Package 'GBClust'

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Title Generalized Bayes clustering
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Description This package is an implementation of several generalized Bayes clustering methods.
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R topics documented:
comp_medoids
kbinary
kbinary_select
kdiss
kdiss_select
kmeans2
kmeans_gibbs
Minkowski_gibbs
Index

2 kbinary

Description

Compute the medoids of a given clustering solution based on the corresponding dissimilarity matrix.

Usage

```
comp_medoids(D, cluster)
```

Arguments

D A n x n numeric matrix containing the dissimilarities, i.e. the output of the func-

tions dist or daisy.

cluster A clustering solution, i.e. the output of kdiss.

Value

medoids Indexes of the medoids.

kbinary K-binary clustering

Description

Perform the so-called k-binary clustering algorithm, for obtaining groups when the data are binary observations.

Usage

```
kbinary(x, k, nstart = 1, trace = FALSE)
```

Arguments

Х	binary matrix of data, or an object that can be coerced to such a matrix (such as a numeric vector or a data frame with all numeric columns).
k	The number of clusters to be considered. A random set of (distinct) rows in x is chosen as the initial centres.
nstart	Number of random sets that has been chosen
trace	logical: if true, tracing information on the progress of the algorithm is produced.

Value

```
cluster Labels of the clusters at convergence
centers The value of the centroids at convergence
loss Numeric value of the loss function at convergence
```

kbinary_gibbs 3

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kbinary_gibbs	K-dissimilarities algorithm with uncertain	ну дианијісанон

Description

Perform the Gibbs-sampling for the k-dissimilarities algorithm using the Minkowski distance; see dist.

Usage

```
kbinary_gibbs(
    x,
    k,
    lambda = 1,
    R = 1000,
    burn_in = 1000,
    nstart = 10,
    trace = FALSE
)
```

Arguments

X	numeric matrix of of the data
k	The number of clusters to be considered.
lambda	Gibbs posterior tuning parameter
R	Number of MCMC samples after burn-in
burn_in	Number of MCMC samples to be discarded as burn-in period
nstart	Number of random initializations for the k-means algorithm
trace	logical: if true, tracing information on the progress of the algorithm is produced.

Value

```
G The letters of the alphabet

lambda A vector of numbers

loss A vector of numbers

G_map A vector of numbers

loss_map A vector of numbers
```

4 kdiss

kbinary_select	Selection of the number of cluster for the k-binary algorithm

Description

It displays the value of the loss function for various choices of k

Usage

```
kbinary_select(x, k_max, nstart = 1)
```

Arguments

X	numeric matrix of data, or an object that can be coerced to such a matrix (such as a numeric vector or a data frame with all numeric columns).
k_max	The maximum number of clusters to be considered. A random set of (distinct) rows in x is chosen as the initial centres.
nstart	Number of random sets that has been chosen

Value

It plots the loss function for different clustering solutions

|--|

Description

Perform the k-dissimilarities algorithm described.

Usage

```
kdiss(D, k, nstart = 1, trace = FALSE)
```

Arguments

D	A n x n numeric matrix with the dissimilarities, typically the output of ${\tt dist}$ or ${\tt daisy}$.
k	The number of clusters to be considered. See kdiss_select for selection criteria.
nstart	Number of random initializations.
trace	logical: if true, tracing information on the progress of the algorithm is produced

Value

cluster Labels of the clusters at convergence
loss Numeric value of the loss function at convergence

kdiss_select 5

ŀ	kdiss_select	Selection of the number of cluster for the k-dissimilarities algorithm

Description

It displays the value of the loss function / average silhouette width, for different values of k

Usage

```
kdiss_select(D, k_max, nstart = 1, method = "elbow")
```

Arguments

D	A n \times n numeric matrix with the dissimilarities, typically the output of dist or daisy.
k_max	Maximum number of clusters to be considered.
nstart	Number of random initializations.
method	The graph that will be displayed. Supported options are method="elbow", which displays the loss function, or method="silhouette". See silhouette for details about the latter.

Value

It return a ggplot2 graph of the loss function / average silhouette width, for $k=1,\ldots,k_max$.

kmeans2	K-Means^2 clustering	

Description

Perform k-means and k-means^2 on a data matrix.

