Package 'missSBM'

May 8, 2026

Type Package

Title Handling Missing Data in Stochastic Block Models

Version 1.0.5

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Description
When a network is partially observed (here, NAs in the adjacency matrix rather than 1 or 0 due to missing information between node pairs), it is possible to account for the underlying process that generates those NAs. 'missSBM', presented in 'Barbillon, Chiquet and Tabouy' (2022) <doi:10.18637/jss.v101.i12>, adjusts the popular stochastic block model from network data sampled under various missing data conditions, as described in 'Tabouy, Barbillon and Chiquet' (2019) <doi:10.1080/01621459.2018.1562934>.

URL https://grosssbm.github.io/missSBM/

BugReports https://github.com/grossSBM/missSBM/issues

License GPL-3

Encoding UTF-8

LazyData true

RoxygenNote 7.3.2

Depends R (>= 3.4.0)

Imports Rcpp, methods, igraph, nloptr, ggplot2, future.apply, R6, rlang, sbm, magrittr, Matrix, RSpectra

LinkingTo Rcpp, RcppArmadillo, nloptr

Collate 'utils_missSBM.R' 'R6Class-networkSampling.R' 'R6Class-networkSampling_fit.R' 'R6Class-simpleSBM_fit.R' 'R6Class-missSBM_fit.R' 'R6Class-missSBM_collection.R' 'R6Class-networkSampler.R' 'R6Class-partlyObservedNetwork.R' 'RcppExports.R' 'er_network.R' 'estimateMissSBM.R' 'frenchblog2007.R' 'kmeans.R' 'missSBM-package.R' 'observeNetwork.R' 'war.R'

Suggests aricode, blockmodels, corrplot, future, testthat (>= 2.1.0), covr, knitr, rmarkdown, spelling

VignetteBuilder knitr

Language en-US

NeedsCompilation yes

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Repository CRAN

Date/Publication 2025-03-13 08:30:02 UTC

Contents

`blockDyadSampler` *Class for defining a block dyad sampler*

Description

Class for defining a block dyad sampler

Class for defining a block dyad sampler

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `missSBM::dyadSampler` -> `blockDyadSampler`

Active bindings

`df` the number of parameters of this sampling

Methods

Public methods:

- `blockDyadSampler$new()`
- `blockDyadSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
blockDyadSampler$new(
  parameters = NA,
  nbNodes = NA,
  directed = FALSE,
  clusters = NA
)
```

Arguments:

`parameters` the vector of parameters associated to the sampling at play

`nbNodes` number of nodes in the network

`directed` logical, directed network of not

`clusters` a vector of class memberships

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
blockDyadSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`blockDyadSampling_fit` *Class for fitting a block-dyad sampling*

Description

Class for fitting a block-dyad sampling

Class for fitting a block-dyad sampling

Super classes

`missSBM::networkSampling` -> `missSBM::networkSamplingDyads_fit` -> `blockDyadSampling_fit`

Active bindings

`vExpec` variational expectation of the sampling

`log_lambda` matrix, term for adjusting the imputation step which depends on the type of sampling

Methods

Public methods:

- `blockDyadSampling_fit$new()`
- `blockDyadSampling_fit$update_parameters()`
- `blockDyadSampling_fit$clone()`

Method `new()`**:** constructor

Usage:

```
blockDyadSampling_fit$new(partlyObservedNetwork, blockInit)
```

Arguments:

`partlyObservedNetwork` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`blockInit` n x Q matrix of initial block indicators

Method `update_parameters()`**:** a method to update the estimation of the parameters. By default, nothing to do (corresponds to MAR sampling)

Usage:

```
blockDyadSampling_fit$update_parameters(nu, Z)
```

Arguments:

`nu` the matrix of (uncorrected) imputation for missing entries
`Z` probabilities of block memberships

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
blockDyadSampling_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`blockNodeSampler` *Class for defining a block node sampler*

Description

Class for defining a block node sampler

Class for defining a block node sampler

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `missSBM::nodeSampler` -> `blockNodeSampler`

Methods

Public methods:

- `blockNodeSampler$new()`
- `blockNodeSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
blockNodeSampler$new(
  parameters = NA,
  nbNodes = NA,
  directed = FALSE,
  clusters = NA
)
```

Arguments:

`parameters` the vector of parameters associated to the sampling at play
`nbNodes` number of nodes in the network
`directed` logical, directed network of not
`clusters` a vector of class memberships

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
blockNodeSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`blockNodeSampling_fit` *Class for fitting a block-node sampling*

Description

Class for fitting a block-node sampling

Class for fitting a block-node sampling

Super classes

`missSBM::networkSampling` -> `missSBM::networkSamplingNodes_fit` -> `blockNodeSampling_fit`

Active bindings

`vExpec` variational expectation of the sampling

`log_lambda` double, term for adjusting the imputation step which depends on the type of sampling

Methods

Public methods:

- `blockNodeSampling_fit$new()`
- `blockNodeSampling_fit$update_parameters()`
- `blockNodeSampling_fit$clone()`

Method `new()`**:** constructor

Usage:

```
blockNodeSampling_fit$new(partlyObservedNetwork, blockInit)
```

Arguments:

`partlyObservedNetwork` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`blockInit` n x Q matrix of initial block indicators

Method `update_parameters()`**:** a method to update the estimation of the parameters. By default, nothing to do (corresponds to MAR sampling)

Usage:

```
blockNodeSampling_fit$update_parameters(imputedNet, Z)
```

Arguments:

`imputedNet` an adjacency matrix where missing values have been imputed

`Z` indicator of blocks

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
blockNodeSampling_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

coef.missSBM_fit *Extract model coefficients*

Description

Extracts model coefficients from objects missSBM_fit returned by estimateMissSBM()

Usage

```
## S3 method for class 'missSBM_fit'
coef(
  object,
  type = c("mixture", "connectivity", "covariates", "sampling"),
  ...
)
```

Arguments

object	an R6 object with class missSBM_fit
type	type of parameter that should be extracted. Either "mixture" (default), "connectivity", "covariates" or "sampling"
...	additional parameters for S3 compatibility. Not used

Value

A vector or matrix of coefficients extracted from the missSBM_fit model.

