

Package: mixedClust (via r-universe)

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Type Package

Title Co-Clustering of Mixed Type Data

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Description Implementation of the co-clustering method for mixed type data proposed in M. Selosse, J. Jacques, C. Biernacki (2018) <<https://hal.science/hal-01893457>>. It consists in clustering simultaneously the rows (observations) and the columns (features) of a heterogeneous data set.

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Imports Rcpp (>= 0.12.11), fda, methods

LinkingTo Rcpp, RcppProgress, RcppArmadillo

Suggests rmarkdown, ordinalClust, knitr

VignetteBuilder knitr

LazyData true

Depends R (>= 3.5.0)

NeedsCompilation yes

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Repository <https://cran.r-universe.dev>

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|----|---|
| M1 | <i>Matrix of simulated ordinal data</i> |
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Description

This is a toy dataset for running simple examples.

Usage

M1

Format

A mixed type data matrix with 50 lines and 120 columns. There are 40 categorical variables, 40 continuous variables, and 40 ordinal variables.

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| mixedCoclust | <i>Function to perform a co-clustering</i> |
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Description

This function performs a co-clustering on heterogeneous data sets by using the Multiple Latent Block model (cf references for further details).

Usage

```
mixedCoclust(x=matrix(0,nrow=1,ncol=1), idx_list=c(1), distrib_names,
             kr, kc, init, nbSEM, nbSEMBurn, nbRepeat=1, nbindmini, m=0,
             functionalData=array(0, c(1,1,1)), zrinit= 0 , zcinit=0,
             percentRandomB=0, percentRandomP=0)
```

Arguments

| | |
|---------------|--|
| x | Data matrix, of dimension $N \times J_{tot}$. The features with same type should be aside. The missing values should be coded as NA. |
| idx_list | Vector of length D. This argument is useful when variables are of different types. Element d should indicate where the variables of type d begins in matrix x. |
| distrib_names | Vector of length D. indicates the type of distribution to use. Must be among "Gaussian", "Multinomial", "BOS", "Poisson" or "Functional". Functional data must always be at the end. |
| kr | Number of row classes. |
| kc | Vector of length D. d-th element indicates the number of column clusters. |

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| <code>m</code> | Vector of length D . d -th element defines the ordinal and categorical data's number of levels. |
| <code>functionalData</code> | Data tensor of dimension $N \times J \times T$. |
| <code>nbSEM</code> | Number of SEM-Gibbs iterations realized to estimate parameters. |
| <code>nbSEMBurn</code> | Number of SEM-Gibbs burning iterations for estimating parameters. This parameter must be inferior to <code>nbSEM</code> . |
| <code>nbRepeat</code> | Number of times sampling on rows and on columns will be done at each SEM-Gibbs iteration. |
| <code>nbindmini</code> | Minimum number of cells belonging to a block. |
| <code>init</code> | String that indicates the kind of initialisation. Must be one of the following words : "kmeans", "random", "provided", "randomParams" or "randomBurnin". |
| <code>zrinit</code> | Vector of length N . When <code>init="provided"</code> , indicates the labels of each row. |
| <code>zcinit</code> | Vector of length J_{tot} . When <code>init="provided"</code> , indicates the labels of each column. |
| <code>percentRandomB</code> | Vector of length 2. Indicates the percentage of resampling when <code>init</code> is equal to "randomBurnin". |
| <code>percentRandomP</code> | Vector of length 2. Indicates the percentage of resampling when <code>init</code> is equal to "randomParams". |

Value

| | |
|---------------------------|---|
| <code>@V</code> | Matrix of dimension $N \times k_r$ such that $V[i,g]=1$ if i belongs to cluster g . |
| <code>@icl</code> | ICL value for co-clustering. |
| <code>@name</code> | |
| <code>@paramschain</code> | List of length <code>nbSEMBurn</code> . For each iteration of the SEM-Gibbs algorithm, the parameters of the blocks are stored. |
| <code>@pichain</code> | List of length <code>nbSEM</code> . Item i is a vector of length k_r which contains the row mixing proportions at iteration i . |
| <code>@rhochain</code> | List of length <code>nbSEM</code> . Item i is a list of length D whose d -th contains the column mixing proportions of groups of variables d , at iteration i . |
| <code>@zc</code> | List of length D . d -th item is a vector of length $J[d]$ representing the columns partitions for the group of variables d . |
| <code>@zr</code> | Vector of length N with resulting row partitions. |
| <code>@W</code> | List of length D . Item d is a matrix of dimension $J \times k_c[d]$ such that $W[j,h]=1$ if j belongs to cluster h . |
| <code>@m</code> | Vector of length D . d -th element represents the number of levels of d -th group of variables. |
| <code>@params</code> | List of length D . d -th item represents the blocks parameters for group of variables d . |
| <code>@pi</code> | Vector of length k_r . Row mixing proportions. |
| <code>@rho</code> | List of length D . d -th item represents the column mixing proportion for d -th group of variables. |

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| @xhat | List of length D. d-th item represents the d-th group of variables dataset, with missing values completed. |
| @zrchain | Matrix of dimension nbSEM*N. Row i represents the row cluster partitions at iteration i. |
| @zcchain | List of length D. Item d is a matrix of dimension nbSEM*J[d]. Row i represents the column cluster partitions at iteration i. |

Author(s)

Margot Selosse, Julien Jacques, Christophe Biernacki.

Examples

```

data(M1)
nbSEM=30
nbSEMBurn=20
nbindmini=1
init = "random"

kr=2
kc=c(2,2,2)
m=c(6,3)
d.list <- c(1,41,81)
distributions <- c("Multinomial","Gaussian","Bos")
res <- mixedCoclust(x = M1, idx_list = d.list,distrib_names = distributions,
                   kr = kr, kc = kc, m = m, init = init,nbSEM = nbSEM,
                   nbSEMBurn = nbSEMBurn, nbindmini = nbindmini)

```

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