records for illustrating estimands in stepped wedge CRT with a binary outcome.

### Usage

```
data(sw_b)
```

## Format

A data frame with columns:

- cluster** Cluster identifier (integer).
- period** Period index (integer).
- id** Individual identifier within cluster-period (integer).
- trt** Treatment indicator (0/1).
- x1** Auxiliary covariate (0/1).
- x2** Auxiliary covariate (numeric).
- y** Outcome (0/1, binary).

## Examples

```
data(sw_b)
head(sw_b)
```

---

sw\_c

*Example of stepped wedge CRT dataset for continuous outcome*

---

## Description

A toy dataset with cluster, period, and individual records for illustrating estimands in stepped wedge CRT with a continuous outcome.

## Usage

```
data(sw_c)
```

## Format

A data frame with columns:

- cluster** Cluster identifier (integer).
- period** Period index (integer).
- id** Individual identifier within cluster-period (integer).
- trt** Treatment indicator (0/1).
- x1** Auxiliary covariate (0/1).
- x2** Auxiliary covariate (numeric).
- y** Outcome (numeric, continuous).

## Examples

```
data(sw_c)
head(sw_c)
```

---

**xo\_b***Example crossover cluster-randomized trial dataset with binary outcome*

---

**Description**

A small simulated 2×2 crossover trial dataset with a binary outcome.

**Usage**

```
xo_b
```

**Format**

A tibble/data.frame with one row per subject and the following columns:

- h** Integer cluster ID (hospital)
- p** Integer period (1 or 2)
- k** Integer subject index within cluster-period
- trt** Treatment indicator (0 = control, 1 = treatment)
- x\_c01, x\_c02** Continuous covariates
- x\_b01** Binary covariate (0/1)
- x\_cat1\_2, x\_cat1\_3** Dummy variables for a 3-level categorical covariate (level 1 is reference)
- y\_bin** Observed binary outcome (0/1)

**Examples**

```
data(xo_b)
str(xo_b)
head(xo_b)
```

---

**xo\_c***Example crossover cluster-randomized trial dataset with continuous outcome*

---

**Description**

A small simulated 2×2 crossover trial dataset with a continuous outcome.

**Usage**

```
xo_c
```

## Format

A tibble/data.frame with one row per subject and the following columns:

- h** Integer cluster ID (hospital)
- p** Integer period (1 or 2)
- k** Integer subject index within cluster-period
- trt** Treatment indicator (0 = control, 1 = treatment)
- x\_c01, x\_c02** Continuous covariates
- x\_b01** Binary covariate (0/1)
- x\_cat1\_2, x\_cat1\_3** Dummy variables for a 3-level categorical covariate (level 1 is reference)
- y\_cont** Observed continuous outcome

## Examples

```
data(xo_c)
str(xo_c)
head(xo_c)
```

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