Package 'finbipartite'

February 22, 2023

learn_bipartite_graph_nie

Laplacian matrix of a k-component bipartite graph via Nie's method Computes the Laplacian matrix of a bipartite graph on the basis of an observed similarity matrix.

Description

Laplacian matrix of a k-component bipartite graph via Nie's method

Computes the Laplacian matrix of a bipartite graph on the basis of an observed similarity matrix.

Usage

```
learn_bipartite_graph_nie(
   S,
   r,
   q,
   k,
   learning_rate = 1e-04,
   eta = 1,
   maxiter = 1000,
   reltol = 1e-06,
   verbose = TRUE,
   record_objective = FALSE
)
```

Arguments

S a p x p similarity matrix, where p is the number of nodes in the graph.

number of nodes in the objects set.
 number of nodes in the classes set.
 number of components of the graph.

learning_rate gradient descent parameter.

eta rank constraint hyperparameter.
maxiter maximum number of iterations.

reltol relative tolerance as a convergence criteria.

verbose whether or not to show a progress bar during the iterations.

record_objective

whether or not to record the objective function value during iterations.

A list containing possibly the following elements:

laplacian estimated Laplacian matrix
adjacency estimated adjacency matrix

B estimated graph weights matrix

maxiter number of iterations taken to reach convergence
convergence boolean flag to indicate whether or not the optimization converged

obj_fun objective function value per iteration

References

Feiping Nie, Xiaoqian Wang, Cheng Deng, Heng Huang. "Learning A Structured Optimal Bipartite Graph for Co-Clustering". Advances in Neural Information Processing Systems (NIPS 2017)

Examples

```
library(finbipartite)
library(igraph)
set.seed(42)
r <- 50
q <- 5
p \leftarrow r + q
bipartite <- sample_bipartite(r, q, type="Gnp", p = 1, directed=FALSE)
# randomly assign edge weights to connected nodes
E(bipartite)$weight <- 1
Lw <- as.matrix(laplacian_matrix(bipartite))</pre>
B \leftarrow -Lw[1:r, (r+1):p]
B[,] <- runif(length(B))</pre>
B \leftarrow B / rowSums(B)
# utils functions
from_B_to_laplacian <- function(B) {</pre>
  A <- from_B_to_adjacency(B)
  return(diag(rowSums(A)) - A)
}
from_B_to_adjacency <- function(B) {</pre>
  r <- nrow(B)
  q \leftarrow ncol(B)
  zeros_rxr <- matrix(0, r, r)</pre>
  zeros_qxq <- matrix(0, q, q)</pre>
  return(rbind(cbind(zeros_rxr, B), cbind(t(B), zeros_qxq)))
Ltrue <- from_B_to_laplacian(B)</pre>
X <- MASS::mvrnorm(100*p, rep(0, p), MASS::ginv(Ltrue))</pre>
S \leftarrow cov(X)
bipartite_graph <- learn_bipartite_graph_nie(S = S,</pre>
                                                   r = r,
                                                   q = q,
```

```
k = 1,
learning_rate = 5e-1,
eta = 0,
verbose=FALSE)
```

learn_connected_bipartite_graph_pgd

Laplacian matrix of a connected bipartite graph with Gaussian data Computes the Laplacian matrix of a bipartite graph on the basis of an observed data matrix.

Description

Laplacian matrix of a connected bipartite graph with Gaussian data

Computes the Laplacian matrix of a bipartite graph on the basis of an observed data matrix.

Usage

```
learn_connected_bipartite_graph_pgd(
   S,
   r,
   q,
   init = "naive",
   learning_rate = 1e-04,
   maxiter = 1000,
   reltol = 1e-05,
   verbose = TRUE,
   record_objective = FALSE,
   backtrack = TRUE
)
```

Arguments

S a p x p covariance matrix, where p is the number of nodes in the graph.

r number of nodes in the objects set.
q number of nodes in the classes set.

init string denoting how to compute the initial graph.

learning_rate gradient descent parameter.
maxiter maximum number of iterations.

reltol relative tolerance as a convergence criteria.

verbose whether or not to show a progress bar during the iterations.

record_objective

whether or not to record the objective function value during iterations.

backtrack whether or not to optimize the learning rate via backtracking.

