Package 'MaximinInfer'

October 18, 2021

Title Inference for Maximin Effect in high-dimensional settings

Type Package

Version 0.1.0
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Description Provides functionality for the paper. The function is used to compute the bias corrected estimator of ridge-penalized maximin effect and the point estimator of its linear contrast. It also constructs the confidence interval for the linear contrast.
License GPL-3
Encoding UTF-8
LazyData true
RoxygenNote 7.1.1
Suggests knitr, rmarkdown
VignetteBuilder knitr
Imports MASS, stats, scalreg, flare
Depends CVXR, glmnet, intervals, SIHR
R topics documented:
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Description

'infer' is a generic function for inference for Maximin model.

Usage

```
infer(object, ...)
```

Arguments

object a "Maximin" object
... additional arguments affecting inference

infer.Maximin

Inference method for Maximin

Description

Point estimator and Confidence interval based on Maximin object

Usage

```
## S3 method for class 'Maximin'
infer(object, gen.size = 500, delta = -1, threshold = 2, alpha = 0.01, ...)
```

Arguments

object	Object of class inheriting from "Maximin"
gen.size	The generated sample size
delta	The ridge penalty. If set as negative value, the penalty is decided data-dependently. (Default $= -1$)
threshold	Should generated samples be filter or not? If 0, do not filter; if 1, use chi-square threshold to filter; if 2, use normal threshold to filter. (Default = 2)
alpha	confidence value to select generated samples
	further arguments passed

Value

delta	The ridge penalty used
weight	The weight vector for groups, of length ${\cal L}$
point	The point estimator of the linear contrast
mm.effect	The aggregated maximin effect (coefficients), of length p or $p+1$
CI	Confidence interval for the linear contrast

Maximin 3

Examples

```
## number of groups
L=2
## dimension
p=500
## sample size for each group of source data
ns.source = c(500, 400)
## sample size for target data
n.target=1000
A1gen <- function(rho,p){
  A1=matrix(0,p,p)
  for(i in 1:p){
    for(j in 1:p){
      A1[i,j] < -rho^(abs(i-j))
  }
  return(A1)
}
## mean vector
mean.source = rep(0, p)
mean.target = rep(0, p)
## covariate shifts
cov.source = A1gen(0.6, p)
cov.target = diag(p)
## true coefficients
Bs = matrix(0, p, L)
Bs[1:10,1] = seq(1:10)/40
Bs[1:10,2] = -seq(1:10)/40
## Data
X.source = MASS::mvrnorm(sum(ns.source), mu=mean.source, Sigma=cov.source)
X.target = MASS::mvrnorm(n.target, mu=mean.target, Sigma=cov.target)
idx.source = rep(1:L, times=ns.source)
Y.source = rep(0, sum(ns.source))
for(1 in 1:L){
  idx.l = which(idx.source==1)
  Y.source[idx.l] = X.source[idx.l, ] %*% Bs[,l] + rnorm(ns.source[l])
loading = rep(0, p)
loading[1:5] = 1
mm <- Maximin(X.source, Y.source, idx.source, loading, X.target, covariate.shift = TRUE)</pre>
mmInfer <- infer(mm, gen.size=100, delta=0)</pre>
```

Maximin

Class Maximin

Description

Class Maximin

Usage

Maximin(

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```
X.source,
Y.source,
idx.source,
loading,
X.target = NULL,
cov.target = NULL,
covariate.shift = TRUE,
lam.value = c("CV", "CV.min", "scalreg", "slim"),
intercept = TRUE,
intercept.loading = FALSE
)
```

Arguments

Design matrix for source data, of dimension $n.source \times p$ X.source Y.source Outcome vector for source data, of length n.sourceIndicator vector of groups for source data, of length n.sourceidx.source loading Loading, of length p X.target Design matrix for target data, of dimension $n.target \times p$ (default = 'NULL') Covariance matrix for target data, of dimension $p \times p$. If set as 'NULL', 'cov.target' cov.target is unknown. (default = 'NULL') covariate.shift Covariate shifts or not between source and target data (default = 'TRUE") lam.value The method to be used to obtain each group's intial estimator intercept Should intercept be fitted for the initial estimator (default = 'TRUE')

intercept.loading

Should intercept be included for the loading (default = 'FALSE')

Value

'Maximin' returns an object of class "Maximin". The function 'infer' is used to do further inference.\ An object of class "Maximin" is a list containing the following components.

Gamma.prop The proposed debiased Weight matrix
Coef.est The initial estimators for each group
Point.vec The point estimator for each group

L The number of groups

gen.mu The mean vector for sampling method gen.Cov The variance matrix for sampling method

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