covarDyadSampling_fit *Class for fitting a dyad sampling with covariates*

Description

Class for fitting a dyad sampling with covariates

Class for fitting a dyad sampling with covariates

Super classes

missSBM::networkSampling -> missSBM::networkSamplingDyads_fit -> covarDyadSampling_fit

Active bindings

vExpec variational expectation of the sampling

Methods

Public methods:

- covarDyadSampling_fit$new()
- covarDyadSampling_fit$clone()

Method `new()`**:** constructor

Usage:

```
covarDyadSampling_fit$new(partialNet, ...)
```

Arguments:

`partialNet` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`...` used for compatibility

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
covarDyadSampling_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

---

covarNodeSampling_fit *Class for fitting a node-centered sampling with covariate*

Description

Class for fitting a node-centered sampling with covariate

Class for fitting a node-centered sampling with covariate

Super classes

`missSBM::networkSampling` -> `missSBM::networkSamplingNodes_fit` -> `covarNodeSampling_fit`

Active bindings

`vExpec` variational expectation of the sampling

Methods

Public methods:

- covarNodeSampling_fit$new()
- covarNodeSampling_fit$clone()

Method `new()`**:** constructor

Usage:

```
covarNodeSampling_fit$new(partlyObservedNetwork, ...)
```

Arguments:

`partlyObservedNetwork` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`...` used for compatibility

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
covarNodeSampling_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`degreeSampler` *Class for defining a degree sampler*

Description

Class for defining a degree sampler

Class for defining a degree sampler

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `missSBM::nodeSampler` -> `degreeSampler`

Methods

Public methods:

- `degreeSampler$new()`
- `degreeSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
degreeSampler$new(parameters = NA, degrees = NA, directed = FALSE)
```

Arguments:

`parameters` the vector of parameters associated to the sampling at play

`degrees` vector of nodes' degrees

`directed` logical, directed network of not

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
degreeSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`degreeSampling_fit` *Class for fitting a degree sampling*

Description

Class for fitting a degree sampling

Class for fitting a degree sampling

Super classes

`missSBM::networkSampling` -> `missSBM::networkSamplingNodes_fit` -> `degreeSampling_fit`

Active bindings

`vExpec` variational expectation of the sampling

Methods

Public methods:

- `degreeSampling_fit$new()`
- `degreeSampling_fit$update_parameters()`
- `degreeSampling_fit$update_imputation()`
- `degreeSampling_fit$clone()`

Method `new()`**:** constructor

Usage:

```
degreeSampling_fit$new(partlyObservedNetwork, blockInit, connectInit)
```

Arguments:

`partlyObservedNetwork` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`blockInit` n x Q matrix of initial block indicators

`connectInit` Q x Q matrix of initial block probabilities of connection

Method `update_parameters()`**:** a method to update the estimation of the parameters. By default, nothing to do (corresponds to MAR sampling)

Usage:

```
degreeSampling_fit$update_parameters(imputedNet, ...)
```

Arguments:

`imputedNet` an adjacency matrix where missing values have been imputed

`...` used for compatibility

Method `update_imputation()`**:** a method to update the imputation of the missing entries.

Usage:

```
degreeSampling_fit$update_imputation(PI, ...)
```

Arguments:

`PI` the matrix of inter/intra class probability of connection

`...` use for compatibility

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
degreeSampling_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`doubleStandardSampler` *Class for defining a double-standard sampler*

Description

Class for defining a double-standard sampler

Class for defining a double-standard sampler

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `missSBM::dyadSampler` -> `doubleStandardSampler`

Methods

Public methods:

- `doubleStandardSampler$new()`
- `doubleStandardSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
doubleStandardSampler$new(parameters = NA, adjMatrix = NA, directed = FALSE)
```

Arguments:

`parameters` the vector of parameters associated to the sampling at play

`adjMatrix` matrix of adjacency

`directed` logical, directed network of not

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
doubleStandardSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

doubleStandardSampling_fit

Class for fitting a double-standard sampling

Description

Class for fitting a double-standard sampling

Class for fitting a double-standard sampling

Super classes

`missSBM::networkSampling` -> `missSBM::networkSamplingDyads_fit` -> `doubleStandardSampling_fit`

Active bindings

`vExpec` variational expectation of the sampling

Methods

Public methods:

- `doubleStandardSampling_fit$new()`
- `doubleStandardSampling_fit$update_parameters()`
- `doubleStandardSampling_fit$update_imputation()`
- `doubleStandardSampling_fit$clone()`

Method `new()`**:** constructor

Usage:

```
doubleStandardSampling_fit$new(partlyObservedNetwork, ...)
```

Arguments:

`partlyObservedNetwork` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`...` used for compatibility

Method `update_parameters()`**:** a method to update the estimation of the parameters. By default, nothing to do (corresponds to MAR sampling)

Usage:

```
doubleStandardSampling_fit$update_parameters(nu, ...)
```

Arguments:

`nu` an adjacency matrix with imputed values (only)

`...` use for compatibility

Method `update_imputation()`**:** a method to update the imputation of the missing entries.

Usage:

```
doubleStandardSampling_fit$update_imputation(nu)
```

Arguments:

`nu` the matrix of (uncorrected) imputation for missing entries

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
doubleStandardSampling_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`dyadSampler` *Virtual class for all dyad-centered samplers*

Description

Virtual class for all dyad-centered samplers

Virtual class for all dyad-centered samplers

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `dyadSampler`

Methods

Public methods:

- `dyadSampler$new()`
- `dyadSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
dyadSampler$new(type = NA, parameters = NA, nbNodes = NA, directed = FALSE)
```

Arguments:

`type` character for the type of sampling. must be in ("dyad", "covar-dyad", "node", "covar-node", "block-node", "block-dyad", "double-standard", "degree")

`parameters` the vector of parameters associated to the sampling at play

`nbNodes` number of nodes in the network

`directed` logical, directed network of not

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
dyadSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

dyadSampling_fit *Class for fitting a dyad sampling*

Description

Class for fitting a dyad sampling

Class for fitting a dyad sampling

Super classes

missSBM::networkSampling -> missSBM::networkSamplingDyads_fit -> dyadSampling_fit

Active bindings

vExpec variational expectation of the sampling

Methods

Public methods:

- dyadSampling_fit$new()
- dyadSampling_fit$clone()

Method new(): constructor

Usage:

```
dyadSampling_fit$new(partlyObservedNetwork, ...)
```

Arguments:

partlyObservedNetwork a object with class partlyObservedNetwork representing the observed data with possibly missing entries

... used for compatibility

Method clone(): The objects of this class are cloneable with this method.

Usage:

```
dyadSampling_fit$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

er_network *ER ego centered network*

Description

A dataset containing the weighted PPI network centered around the ESR1 (ER) protein

Usage

```
er_network
```

Format

A sparse symmetric matrix with 741 rows and 741 columns ESR1

Source

https://string-db.org/

Examples

```
data("er_network")
class(er_network)
```

estimateMissSBM *Estimation of simple SBMs with missing data*

Description

Variational EM inference of Stochastic Block Models indexed by block number from a partially observed network.

