

## Package ‘kmodR’

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### Type Package

## Title K-Means with Simultaneous Outlier Detection

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**Author** David Charles Howe [aut, cre]

Maintainer: David Charles Howe <kmodR@edgecondition.com>

**Description** An implementation of the 'k-means--' algorithm proposed by Chawla and Giannis, 2013 in their paper, ``k-means-- : A unified approach to clustering and outlier detection. SIAM International Conference on Data Mining (SDM13)'', and using 'ordering' described by Howe, 2013 in the thesis, ``Clustering and anomaly detection in tropical cyclones''. Useful for creating (potentially) tighter clusters than standard k-means and simultaneously finding outliers inexpensively in multidimensional space.

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**LazyData** TRUE

Suggests test that

## NeedsCompilation no

## Repository CRAN

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kmod*K-Means clustering with simultaneous Outlier Detection*

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## Description

K-Means clustering with simultaneous Outlier Detection

## Usage

```
kmod(X, k = 5, l = 0, i_max = 100, conv_method = "delta_C",
conv_error = 0, allow_empty_c = FALSE)
```

## Arguments

X	matrix of numeric data or an object that can be coerced to such a matrix (such as a data frame with numeric columns only).
k	the number of clusters (default = 5)
l	the number of outliers (default = 0)
i_max	the maximum number of iterations permissible (default = 100)
conv_method	character: the method used to assess if kmod has converged (default = "delta_C")
conv_error	numeric: the tolerance permissible when assessing convergence (default = 0)
allow_empty_c	logical: set whether empty clusters are permissible (default = FALSE)

## Value

kmod returns a list comprising the following components

- @return k the number of clusters specified
- l the number of outliers specified
- C the set of cluster centroids
- C\_sizes cluster sizes
- C\_ss the sum of squares for each cluster
- L the set of outliers
- L\_dist\_sqr the distance squares for each outlier to C
- L\_index the index of each outlier in the supplied dataset
- XC\_dist\_sqr\_assign the distance square and cluster assignment of each point in the supplied dataset
- within\_ss the within cluster sum of squares (excludes outliers)
- between\_ss the between cluster sum of squares
- tot\_ss the total sum of squares
- iterations the number of iterations taken to converge

**Examples**

```
# a 2-dimensional example with 2 clusters and 5 outliers
x <- rbind(matrix(rnorm(100, sd = 0.3), ncol = 2),
            matrix(rnorm(100, mean = 1, sd = 0.3), ncol = 2))
colnames(x) <- c("x", "y")
(cl <- kmod(x, 2, 5))

# cluster a dataset with 8 clusters and 0 outliers
x <- kmod(x, 8)
```

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