Usage

```
kmeans2(x, k, nstart = 1, algorithm = "kmeans", trace = FALSE)
```

Arguments

X	numeric matrix of data, or an object that can be coerced to such a matrix (such as a numeric vector or a data frame with all numeric columns).
k	The number of clusters to be considered. A random set of (distinct) rows in x is chosen as the initial centres.
nstart	Number of random sets that has been chosen
algorithm	The algorithm to be used
trace	logical: if true, tracing information on the progress of the algorithm is produced.

6 kmeans_gibbs

Value

cluster Labels of the clusters at convergence centers The value of the centroids at convergence loss Numeric value of the loss function at convergence

kmeans2_select

Selection of the number of cluster for the k-dissimilarities algorithm

Description

It displays the value of the loss function for various choices of k

Usage

```
kmeans2_select(x, k_max, nstart = 1, algorithm = "kmeans")
```

Arguments

numeric matrix of data, or an object that can be coerced to such a matrix (such as a numeric vector or a data frame with all numeric columns).
 k_max
 The maximum number of clusters to be considered. A random set of (distinct) rows in x is chosen as the initial centres.
 nstart
 Number of random sets that has been chosen

algorithm The algorithm to be used, either kmeans or kmeans2

Value

It plots the loss function for different clustering solutions

kmeans_gibbs

K-means clustering with uncertainty quantification

Description

Perform the Gibbs-sampling for the k-means algorithm.

Usage

```
kmeans_gibbs(
    x,
    k,
    a_lambda,
    b_lambda,
    R = 1000,
    burn_in = 1000,
    nstart = 10,
    trace = FALSE
)
```

Minkowski_gibbs 7

Arguments

Χ	A n x d numeric matrix of the data.
k	The number of clusters to be considered.
a_lambda	Hyperparameter of the Gamma prior on the scale parameter
b_lambda	Hyperparameter of the Gamma prior on on the scale parameter
R	Number of MCMC samples after burn-in
burn_in	Number of MCMC samples to be discarded as burn-in period
nstart	Number of random initializations for the k-means algorithm
trace	logical: if true, tracing information on the progress of the algorithm is produced.

Value

G ARxn matrix including the cluster labels for each MCMC iteration lambda ARvector of numbers loss Avector of numbers G_map Avector of numbers loss_map Avector of numbers

Minkowski_gibbs

K-dissimilarities algorithm with uncertainty quantification

Description

Perform the Gibbs-sampling for the k-dissimilarities algorithm using the Minkowski distance. This function is complementary to kdiss, which is used instead to get a point estimate.

Usage

```
Minkowski_gibbs(
    x,
    k,
    p,
    a_lambda = 0,
    b_lambda = 0,
    R = 1000,
    burn_in = 1000,
    nstart = 10,
    trace = FALSE
)
```

Arguments

x	numeric matrix of of the data
k	The number of clusters to be considered.
р	Power of the Minkowski distance
a_lambda	Hyperparameter of the Gamma prior on the scale parameter. The default a_lambda = 0 leads to an improper prior.

8 Minkowski_gibbs

b_lambda Hyperparameter of the Gamma prior on on the scale parameter. The default

 a_1 ambda = 0 leads to an improper prior.

R Number of MCMC samples after burn-in.

burn_in Number of MCMC samples to be discarded as burn-in period.

nstart Number of random initializations for the kdiss algorithm, used to initialize the

MCMC chain.

trace logical: if true, tracing information on the progress of the algorithm is produced.

Value

G Labels of the clusters at each MCMC iteration.

lambda Numeric vector of the values of lambda at each MCMC iteration.

loss Numeric vector of the loss function at each MCMC iteration.

G_map Labels of the clusters obtained using kdiss, representing the maximum a posteriori.

loss_map Numeric value of the loss function obtained using kdiss, representing the maximized loss.

Index

```
comp_medoids, 2
daisy, 2, 4, 5
dist, 2-5
ggplot2, 5
kbinary, 2
kbinary_gibbs, 3
kbinary_select, 4
kdiss, 2, 4, 7, 8
kdiss_select, 4, 5
kmeans2, 5
kmeans2_select, 6
kmeans_gibbs, 6
Minkowski_gibbs, 7
silhouette, 5
```