Usage

```
estimateMissSBM(
  adjacencyMatrix,
  vBlocks,
  sampling,
  covariates = list(),
  control = list()
)
```

Arguments

`adjacencyMatrix`	The N x N adjacency matrix of the network data. If `adjacencyMatrix` is symmetric, we assume an undirected network with no loop; otherwise the network is assumed to be directed.
`vBlocks`	The vector of number of blocks considered in the collection.
`sampling`	The model used to described the process that originates the missing data: MAR designs ("dyad", "node","covar-dyad","covar-node","snowball") and MNAR designs ("double-standard", "block-dyad", "block-node" , "degree") are available. See details.
`covariates`	An optional list with M entries (the M covariates). If the covariates are node-centered, each entry of `covariates` must be a size-N vector; if the covariates are dyad-centered, each entry of `covariates` must be N x N matrix.
`control`	a list of parameters controlling advanced features. See details.

Details

Internal functions use `future_lapply`, so set your plan to `'multisession'` or `'multicore'` to use several cores/workers. The list of parameters `control` tunes more advanced features, such as the initialization, how covariates are handled in the model, and the variational EM algorithm:

- useCov logical. If `covariates` is not null, should they be used for the for the SBM inference (or just for the sampling)? Default is TRUE.
- clusterInit Initial method for clustering: either a character ("spectral") or a list with `length(vBlocks)` vectors, each with size `ncol(adjacencyMatrix)`, providing a user-defined clustering. Default is "spectral". similarity An R x R -> R function to compute similarities between node covariates. Default is `l1_similarity`, that is, -abs(x-y). Only relevant when the covariates are node-centered (i.e. `covariates` is a list of size-N vectors).
- threshold V-EM algorithm stops stop when an optimization step changes the objective function or the parameters by less than threshold. Default is 1e-2.
- maxIter V-EM algorithm stops when the number of iteration exceeds maxIter. Default is 50.
- fixPointIter number of fix-point iterations in the V-E step. Default is 3.
- exploration character indicating the kind of exploration used among "forward", "backward", "both" or "none". Default is "both".
- iterates integer for the number of iterations during exploration. Only relevant when `exploration` is different from "none". Default is 1.
- trace logical for verbosity. Default is TRUE.

The different sampling designs are split into two families in which we find dyad-centered and node-centered samplings. See doi:10.1080/01621459.2018.1562934 for a complete description.

- Missing at Random (MAR)
  - dyad parameter = p = Prob(Dyad(i,j) is observed)
  - node parameter = p = Prob(Node i is observed)
  - covar-dyad": parameter = beta in R^M, such that Prob(Dyad (i,j) is observed) = logistic(parameter' covarArray (i,j, .))

  - covar-node": parameter = nu in R^M such that Prob(Node i is observed) = logistic(parameter' covarMatrix (i,)
  - snowball": parameter = number of waves with Prob(Node i is observed in the 1st wave)
- Missing Not At Random (MNAR)
  - double-standard parameter = (p0,p1) with p0 = Prob(Dyad (i,j) is observed | the dyad is equal to 0), p1 = Prob(Dyad (i,j) is observed | the dyad is equal to 1)
  - block-node parameter = c(p(1),...,p(Q)) and p(q) = Prob(Node i is observed | node i is in cluster q)
  - block-dyad parameter = c(p(1,1),...,p(Q,Q)) and p(q,l) = Prob(Edge (i,j) is observed | node i is in cluster q and node j is in cluster l)

Value

Returns an R6 object with class `missSBM_collection`.

See Also

`observeNetwork`, `missSBM_collection` and `missSBM_fit`.

Examples

```
## SBM parameters
N <- 100 # number of nodes
Q <- 3   # number of clusters
pi <- rep(1,Q)/Q     # block proportion
theta <- list(mean = diag(.45,Q) + .05 ) # connectivity matrix

## Sampling parameters
samplingParameters <- .75 # the sampling rate
sampling  <- "dyad"      # the sampling design

## generate a undirected binary SBM with no covariate
sbm <- sbm::sampleSimpleSBM(N, pi, theta)

## Uncomment to set parallel computing with future
## future::plan("multicore", workers = 2)

## Sample some dyads data + Infer SBM with missing data
collection <-
   observeNetwork(sbm$networkData, sampling, samplingParameters) %>%
   estimateMissSBM(vBlocks = 1:4, sampling = sampling)
plot(collection, "monitoring")
plot(collection, "icl")

collection$ICL
coef(collection$bestModel$fittedSBM, "connectivity")

myModel <- collection$bestModel
plot(myModel, "expected")
plot(myModel, "imputed")
plot(myModel, "meso")
```

```
coef(myModel, "sampling")
coef(myModel, "connectivity")
predict(myModel)[1:5, 1:5]
```

`fitted.missSBM_fit` *Extract model fitted values from object* `missSBM_fit`*, return by* `estimateMissSBM()`

Description

Extract model fitted values from object `missSBM_fit`, return by `estimateMissSBM()`

Usage

```
## S3 method for class 'missSBM_fit'
fitted(object, ...)
```

Arguments

`object`	an R6 object with class `missSBM_fit`
`...`	additional parameters for S3 compatibility.

Value

A matrix of estimated probabilities of connection

`frenchblog2007` *Political Blogosphere network prior to 2007 French presidential election*

Description

French Political Blogosphere network dataset consists of a single day snapshot of over 200 political blogs automatically extracted the 14 October 2006 and manually classified by the "Observatoire Présidentielle" project. Originally part of the 'mixer' package

Usage

```
frenchblog2007
```

Format

An igraph object with 196 nodes. The vertex attribute "party" provides a possible clustering of the nodes.

