

Package ‘cobiclust’

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Type Package

Title Biclustering via Latent Block Model Adapted to Overdispersed Count Data

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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.

Imports cluster

License GPL-2

Encoding UTF-8

LazyData true

RoxxygenNote 6.0.1

NeedsCompilation no

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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)
```

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 colums.
<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).

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(
  rnbinom(
```

```
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 <- biclust(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 biclust function. |
| K | The number of groups in rows. |
| G | The number of groups in columns. |

Value

A data frame 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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