

# Package ‘cobiclust’

October 12, 2022

**Type** Package

**Title** Biclustering via Latent Block Model Adapted to Overdispersed  
Count Data

**Version** 0.1.0

**Author** Julie Aubert

**Maintainer** Julie Aubert <julie.aubert@agroparistech.fr>

**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

**RoxygenNote** 6.0.1

**NeedsCompilation** no

**Repository** CRAN

**Date/Publication** 2018-10-15 18:50:07 UTC

## R topics documented:

cobiclust . . . . .	2
selection_criteria . . . . .	3

<b>Index</b>	<b>4</b>
--------------	----------

---

 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

x	the input matrix of observed data.
K	an integer specifying the number of groups in rows.
G	an integer specifying the number of groups in columns.
nu_j	a vector of . The length is equal to the number of columns.
a	an numeric.
akg	a logical variable indicating whether to use a common dispersion parameter (akg = FALSE) or a dispersion parameter per cocluster (akg = TRUE).
cvg_lim	a number specifying the threshold used for convergence criterion (cvg_lim = 1e-05 by default).
nbiter	the maximal number of iterations for the global loop of variational EM algorithm (nbiter = 5000 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]) %% matrix(1, npc[1], 1)
W <- diag(KG[2]) %% 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 <- 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.

# Index

cobiclust, 2

cobiclustering, 2

selection\_criteria, 3