Package ‘space’

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space.joint

A function to estimate partial correlations using the Joint Sparse Regression Model

Description

A function to estimate partial correlations using the Joint Sparse Regression Model

Usage

space.joint(Y.m, lam1, lam2=0, sig=NULL, weight=NULL, iter=2)
Arguments

- **Y.m**: numeric matrix. Columns are for variables and rows are for samples. Missing values are not allowed. It's recommended to first standardize each column to have mean 0 and $l_2$ norm 1.

- **lam1**: numeric value. This is the $l_1$ norm penalty parameter. If the columns of Y.m have norm one, then the suggested range of lam1 is: $O(n^{3/2}p^{-1}(1 - \alpha/(2p^2)))$ for small $\alpha$ such as 0.1.

- **lam2**: numeric value. If not specified, lasso regression is used in the Joint Sparse Regression Model (JSRM). Otherwise, elastic net regression is used in JSRM and lam2 serves as the $l_2$ norm penalty parameter.

- **sig**: numeric vector. Its length should be the same as the number of columns of Y.m. It is the vector of $\sigma_{ii}$ (the diagonal of the inverse covariance matrix). If not specified, $\sigma_{ii}$ will be estimated during the model fitting with initial values rep(1,p). The number of the iteration of the model fitting (iter) will then be at least 2. Note, the scale of sig does not matter.

- **weight**: numeric value or vector. It specifies the weights or the type of weights used for each regression in JSRM. The default value is NULL, which means all regressions will be weighted equally in the joint model. If weight=1, residue variances will be used for weights. If weight=2, the estimated degree of each variable will be used for weights. Otherwise, it should be a positive numeric vector, whose length is equal to the number of columns of Y.m.

- **iter**: integer. It is the total number of interactions in JSRM for estimating $\sigma_{ii}$ and partial correlations. When sig= NULL and/or weight= NULL or 2, iter should be at least 2.

Details

space.joint uses a computationally efficient approach for selecting non-zero partial correlations under the high-dimension-low-sample-size setting (Peng and et.al., 2007).

Value

A list with two components

- **ParCor**: the estimated partial correlation matrix.
- **sig.fit**: numeric vector of the estimated diagonal $\sigma_{ii}$.

Author(s)

J. Peng, P. Wang, Nengfeng Zhou, Ji Zhu

References


Examples

(A) The simulated Hub.net example in Peng et al. (2007).

```r
data(spaceSimu)

n=nrow(spaceSimu$y.data)
p=ncol(spaceSimu$y.data)
true.adj=abs(spaceSimu$ParCor.true)>1e-6

view the network corresponding to the parcial correlation matrix in the simulation example
the following code can run only if the "igraph" is installed in the system.

library(igraph)
plot.adj=true.adj
diag(plot.adj)=0
temp=graph.adjacency(adjmatrix=plot.adj, mode="undirected")
temp.degree=apply(plot.adj, 2, sum)
V(temp)$color=(temp.degree>9)+3
plot(temp, vertex.size=3, vertex.frame.color="white", layout=layout.fruchterman.reingold, vertex.label=NA, edge

estimate the parcial correlation matrix with various methods
alpha=1
l1=1/sqrt(n)*qnorm(1-alpha/(2*p^2))
iter=3

the values of lam1 were selected to make the results of different methods comparable.

1. MB method
result1=space.neighbor(spaceSimu$y.data, lam1=l1*x0.7, lam2=0)
fit.adj=abs(result1$ParCor)>1e-6
sum(fit.adj==1)/2 #total number of edges detected
sum(fit.adj[true.adj==1]==1)/2 #total number of true edges detected

2. Joint method with no weight
result2=space.joint(spaceSimu$y.data, lam1=l1*x1.56, lam2=0, iter=iter)
fit.adj=abs(result2$ParCor)>1e-6
sum(fit.adj==1)/2 #total number of edges detected
sum(fit.adj[true.adj==1]==1)/2 #total number of true edges detected

3. Joint method with residue variance based weights
result3=space.joint(spaceSimu$y.data, lam1=l1*x1.86, lam2=0, weight=1, iter=iter)
fit.adj=abs(result3$ParCor)>1e-6
sum(fit.adj==1)/2 #total number of edges detected
sum(fit.adj[true.adj==1]==1)/2 #total number of true edges detected

4. Joint method with degree based weights
result4=space.joint(spaceSimu$y.data, lam1=l1*x1.61, lam2=0, weight=2, iter=iter)
fit.adj=abs(result4$ParCor)>1e-6
sum(fit.adj==1)/2 #total number of edges detected
sum(fit.adj[true.adj==1]==1)/2 #total number of true edges detected
```
space.neighbor  A function to estimate partial correlations using the neighborhood selection approach