Source

https://www.meltwater.com/en/suite/consumer-intelligence?utm_source=direct&utm_medium=linkfluence

Examples

```
data(frenchblog2007)
igraph::V(frenchblog2007)$party
igraph::plot.igraph(frenchblog2007,
  vertex.color = factor(igraph::V(frenchblog2007)$party),
  vertex.label = NA
 )
```

`l1_similarity` *L1-similarity*

Description

Compute l1-similarity between two vectors

Usage

```
l1_similarity(x, y)
```

Arguments

x	a vector
y	a vector

Value

a vector equal to -abs(x-y)

`missSBM_collection` *An R6 class to represent a collection of SBM fits with missing data*

Description

The function `estimateMissSBM()` fits a collection of SBM with missing data for a varying number of block. These models with class `missSBM_fit` are stored in an instance of an object with class `missSBM_collection`, described here.

Fields are accessed via active binding and cannot be changed by the user.

This class comes with a set of R6 methods, some of them being useful for the user and exported as S3 methods. See the documentation for `show()` and `print()`

Active bindings

`models` a list of models

`ICL` the vector of Integrated Classification Criterion (ICL) associated to the models in the collection (the smaller, the better)

`bestModel` the best model according to the ICL

`vBlocks` a vector with the number of blocks

`optimizationStatus` a data.frame summarizing the optimization process for all models

Methods

Public methods:

- `missSBM_collection$new()`
- `missSBM_collection$estimate()`
- `missSBM_collection$explore()`
- `missSBM_collection$plot()`
- `missSBM_collection$show()`
- `missSBM_collection$print()`
- `missSBM_collection$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
missSBM_collection$new(partlyObservedNet, sampling, clusterInit, control)
```

Arguments:

`partlyObservedNet` An object with class `partlyObservedNetwork`.

`sampling` The sampling design for the modelling of missing data: MAR designs ("dyad", "node") and MNAR designs ("double-standard", "block-dyad", "block-node" ,"degree")

`clusterInit` Initial clustering: a list of vectors, each with size `ncol(adjacencyMatrix)`.

`control` a list of parameters controlling advanced features. Only 'trace' and 'useCov' are relevant here. See `estimateMissSBM()` for details.

Method `estimate()`**:** method to launch the estimation of the collection of models

Usage:

```
missSBM_collection$estimate(control)
```

Arguments:

`control` a list of parameters controlling the variational EM algorithm. See details of function `estimateMissSBM()`

Method `explore()`**:** method for performing exploration of the ICL

Usage:

```
missSBM_collection$explore(control)
```

Arguments:

`control` a list of parameters controlling the exploration, similar to those found in the regular function `estimateMissSBM()`

Method `plot()`**:** plot method for missSBM_collection

Usage:

```
missSBM_collection$plot(type = c("icl", "elbo", "monitoring"))
```

Arguments:

`type` the type specifies the field to plot, either "icl", "elbo" or "monitoring". Default is "icl"

Method `show()`**:** show method for missSBM_collection

Usage:

```
missSBM_collection$show()
```

Method `print()`**:** User friendly print method

Usage:

```
missSBM_collection$print()
```

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
missSBM_collection$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

Examples

```
## Uncomment to set parallel computing with future
## future::plan("multicore", workers = 2)

## Sample 75% of dyads in  French political Blogosphere's network data
adjacencyMatrix <- missSBM::frenchblog2007 %>%
  igraph::delete.vertices(1:100) %>%
  igraph::as_adj () %>%
  missSBM::observeNetwork(sampling = "dyad", parameters = 0.75)
collection <- estimateMissSBM(adjacencyMatrix, 1:5, sampling = "dyad")
class(collection)
```

`missSBM_fit` *An R6 class to represent an SBM fit with missing data*

Description

The function `estimateMissSBM()` fits a collection of SBM for varying number of block. Each fitted SBM is an instance of an R6 object with class `missSBM_fit`, described here.

Fields are accessed via active binding and cannot be changed by the user.

This class comes with a set of R6 methods, some of them being useful for the user and exported as S3 methods. See the documentation for `show()`, `print()`, `fitted()`, `predict()`, `plot()`.

Active bindings

`fittedSBM` the fitted SBM with class `SimpleSBM_fit_noCov`, `SimpleSBM_fit_withCov` or `SimpleSBM_fit_MNAR` inheriting from class `sbm::SimpleSBM_fit`

`fittedSampling` the fitted sampling, inheriting from class `networkSampling` and corresponding fits

`imputedNetwork` The network data as a matrix with NAs values imputed with the current model

`monitoring` a list carrying information about the optimization process

`entropyImputed` the entropy of the distribution of the imputed dyads

`entropy` the entropy due to the distribution of the imputed dyads and of the clustering

`vExpec` double: variational expectation of the complete log-likelihood

`penalty` double, value of the penalty term in ICL

`loglik` double: approximation of the log-likelihood (variational lower bound) reached

`ICL` double: value of the integrated classification log-likelihood

Methods

Public methods:

- `missSBM_fit$new()`
- `missSBM_fit$doVEM()`
- `missSBM_fit$show()`
- `missSBM_fit$print()`
- `missSBM_fit$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
missSBM_fit$new(partlyObservedNet, netSampling, clusterInit, useCov = TRUE)
```

Arguments:

`partlyObservedNet` An object with class `partlyObservedNetwork`.

`netSampling` The sampling design for the modelling of missing data: MAR designs ("dyad", "node") and MNAR designs ("double-standard", "block-dyad", "block-node" ,"degree")

`clusterInit` Initial clustering: a vector with size `ncol(adjacencyMatrix)`, providing a user-defined clustering. The number of blocks is deduced from the number of levels in with `clusterInit`.

`useCov` logical. If covariates are present in partlyObservedNet, should they be used for the inference or of the network sampling design, or just for the SBM inference? default is TRUE.

