

Package: cobiclust (via r-universe)

August 20, 2024

Type Package

Title Biclustering via Latent Block Model Adapted to Overdispersed Count Data

Version 0.1.0

Description Implementation of a probabilistic method for biclustering adapted to overdispersed count data. It is a Gamma-Poisson Latent Block Model. It also implements two selection criteria in order to select the number of biclusters.

License GPL-3

URL <https://github.com/julieaubert/cobiclust>

BugReports <https://github.com/julieaubert/cobiclust/issues>

Imports cluster, stats

Suggests spelling, testthat

Encoding UTF-8

Language en-US

LazyData true

RoxxygenNote 7.1.1

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

RemoteUrl <https://github.com/julieaubert/cobiclust>

RemoteRef HEAD

RemoteSha 834fdb2a4e429a27315b4f3e3f87b0898609408a

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cobiclust*Perform a biclustering adapted to overdispersed count data.***Description**

Perform a biclustering adapted to overdispersed count data.

Usage

```
cobiclust(
  x,
  K = 2,
  G = 3,
  nu_j = NULL,
  a = NULL,
  akg = FALSE,
  cvg_lim = 1e-05,
  nbiter = 5000,
  tol = 1e-04
)
```

Arguments

<code>x</code>	the input matrix of observed data.
<code>K</code>	an integer specifying the number of groups in rows.
<code>G</code>	an integer specifying the number of groups in columns.
<code>nu_j</code>	a vector of . The length is equal to the number of columns.
<code>a</code>	an numeric.
<code>akg</code>	a logical variable indicating whether to use a common dispersion parameter (<code>akg = FALSE</code>) or a dispersion parameter per cocluster (<code>akg = TRUE</code>).
<code>cvg_lim</code>	a number specifying the threshold used for convergence criterion (<code>cvg_lim = 1e-05</code> by default).
<code>nbiter</code>	the maximal number of iterations for the global loop of variational EM algorithm (<code>nbiter = 5000</code> by default).
<code>tol</code>	the level of relative iteration convergence tolerance (<code>tol = 1e-04</code> by default).

Value

An object of class `cobiclustering`

See Also

[cobiclustering](#) for the `cobiclustering` class.

Examples

```

npc <- c(50, 40) # nodes per class
KG <- c(2, 3) # classes
nm <- npc * KG # nodes
Z <- diag(KG[1]) %x% matrix(1, npc[1], 1)
W <- diag(KG[2]) %x% matrix(1, npc[2], 1)
L <- 70 * matrix(runif(KG[1] * KG[2]), KG[1], KG[2])
M_in_expectation <- Z %*% L %*% t(W)
size <- 50
M<-matrix(
  rnbnom(
    n = length(as.vector(M_in_expectation)),
    mu = as.vector(M_in_expectation), size = size)
  , nm[1], nm[2])
rownames(M) <- paste("OTU", 1:nrow(M), sep = "_")
colnames(M) <- paste("S", 1:ncol(M), sep = "_")
res <- cobiclust(M, K = 2, G = 3, nu_j = rep(1,120), a = 1/size, cvg_lim = 1e-5)

```

selection_criteria *Calculate selection criteria.*

Description

Calculate selection criteria.

Usage

```
selection_criteria(x, K, G)
```

Arguments

- x The output of the cobiclust function.
- K The number of groups in rows.
- G The number of groups in columns.

Value

- A dataframe with 7 columns.
- vICL the vICL selection criterion.
- BIC the BIC selection criterion.
- penKG the value of the BIC penalty.
- lb the value of the lower bound of the log-likelihood.
- entZW the value of the entropy of the latent variables Z and W.
- K the number of groups in rows.
- G the number of groups in columns.

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