Description

A function to estimate partial correlations using the neighborhood selection approach

Usage

space.neighbor(Y.m, lam1, lam2=0)

Arguments

Y.m numeric matrix. Each column is for one variable and each row is for one sample. Missing values are not allowed. It’s recommended to first standardize each column to have mean 0 and norm 1.

lam1 numeric value. This is the $l_1$ norm penalty parameter. If the columns of Y.m have norm one, then the suggested range of lam1 is: $O(n^{1/2} \Phi^{-1}(1 - \alpha/(2p^2)))$ for small $\alpha$ such as 0.1.

lam2 numeric value. If not specified, lasso regression is used in the neighborhood selection. Otherwise, elastic net regression is used and lam2 serves as the $l_2$ norm penalty parameter.

Details

space.neighbor estimate partial correlations using the neighborhood selection approach (Meinshausen and Buhlmann, 2006).

Value

A list with two components

ParCor the estimated partial correlation matrix.

sig.fit numeric vector of the estimated $\sigma_{ii}$

Author(s)

J. Peng, P. Wang, N. Zhou, J. Zhu

References


Examples

# The simulated Hub.net example in Peng et. al. (2007).
data(spaceSimu)

n = nrow(spaceSimu$Y.data)
p = ncol(spaceSimu$Y.data)
true.adj = abs(spaceSimu$ParCor.true) > 1e-6

# view the network corresponding to the parcial correlation matrix in the simulation example
# view the network corresponding to the parcial correlation matrix in the simulation example
# the following code can run only if the "igraph" is installed in the system.
library(igraph)
plot.adj = true.adj
diag(plot.adj) = 0

# estimate the parcial correlation matrix with various methods
alpha = 1
l1 = 1/sqrt(n) * qnorm(1 - alpha / (2 * p^2))
iter = 3

# the values of lam1 were selected to make the results of different methods comparable.
### 1. MB method
result1 = space.neighbor(spaceSimu$Y.data, lam1 = l1 * 0.7, lam2 = 0)
fit.adj = abs(result1$ParCor) > 1e-6
sum(fit.adj == 1) / 2 # total number of edges detected
sum(fit.adj[true.adj == 1] == 1) / 2 # total number of true edges detected

### 2. Joint method with no weight
result2 = space.join(spaceSimu$Y.data, lam1 = l1 * n^0.56, lam2 = 0, iter = iter)
fit.adj = abs(result2$ParCor) > 1e-6
sum(fit.adj == 1) / 2 # total number of edges detected
sum(fit.adj[true.adj == 1] == 1) / 2 # total number of true edges detected

### 3. Joint method with residue variance based weights
result3 = space.join(spaceSimu$Y.data, lam1 = l1 * n^0.86, lam2 = 0, weight = 1, iter = iter)
fit.adj = abs(result3$ParCor) > 1e-6
sum(fit.adj == 1) / 2 # total number of edges detected
sum(fit.adj[true.adj == 1] == 1) / 2 # total number of true edges detected

### 4. Joint method with degree based weights
result4 = space.join(spaceSimu$Y.data, lam1 = l1 * n^0.61, lam2 = 0, weight = 2, iter = iter)
fit.adj = abs(result4$ParCor) > 1e-6
sum(fit.adj == 1) / 2 # total number of edges detected
sum(fit.adj[true.adj == 1] == 1) / 2 # total number of true edges detected
spaceSimu is a simulated data example for inferring the partial correlation matrix

Description
A list containing a simulated data example for package space

Usage
data(spaceSimu)

Details

Y.data is a simulated array data with columns corresponding to samples and rows corresponding genes/clones. ParCor.true is the true partial correlation matrix used to simulate Y.data.

Value
spaceSimu is a list of two components:

- Y.data a numeric matrix consisting of 500 rows and 250 columns.
- ParCor.true a numeric matrix consisting of 500 rows and 500 columns.

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