Method `doVEM()`**:** a method to perform inference of the current missSBM fit with variational EM

Usage:

```
missSBM_fit$doVEM(
  control = list(threshold = 0.01, maxIter = 100, fixPointIter = 3, trace = TRUE)
)
```

Arguments:

`control` a list of parameters controlling the variational EM algorithm. See details of function `estimateMissSBM()`

Method `show()`**:** show method for missSBM_fit

Usage:

```
missSBM_fit$show()
```

Method `print()`**:** User friendly print method

Usage:

```
missSBM_fit$print()
```

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
missSBM_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

Examples

```
## Sample 75% of dyads in  French political Blogosphere's network data
adjMatrix <- missSBM::frenchblog2007 %>%
  igraph::as_adj (sparse = FALSE) %>%
  missSBM::observeNetwork(sampling = "dyad", parameters = 0.75)
collection <- estimateMissSBM(adjMatrix, 3:5, sampling = "dyad")
my_missSBM_fit <- collection$bestModel
class(my_missSBM_fit)
plot(my_missSBM_fit, "imputed")
```

`networkSampler` *Definition of R6 Class 'networkSampling_sampler'*

Description

Definition of R6 Class 'networkSampling_sampler'

Definition of R6 Class 'networkSampling_sampler'

Details

This class is use to define a sampling model for a network. Inherits from 'networkSampling'. Owns a rSampling method which takes an adjacency matrix as an input and send back an object with class partlyObservedNetwork.

Super class

`missSBM::networkSampling` -> `networkSampler`

Active bindings

`samplingMatrix` a matrix of logical indicating observed entries

Methods

Public methods:

- `networkSampler$new()`
- `networkSampler$rSamplingMatrix()`
- `networkSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
networkSampler$new(type = NA, parameters = NA, nbNodes = NA, directed = FALSE)
```

Arguments:

`type` character for the type of sampling. must be in ("dyad", "covar-dyad", "node", "covar-node", "block-node", "block-dyad", "double-standard", "degree")

`parameters` the vector of parameters associated to the sampling at play

`nbNodes` number of nodes in the network

`directed` logical, directed network of not

Method `rSamplingMatrix()`**:** a method for drawing a sampling matrix according to the current sampling design

Usage:

```
networkSampler$rSamplingMatrix()
```

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
networkSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

See Also

`partlyObservedNetwork`

networkSampling *Definition of R6 Class 'networkSampling'*

Description

Definition of R6 Class 'networkSampling'

Definition of R6 Class 'networkSampling'

Details

this virtual class is the mother of all subtypes of networkSampling (either sampler or fit) It is used to define a sampling model for a network. It has a rSampling method which takes an adjacency matrix as an input and send back an object with class partlyObservedNetwork.

Active bindings

`type` a character for the type of sampling

`parameters` the vector of parameters associated with the sampling at play

`df` the number of entries in the vector of parameters

Methods

Public methods:

- `networkSampling$new()`
- `networkSampling$show()`
- `networkSampling$print()`
- `networkSampling$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
networkSampling$new(type = NA, parameters = NA)
```

Arguments:

`type` character for the type of sampling. must be in ("dyad", "covar-dyad", "node", "covar-node", "block-node", "block-dyad", "double-standard", "degree")

`parameters` the vector of parameters associated to the sampling at play

Method `show()`**:** show method

Usage:

```
networkSampling$show(
  type = paste0(private$name, "-model for network sampling\n")
)
```

Arguments:

`type` character used to specify the type of sampling

Method `print()`**:** User friendly print method

Usage:

```
networkSampling$print()
```

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
networkSampling$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`networkSamplingDyads_fit` *Virtual class used to define a family of networkSamplingDyads_fit*

Description

Virtual class used to define a family of networkSamplingDyads_fit

Virtual class used to define a family of networkSamplingDyads_fit

Super class

`missSBM::networkSampling` -> `networkSamplingDyads_fit`

Active bindings

`penalty` double, value of the penalty term in ICL

`log_lambda` double, term for adjusting the imputation step which depends on the type of sampling

Methods

Public methods:

- `networkSamplingDyads_fit$new()`
- `networkSamplingDyads_fit$show()`
- `networkSamplingDyads_fit$update_parameters()`
- `networkSamplingDyads_fit$update_imputation()`
- `networkSamplingDyads_fit$clone()`

Method `new()`**:** constructor for networkSampling_fit

Usage:

```
networkSamplingDyads_fit$new(partlyObservedNetwork, name)
```

Arguments:

`partlyObservedNetwork` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`name` a character for the name of sampling to fit on the partlyObservedNetwork

Method `show()`: show method

Usage:

```
networkSamplingDyads_fit$show()
```

Method `update_parameters()`: a method to update the estimation of the parameters. By default, nothing to do (corresponds to MAR sampling)

Usage:

```
networkSamplingDyads_fit$update_parameters(...)
```

Arguments:

`...` use for compatibility

Method `update_imputation()`: a method to update the imputation of the missing entries.

Usage:

```
networkSamplingDyads_fit$update_imputation(nu)
```

Arguments:

`nu` the matrix of (uncorrected) imputation for missing entries

Method `clone()`: The objects of this class are cloneable with this method.

Usage:

```
networkSamplingDyads_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`networkSamplingNodes_fit` *Virtual class used to define a family of networkSamplingNodes_fit*

Description

Virtual class used to define a family of networkSamplingNodes_fit

Virtual class used to define a family of networkSamplingNodes_fit

Super class

`missSBM::networkSampling` -> `networkSamplingNodes_fit`

Active bindings

`penalty` double, value of the penalty term in ICL

`log_lambda` double, term for adjusting the imputation step which depends on the type of sampling

Methods

Public methods:

- networkSamplingNodes_fit$new()
- networkSamplingNodes_fit$show()
- networkSamplingNodes_fit$update_parameters()
- networkSamplingNodes_fit$update_imputation()
- networkSamplingNodes_fit$clone()

Method `new()`**:** constructor

Usage:

```
networkSamplingNodes_fit$new(partlyObservedNetwork, name)
```

Arguments:

`partlyObservedNetwork` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`name` a character for the name of sampling to fit on the partlyObservedNetwork

Method `show()`**:** show method

Usage:

```
networkSamplingNodes_fit$show()
```

Method `update_parameters()`**:** a method to update the estimation of the parameters. By default, nothing to do (corresponds to MAR sampling)

Usage:

```
networkSamplingNodes_fit$update_parameters(...)
```

Arguments:

`...` use for compatibility

Method `update_imputation()`**:** a method to update the imputation of the missing entries.

Usage:

```
networkSamplingNodes_fit$update_imputation(nu)
```

Arguments:

`nu` the matrix of (uncorrected) imputation for missing entries

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
networkSamplingNodes_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

nodeSampler *Virtual class for all node-centered samplers*

Description

Virtual class for all node-centered samplers

Virtual class for all node-centered samplers

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `nodeSampler`

Methods

Public methods:

- `nodeSampler$clone()`

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
nodeSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

nodeSampling_fit *Class for fitting a node sampling*

Description

Class for fitting a node sampling

Class for fitting a node sampling

Super classes

`missSBM::networkSampling` -> `missSBM::networkSamplingNodes_fit` -> `nodeSampling_fit`

Active bindings

`vExpec` variational expectation of the sampling

Methods

Public methods:

- nodeSampling_fit$new()
- nodeSampling_fit$clone()

Method `new()`**:** constructor

Usage:

```
nodeSampling_fit$new(partlyObservedNetwork, ...)
```

Arguments:

`partlyObservedNetwork` a object with class partlyObservedNetwork representing the observed data with possibly missing entries

`...` used for compatibility

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
nodeSampling_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`observeNetwork` *Observe a network partially according to a given sampling design*

Description

This function draws observations in an adjacency matrix according to a given network sampling design.

Usage

```
observeNetwork(
  adjacencyMatrix,
  sampling,
  parameters,
  clusters = NULL,
  covariates = list(),
  similarity = l1_similarity,
  intercept = 0
)
```

Arguments

adjacencyMatrix	The N x N adjacency matrix of the network to sample.
sampling	The sampling design used to observe the adjacency matrix, see details.
parameters	The sampling parameters (adapted to each sampling, see details).
clusters	An optional clustering membership vector of the nodes. Only necessary for block samplings.
covariates	An optional list with M entries (the M covariates). If the covariates are node-centered, each entry of covariates. must be a size-N vector; if the covariates are dyad-centered, each entry of covariates must be N x N matrix.
similarity	An optional function to compute similarities between node covariates. Default is l1_similarity, that is, -abs(x-y). Only relevant when the covariates are node-centered.
intercept	An optional intercept term to be added in case of the presence of covariates. Default is 0.

Details

Internal functions use future_lapply, so set your plan to 'multisession' or 'multicore' to use several cores/workers. The list of parameters control tunes more advanced features, such as the initialization, how covariates are handled in the model, and the variational EM algorithm:

- useCov logical. If covariates is not null, should they be used for the for the SBM inference (or just for the sampling)? Default is TRUE.
- clusterInit Initial method for clustering: either a character ("spectral") or a list with length(vBlocks) vectors, each with size ncol(adjacencyMatrix), providing a user-defined clustering. Default is "spectral". similarity An R x R -> R function to compute similarities between node covariates. Default is l1_similarity, that is, -abs(x-y). Only relevant when the covariates are node-centered (i.e. covariates is a list of size-N vectors).
- threshold V-EM algorithm stops stop when an optimization step changes the objective function or the parameters by less than threshold. Default is 1e-2.
- maxIter V-EM algorithm stops when the number of iteration exceeds maxIter. Default is 50.
- fixPointIter number of fix-point iterations in the V-E step. Default is 3.
- exploration character indicating the kind of exploration used among "forward", "backward", "both" or "none". Default is "both".
- iterates integer for the number of iterations during exploration. Only relevant when exploration is different from "none". Default is 1.
- trace logical for verbosity. Default is TRUE.

The different sampling designs are split into two families in which we find dyad-centered and node-centered samplings. See doi:10.1080/01621459.2018.1562934 for a complete description.

- Missing at Random (MAR)
  - dyad parameter = p = Prob(Dyad(i,j) is observed)
  - node parameter = p = Prob(Node i is observed)

  - covar-dyad": parameter = beta in R^M, such that Prob(Dyad (i,j) is observed) = logistic(parameter' covarArray (i,j, .))
  - covar-node": parameter = nu in R^M such that Prob(Node i is observed) = logistic(parameter' covarMatrix (i,)
  - snowball": parameter = number of waves with Prob(Node i is observed in the 1st wave)

- Missing Not At Random (MNAR)
  - double-standard parameter = (p0,p1) with p0 = Prob(Dyad (i,j) is observed | the dyad is equal to 0), p1 = Prob(Dyad (i,j) is observed | the dyad is equal to 1)
  - block-node parameter = c(p(1),...,p(Q)) and p(q) = Prob(Node i is observed | node i is in cluster q)
  - block-dyad parameter = c(p(1,1),...,p(Q,Q)) and p(q,l) = Prob(Edge (i,j) is observed | node i is in cluster q and node j is in cluster l)

Value

an adjacency matrix with the same dimension as the input, yet with additional NAs.

Examples

```
## SBM parameters
N <- 300 # number of nodes
Q <- 3   # number of clusters
pi <- rep(1,Q)/Q     # block proportion
theta <- list(mean = diag(.45,Q) + .05 ) # connectivity matrix

## simulate an unidrected binary SBM without covariate
sbm <- sbm::sampleSimpleSBM(N, pi, theta)

## Sample network data

# some sampling design and their associated parameters
sampling_parameters <- list(
   "dyad" = .3,
   "node" = .3,
   "double-standard" = c(0.4, 0.8),
   "block-node" = c(.3, .8, .5),
   "block-dyad" = theta$mean,
   "degree" = c(.01, .01),
   "snowball" = c(2,.1)
 )

observed_networks <- list()

for (sampling in names(sampling_parameters)) {
  observed_networks[[sampling]] <-
     missSBM::observeNetwork(
       adjacencyMatrix = sbm$networkData,
       sampling        = sampling,
       parameters      = sampling_parameters[[sampling]],
       cluster         = sbm$memberships
```

```
    )
}
```

`partlyObservedNetwork` *An R6 Class used for internal representation of a partially observed network*

Description

An R6 Class used for internal representation of a partially observed network

An R6 Class used for internal representation of a partially observed network

Details

This class is not exported to the user

Active bindings

`samplingRate` The percentage of observed dyads

`nbNodes` The number of nodes

`nbDyads` The number of dyads

`is_directed` logical indicating if the network is directed or not

`networkData` The adjacency matrix of the network

`covarArray` the array of covariates

`covarMatrix` the matrix of covariates

`samplingMatrix` matrix of observed and non-observed edges

`samplingMatrixBar` matrix of observed and non-observed edges

`observedNodes` a vector of observed and non-observed nodes (observed means at least one non NA value)

Methods

Public methods:

- `partlyObservedNetwork$new()`
- `partlyObservedNetwork$clustering()`
- `partlyObservedNetwork$imputation()`
- `partlyObservedNetwork$clone()`

Method `new()`**:** constructor

Usage:

```
partlyObservedNetwork$new(
  adjacencyMatrix,
  covariates = list(),
  similarity = l1_similarity
)
```

Arguments:

`adjacencyMatrix` The adjacency matrix of the network

`covariates` A list with M entries (the M covariates), each of whom being either a size-N vector or N x N matrix.

`similarity` An R x R -> R function to compute similarities between node covariates. Default is `l1_similarity`, that is, -abs(x-y).

Method `clustering()`**:** method to cluster network data with missing value

Usage:

```
partlyObservedNetwork$clustering(
  vBlocks,
  imputation = ifelse(is.null(private$phi), "median", "average")
)
```

Arguments:

`vBlocks` The vector of number of blocks considered in the collection.

`imputation` character indicating the type of imputation among "median", "average"

Method `imputation()`**:** basic imputation from existing clustering

Usage:

```
partlyObservedNetwork$imputation(type = c("median", "average", "zero"))
```

Arguments:

`type` a character, the type of imputation. Either "median" or "average"

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
partlyObservedNetwork$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`plot.missSBM_fit` *Visualization for an object* `missSBM_fit`

Description

Plot function for the various fields of a `missSBM_fit`: the fitted SBM (network or connectivity), and a plot monitoring the optimization.

Usage

```
## S3 method for class 'missSBM_fit'
plot(
  x,
  type = c("imputed", "expected", "meso", "monitoring"),
  dimLabels = list(row = "node", col = "node"),
  ...
)
```

Arguments

x	an object with class missSBM_fit
type	the type specifies the field to plot, either "imputed", "expected", "meso", or "monitoring"
dimLabels	: a list of two characters specifying the labels of the nodes. Default to list(row= 'node',col = 'node'))
...	additional parameters for S3 compatibility. Not used

Value

a ggplot object

---

predicted.missSBM_fit *Prediction of a missSBM_fit (i.e. network with imputed missing dyads)*

Description

Prediction of a missSBM_fit (i.e. network with imputed missing dyads)

Usage

```
## S3 method for class 'missSBM_fit'
predict(object, ...)
```

Arguments

object	an R6 object with class missSBM_fit
...	additional parameters for S3 compatibility.

Value

an adjacency matrix between pairs of nodes. Missing dyads are imputed with their expected values, i.e. by there estimated probabilities of connection under the missing SBM.

`simpleDyadSampler` *Class for defining a simple dyad sampler*

Description

Class for defining a simple dyad sampler

Class for defining a simple dyad sampler

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `missSBM::dyadSampler` -> `simpleDyadSampler`

Methods

Public methods:

- `simpleDyadSampler$new()`
- `simpleDyadSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
simpleDyadSampler$new(
  parameters = NA,
  nbNodes = NA,
  directed = FALSE,
  covarArray = NULL,
  intercept = 0
)
```

Arguments:

`parameters` the vector of parameters associated to the sampling at play

`nbNodes` number of nodes in the network

`directed` logical, directed network of not

`covarArray` an array of covariates used

`intercept` double, intercept term used to compute the probability of sampling in the presence of covariates. Default 0.

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
simpleDyadSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`simpleNodeSampler` *Class for defining a simple node sampler*

Description

Class for defining a simple node sampler

Class for defining a simple node sampler

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `missSBM::nodeSampler` -> `simpleNodeSampler`

Methods

Public methods:

- `simpleNodeSampler$new()`
- `simpleNodeSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
simpleNodeSampler$new(
  parameters = NA,
  nbNodes = NA,
  directed = FALSE,
  covarMatrix = NULL,
  intercept = 0
)
```

Arguments:

`parameters` the vector of parameters associated to the sampling at play

`nbNodes` number of nodes in the network

`directed` logical, directed network of not

`covarMatrix` a matrix of covariates used

`intercept` double, intercept term used to compute the probability of sampling in the presence of covariates. Default 0.

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
simpleNodeSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

SimpleSBM_fit *This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.*

Description

This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.

This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.

Details

It is not designed not be call by the user

Super classes

sbm::SBM -> sbm::SimpleSBM -> SimpleSBM_fit

Active bindings

`type` the type of SBM (distribution of edges values, network type, presence of covariates)

`penalty` double, value of the penalty term in ICL

`entropy` double, value of the entropy due to the clustering distribution

`loglik` double: approximation of the log-likelihood (variational lower bound) reached

`ICL` double: value of the integrated classification log-likelihood

Methods

Public methods:

- `SimpleSBM_fit$new()`
- `SimpleSBM_fit$doVEM()`
- `SimpleSBM_fit$reorder()`
- `SimpleSBM_fit$clone()`

Method `new()`**:** constructor for simpleSBM_fit for missSBM purpose

Usage:

```
SimpleSBM_fit$new(networkData, clusterInit, covarList = list())
```

Arguments:

`networkData` a structure to store network under missing data condition: either a matrix possibly with NA, or a missSBM:::partlyObservedNetwork

`clusterInit` Initial clustering: a vector with size `ncol(adjacencyMatrix)`, providing a user-defined clustering with `nbBlocks` levels.

`covarList` An optional list with M entries (the M covariates).

Method `doVEM()`**:** method to perform estimation via variational EM

Usage:

```
SimpleSBM_fit$doVEM(
  threshold = 0.01,
  maxIter = 100,
  fixPointIter = 3,
  trace = FALSE
)
```

Arguments:

`threshold` stop when an optimization step changes the objective function by less than threshold. Default is 1e-4.

`maxIter` V-EM algorithm stops when the number of iteration exceeds maxIter. Default is 10

`fixPointIter` number of fix-point iterations in the Variational E step. Default is 5.

`trace` logical for verbosity. Default is `FALSE`.

Method `reorder()`**:** permute group labels by order of decreasing probability

Usage:

```
SimpleSBM_fit$reorder()
```

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
SimpleSBM_fit$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`SimpleSBM_fit_MNAR` *This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.*

Description

This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.

This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.

Details

It is not designed not be call by the user

Super classes

`sbm::SBM` -> `sbm::SimpleSBM` -> `missSBM::SimpleSBM_fit` -> `missSBM::SimpleSBM_fit_noCov` -> `SimpleSBM_MNAR_noCov`

Active bindings

`imputation` the matrix of imputed values

`vExpec` double: variational approximation of the expectation complete log-likelihood

Methods

Public methods:

- `SimpleSBM_fit_MNAR$new()`
- `SimpleSBM_fit_MNAR$update_parameters()`
- `SimpleSBM_fit_MNAR$update_blocks()`
- `SimpleSBM_fit_MNAR$clone()`

Method `new()`**:** constructor for simpleSBM_fit for missSBM purpose

Usage:

```
SimpleSBM_fit_MNAR$new(networkData, clusterInit)
```

Arguments:

`networkData` a structure to store network under missing data condition: either a matrix possibly with NA, or a missSBM:::partlyObservedNetwork

`clusterInit` Initial clustering: a vector with size `ncol(adjacencyMatrix)`, providing a user-defined clustering with `nbBlocks` levels.

Method `update_parameters()`**:** update parameters estimation (M-step)

Usage:

```
SimpleSBM_fit_MNAR$update_parameters(nu = NULL)
```

Arguments:

`nu` currently imputed values

Method `update_blocks()`**:** update variational estimation of blocks (VE-step)

Usage:

```
SimpleSBM_fit_MNAR$update_blocks(log_lambda = 0)
```

Arguments:

`log_lambda` additional term sampling dependent used to de-bias estimation of tau

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
SimpleSBM_fit_MNAR$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

SimpleSBM_fit_noCov *This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.*

Description

This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.

This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.

Details

It is not designed not be call by the user

Super classes

sbm::SBM -> sbm::SimpleSBM -> missSBM::SimpleSBM_fit -> SimpleSBM_fit_noCov

Active bindings

`imputation` the matrix of imputed values

`vExpec` double: variational approximation of the expectation complete log-likelihood

`vExpec_corrected` double: variational approximation of the expectation complete log-likelihood with correction to be comparable with MNAR criteria

Methods

Public methods:

- `SimpleSBM_fit_noCov$update_parameters()`
- `SimpleSBM_fit_noCov$update_blocks()`
- `SimpleSBM_fit_noCov$clone()`

Method `update_parameters()`**:** update parameters estimation (M-step)

Usage:

```
SimpleSBM_fit_noCov$update_parameters(...)
```

Arguments:

`...` additional arguments, only required for MNAR cases

Method `update_blocks()`**:** update variational estimation of blocks (VE-step)

Usage:

```
SimpleSBM_fit_noCov$update_blocks(...)
```

Arguments:

`...` additional arguments, only required for MNAR cases

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
SimpleSBM_fit_noCov$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`SimpleSBM_fit_withCov` *This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.*

Description

This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.

This internal class is designed to adjust a binary Stochastic Block Model in the context of missSBM.

Details

It is not designed not be call by the user

Super classes

`sbm::SBM` -> `sbm::SimpleSBM` -> `missSBM::SimpleSBM_fit` -> `SimpleSBM_fit_withCov`

Active bindings

`imputation` the matrix of imputed values

`vExpec` double: variational approximation of the expectation complete log-likelihood

`vExpec_corrected` double: variational approximation of the expectation complete log-likelihood with correction to be comparable with MNAR criteria

Methods

Public methods:

- `SimpleSBM_fit_withCov$update_parameters()`
- `SimpleSBM_fit_withCov$update_blocks()`
- `SimpleSBM_fit_withCov$clone()`

Method `update_parameters()`**:** update parameters estimation (M-step)

Usage:

```
SimpleSBM_fit_withCov$update_parameters(...)
```

Arguments:

`...` use for compatibility

`control` a list to tune nlopt for optimization, see documentation of nloptr

Method `update_blocks()`**:** update variational estimation of blocks (VE-step)

Usage:

```
SimpleSBM_fit_withCov$update_blocks(...)
```

Arguments:

`...` use for compatibility

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
SimpleSBM_fit_withCov$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

`snowballSampler` *Class for defining a snowball sampler*

Description

Class for defining a snowball sampler

Class for defining a snowball sampler

Super classes

`missSBM::networkSampling` -> `missSBM::networkSampler` -> `missSBM::nodeSampler` -> `snowballSampler`

Methods

Public methods:

- `snowballSampler$new()`
- `snowballSampler$clone()`

Method `new()`**:** constructor for networkSampling

Usage:

```
snowballSampler$new(parameters = NA, adjacencyMatrix = NA, directed = FALSE)
```

Arguments:

`parameters` the vector of parameters associated to the sampling at play

`adjacencyMatrix` the adjacency matrix of the network

`directed` logical, directed network of not

Method `clone()`**:** The objects of this class are cloneable with this method.

Usage:

```
snowballSampler$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

summary.missSBM_fit *Summary method for a* missSBM_fit

Description

Summary method for a missSBM_fit

Usage

```
## S3 method for class 'missSBM_fit'
summary(object, ...)
```

Arguments

object	an R6 object with class missSBM_fit
...	additional parameters for S3 compatibility.

Value

a basic printing output

war *War data set*

Description

This dataset contains two networks where the nodes are countries and an edge in network "belligerent" means that the two countries have been at least once at war between years 1816 to 2007 while an edge in network "alliance" means that the two countries have had a formal alliance between years 1816 to 2012. The network belligerent have less nodes since countries which have not been at war are not considered.

Usage

```
war
```

Format

A list with 2 two igraph objects, alliance and belligerent. Each graph have three attributes: 'name' (the country name), 'power' (a score related to military power: the higher, the better) and 'trade' (a score related to the trade effort between pairs of countries).

Source

networks were extracted from https://correlatesofwar.org/

References

Sarkees, Meredith Reid and Frank Wayman (2010). Resort to War: 1816 - 2007. Washington DC: CQ Press.

Gibler, Douglas M. 2009. International military alliances, 1648-2008. CQ Press

Examples

```
data(war)
class(war$belligerent)
igraph::gorder(war$alliance)
igraph::gorder(war$belligerent)
igraph::edges(war$alliance)
igraph::get.graph.attribute(war$alliance)
```

Index