

# Package: MaximInfer (via r-universe)

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**Type** Package

**Title** Inference for Maximin Effects in High-Dimensional Settings

**Version** 2.1.0

**Description** Implementation of the sampling and aggregation method for the covariate shift maximin effect, which was proposed in [doi:10.48550/arXiv.2011.07568](https://doi.org/10.48550/arXiv.2011.07568). It constructs the confidence interval for any linear combination of the high-dimensional maximin effect.

**License** GPL-3

**Encoding** UTF-8

**RoxygenNote** 7.2.3

**Suggests** knitr, rmarkdown

**Imports** MASS, stats, CVXR (>= 1.8), glmnet, intervals, SIHR

**Depends** R (>= 4.3.0)

**NeedsCompilation** no

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**Config/pak/sysreqs** cmake libgmp3-dev make pkg-config libclang-dev

**Repository** <https://anonymous-researcher-123243.r-universe.dev>

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## Contents

|                               |   |
|-------------------------------|---|
| decide_delta . . . . .        | 2 |
| Infer . . . . .               | 2 |
| Maximin . . . . .             | 3 |
| measure_instability . . . . . | 5 |

|              |          |
|--------------|----------|
| <b>Index</b> | <b>6</b> |
|--------------|----------|

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|              |  |
|--------------|--|
| decide_delta | <i>Decide ridge penalty data-dependently</i> |
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### Description

To tell if the estimator is stable or not without ridge penalty at first. If instable, it picks a ridge penalty data-dependently.

### Usage

```
decide_delta(
  obj,
  gen.size = 500,
  step_delta = 0.1,
  MAX_iter = 100,
  verbose = FALSE
)
```

### Arguments

|            |  |
|------------|--|
| obj        | The returned list of Maximin                               |
| gen.size   | The generating sample size (Default = 500)                 |
| step_delta | The step size of searching delta (Default = 0.1)           |
| MAX_iter   | Maximum of iterations for searching (Default = 100)        |
| verbose    | Print information about delta and reward (Default = FALSE) |

### Value

|              |   |
|--------------|---|
| delta        | The data-dependent ridge penalty                        |
| reward.ratio | The ratio of penalized reward over non-penalized reward |

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|       |                         |
|-------|-------------------------|
| Infer | <i>Inference method</i> |
|-------|-------------------------|

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### Description

Given the returned list of Maximin, compute the Point estimator and Confidence interval.

**Usage**

```
Infer(
  obj,
  delta = 0,
  gen.size = 500,
  threshold = 0,
  alpha = 0.05,
  alpha.thres = 0.01
)
```

**Arguments**

|             |  |
|-------------|--|
| obj         | returned list of Maximin   |
| delta       | The ridge penalty (Default = 0)  |
| gen.size    | The generating sample size (Default = 500)   |
| threshold   | Should generated samples be filtered or not? if 0, use normal threshold to filter; if 1, use chi-square threshold to filter; if 2, do not filter (Default = 0) |
| alpha       | confidence value to construct confidence interval (Default = 0.05)   |
| alpha.thres | confidence value to select generated samples (Default = 0.01)  |

**Value**

|           |  |
|-----------|--|
| weight    | The weight vector for groups, of length $L$  |
| mm.effect | The aggregated maximin effect (coefficients), of length $p$ or $p + 1$                     |
| mminfer   | The list of length $n.loading$ , each contains the point estimator and confidence interval |

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Maximin

*Returns a list that provides materials for later inference method.*


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**Description**

Given list of observations, compute the bias-corrected initial estimators and do bias-correction to the regressopm covariance matrix.

**Usage**

```
Maximin(
  Xlist,
  Ylist,
  loading.mat,
  X0 = NULL,
  cov.shift = TRUE,
  cov0 = NULL,
  intercept = TRUE,
```

```

    intercept.loading = FALSE,
    lambda = NULL,
    verbose = FALSE
)

```

### Arguments

|                                |   |
|--------------------------------|---|
| <code>Xlist</code>             | list of design matrix for source data, of length $L$  |
| <code>Ylist</code>             | list of outcome vector for source data, of length $L$   |
| <code>loading.mat</code>       | Loading matrix, of dimension $n.loading \times p$ , each column corresponds to a loading of interest            |
| <code>X0</code>                | design matrix for target data, of dimension $n0 \times p$ (default = NULL)                                      |
| <code>cov.shift</code>         | Covariate shifts or not between source and target data (default = TRUE)   |
| <code>cov0</code>              | Covariance matrix for target data, of dimension $p \times p$ (default = NULL)                                   |
| <code>intercept</code>         | Should intercept be fitted for the initial estimator (default = TRUE)   |
| <code>intercept.loading</code> | Should intercept term be included for the loading (default = FALSE)   |
| <code>lambda</code>            | The tuning parameter in fitting initial model. If NULL, it will be picked by cross-validation. (default = NULL) |
| <code>verbose</code>           | Should intermediate message(s) be printed. (default = FALSE)  |

### Details

The algorithm implemented scenarios with or without covariate shift. If `cov0` is specified, the `X0` will be ignored; if not, while `X0` is specified, `cov0` will be estimated by `X0`. If both are not specified, the algorithm will automatically set `cov.shift` as FALSE.

### Value

The returned list contains the following components:

|                           |   |
|---------------------------|---|
| <code>Gamma.plugin</code> | The plugin regression covariance matrix   |
| <code>Gamma.debias</code> | The proposed debiased regression covariance matrix  |
| <code>Var.Gamma</code>    | The variance matrix for sampling the regression covariance matrix   |
| <code>fits.info</code>    | The list of length $L$ , that contains the initial coefficient estimators and variance of fitted residuals.                         |
| <code>Points.info</code>  | The list of length $L$ , that contains the initial debiased estimator for linear combinations and its corresponding standard error. |

### Examples

```

L = 2
n1 = n2 = 100; p = 4
X1 = MASS::mvrnorm(n1, rep(0,p), Sigma=diag(p))
X2 = MASS::mvrnorm(n2, rep(0,p), Sigma=0.5*diag(p))
b1 = seq(1,4)/10; b2 = rep(0.2, p)

```

```
y1 = as.vector(X1*%b1+rnorm(n1)); y2 = as.vector(X2*%b2+rnorm(n2))
loading1 = rep(0.4, p)
loading2 = c(-0.5, -0.5, rep(0,p-2))
loading.mat = cbind(loading1, loading2)
cov0 = diag(p)
mm = Maximin(list(X1,X2),list(y1,y2),loading.mat,cov0=cov0)

# inference
out = Infer(mm, gen.size=10)
```

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measure\_instability    *measurement of instability*

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## Description

compute the instability measurement given a specific ridge penalty

## Usage

```
measure_instability(
  obj,
  delta = 0,
  gen.size = 500,
  threshold = 0,
  alpha.thres = 0.01
)
```

## Arguments

|             |   |
|-------------|---|
| obj         | The returned list of Maximin  |
| delta       | The ridge penalty (Default = 0)   |
| gen.size    | The generating sample size (Default = 500)  |
| threshold   | Should generated samples be filtered or not? if 0, use normal threshold to filter; if 1, use chi-square threshold to filter; if 2, do not filter. (Default = 0) |
| alpha.thres | The confidence value to select generated samples (Default = 0.01)   |

## Value

The measurement of instability

# Index

decide\_delta, [2](#)

Infer, [2](#)

Maximin, [3](#)

measure\_instability, [5](#)