A list containing possibly the following elements:

estimated Laplacian matrix laplacian estimated adjacency matrix adjacency В estimated graph weights matrix number of iterations taken to reach convergence maxiter convergence boolean flag to indicate whether or not the optimization converged learning rate value per iteration lr_seq obj_seq objective function value per iteration time taken per iteration until convergence is reached

Examples

elapsed_time

```
library(finbipartite)
library(igraph)
set.seed(42)
r <- 50
q <- 5
p \leftarrow r + q
bipartite <- sample_bipartite(r, q, type="Gnp", p = 1, directed=FALSE)</pre>
# randomly assign edge weights to connected nodes
E(bipartite)$weight <- 1
Lw <- as.matrix(laplacian_matrix(bipartite))</pre>
B \leftarrow -Lw[1:r, (r+1):p]
B[,] <- runif(length(B))</pre>
B \leftarrow B / rowSums(B)
# utils functions
from_B_to_laplacian <- function(B) {</pre>
  A <- from_B_to_adjacency(B)
  return(diag(rowSums(A)) - A)
}
from_B_to_adjacency <- function(B) {</pre>
  r <- nrow(B)
  q \leftarrow ncol(B)
  zeros_rxr <- matrix(0, r, r)</pre>
  zeros_qxq <- matrix(0, q, q)</pre>
  return(rbind(cbind(zeros_rxr, B), cbind(t(B), zeros_qxq)))
Ltrue <- from_B_to_laplacian(B)</pre>
X <- MASS::mvrnorm(100*p, rep(0, p), MASS::ginv(Ltrue))</pre>
bipartite_graph <- learn_connected_bipartite_graph_pgd(S = S,</pre>
                                                              q = q,
                                                              verbose=FALSE)
```

```
learn_heavy_tail_bipartite_graph_pgd
```

Laplacian matrix of a connected bipartite graph with heavy-tailed data Computes the Laplacian matrix of a bipartite graph on the basis of an observed data matrix whose distribution is assumed to be Student-t.

Description

Laplacian matrix of a connected bipartite graph with heavy-tailed data

Computes the Laplacian matrix of a bipartite graph on the basis of an observed data matrix whose distribution is assumed to be Student-t.

Usage

```
learn_heavy_tail_bipartite_graph_pgd(
    X,
    r,
    q,
    nu = 2.001,
    learning_rate = 1e-04,
    maxiter = 1000,
    reltol = 1e-05,
    init = "default",
    verbose = TRUE,
    record_objective = FALSE,
    backtrack = TRUE
)
```

Arguments

X a n x p data matrix, where p is the number of nodes in the graph and n is the

number of observations.

r number of nodes in the objects set.
q number of nodes in the classes set.

nu degrees of freedom of the Student-t distribution.

learning_rate gradient descent parameter.
maxiter maximum number of iterations.

reltol relative tolerance as a convergence criteria.

init string denoting how to compute the initial graph or a r x q matrix with initial

graph weights.

verbose whether or not to show a progress bar during the iterations.

record_objective

whether or not to record the objective function value during iterations.

backtrack whether or not to optimize the learning rate via backtracking.

A list containing possibly the following elements:

laplacian estimated Laplacian matrix
adjacency estimated adjacency matrix

B estimated graph weights matrix

maxiter number of iterations taken to reach convergence
convergence boolean flag to indicate whether or not the optimization converged
lr_seq learning rate value per iteration

