

# Package: akmbiclust (via r-universe)

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**Title** Alternating K-Means Biclustering  
**Version** 0.1.0  
**Description** Implements the alternating k-means biclustering algorithm  
in Fraiman and Li (2020) <[arXiv:2009.04550](#)>.  
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`akmbiclust`*Alternating k-means biclustering*

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## Description

This function uses the alternating k-means biclustering algorithm to extract the k biclusters in the matrix X. See the paper "Biclustering with Alternating K-Means" for more details.

## Usage

```
akmbiclust(X, k, lambda = 0, nstart = 1)
```

## Arguments

X	Data matrix.
k	The number of biclusters.
lambda	Regularization parameter. Default is 0.
nstart	The number of random initializations. Default is 1.

## Value

A list containing three objects:

row_labels	The bicluster labels of every row.
col_labels	The bicluster labels of every column.
loss	The loss of the produced biclusters.

## Author(s)

Nicolas Fraiman and Zichao Li

## References

N. Fraiman and Z. Li (2020). Biclustering with Alternating K-Means. arXiv preprint arXiv:2009.04550.

## Examples

```
# we create a 100 by 100 matrix X which has an underlying 2 by 2 block structure.
# The entries in the two 50 by 50 blocks on the top left and bottom right follow
# i.i.d. normal with mean 0 and variance 4. The entries in the two 50 by 50 blocks
# on the top right and bottom left follow i.i.d. normal with mean 0 and variance 1.

X <- matrix(rnorm(10000, 0, 1), 100, 100)
X[1:50, 1:50] <- matrix(rnorm(2500, 0, 2), 50, 50)
X[51:100, 51:100] <- matrix(rnorm(2500, 0, 2), 50, 50)

# Alternating k-means biclustering
```

```
# Result: perfect
result <- akmbiclust(X, 2, lambda = 0, nstart = 100)
result$row_labels
result$col_labels

# Separate k-means clustering on the rows and columns
# Result: random
kmeans(X, 2)$cluster
kmeans(t(X), 2)$cluster
```

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