elapsed_time time taken per iteration until convergence is reached

objective function value per iteration

Examples

obj_seq

```
library(finbipartite)
library(igraph)
set.seed(42)
r <- 50
q <- 5
p < -r + q
bipartite <- sample_bipartite(r, q, type="Gnp", p = 1, directed=FALSE)</pre>
# randomly assign edge weights to connected nodes
E(bipartite)$weight <- 1
Lw <- as.matrix(laplacian_matrix(bipartite))</pre>
B \leftarrow -Lw[1:r, (r+1):p]
B[,] <- runif(length(B))</pre>
B \leftarrow B / rowSums(B)
# utils functions
from_B_to_laplacian <- function(B) {</pre>
  A <- from_B_to_adjacency(B)
  return(diag(rowSums(A)) - A)
}
from_B_to_adjacency <- function(B) {</pre>
  r <- nrow(B)
  q \leftarrow ncol(B)
  zeros_rxr <- matrix(0, r, r)</pre>
  zeros_qxq <- matrix(0, q, q)</pre>
  return(rbind(cbind(zeros_rxr, B), cbind(t(B), zeros_qxq)))
Ltrue <- from_B_to_laplacian(B)</pre>
X <- MASS::mvrnorm(100*p, rep(0, p), MASS::ginv(Ltrue))</pre>
bipartite_graph <- learn_heavy_tail_bipartite_graph_pgd(X = X,</pre>
                                                              r = r,
                                                               q = q,
                                                               nu = 1e2,
                                                               verbose=FALSE)
```

```
learn_heavy_tail_kcomp_bipartite_graph
```

Laplacian matrix of a k-component bipartite graph with heavy-tailed data Computes the Laplacian matrix of a k-component bipartite graph on the basis of an observed data matrix whose distribution is assumed to be Student-t.

Description

Laplacian matrix of a k-component bipartite graph with heavy-tailed data

Computes the Laplacian matrix of a k-component bipartite graph on the basis of an observed data matrix whose distribution is assumed to be Student-t.

Usage

```
learn_heavy_tail_kcomp_bipartite_graph(
    X,
    r,
    q,
    k,
    nu = 2.001,
    rho = 1,
    learning_rate = 1e-04,
    maxiter = 1000,
    reltol = 1e-05,
    init = "default",
    verbose = TRUE,
    record_objective = FALSE
)
```

Arguments

X	a n x p data matrix, where p is the number of nodes in the graph and n is the number of observations.
r	number of nodes in the objects set.
q	number of nodes in the classes set.
k	number of components of the graph.
nu	degrees of freedom of the Student-t distribution.
rho	ADMM hyperparameter.
learning_rate	gradient descent parameter.
maxiter	maximum number of iterations.
reltol	relative tolerance as a convergence criteria.
init	string denoting how to compute the initial graph or a r x q matrix with initial graph weights.

```
verbose whether or not to show a progress bar during the iterations.

record_objective whether or not to record the objective function value during iterations.
```

A list containing possibly the following elements:

```
laplacian
                  estimated Laplacian matrix
adjacency
                  estimated adjacency matrix
                  estimated graph weights matrix
maxiter
                  number of iterations taken to reach convergence
                  boolean flag to indicate whether or not the optimization converged
convergence
dual_residual
                  dual residual value per iteration
primal_residual
                  primal residual value per iteration
aug_lag
                  augmented Lagrangian value per iteration
                  constraint relaxation hyperparameter value per iteration
rho_seq
elapsed_time
                  time taken per iteration until convergence is reached
```

Examples

```
library(finbipartite)
library(igraph)
set.seed(42)
r <- 50
q <- 5
p < -r + q
bipartite <- sample_bipartite(r, q, type="Gnp", p = 1, directed=FALSE)</pre>
# randomly assign edge weights to connected nodes
E(bipartite)$weight <- 1
Lw <- as.matrix(laplacian_matrix(bipartite))</pre>
B < - -Lw[1:r, (r+1):p]
B[,] <- runif(length(B))</pre>
B \leftarrow B / rowSums(B)
# utils functions
from_B_to_laplacian <- function(B) {</pre>
  A <- from_B_to_adjacency(B)
  return(diag(rowSums(A)) - A)
}
from_B_to_adjacency <- function(B) {</pre>
  r <- nrow(B)
  q <- ncol(B)
  zeros_rxr <- matrix(0, r, r)</pre>
  zeros_qxq <- matrix(0, q, q)</pre>
  return(rbind(cbind(zeros_rxr, B), cbind(t(B), zeros_qxq)))
}
```

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