

Package ‘cstab’

October 12, 2022

Type Package

Title Selection of Number of Clusters via Normalized Clustering
Instability

Version 0.2-2

Date 2018-06-19

Author Jonas M. B. Haslbeck, Dirk U. Wulff

Maintainer Jonas M. B. Haslbeck <jonas.haslbeck@gmail.com>

Description Selection of the number of clusters in cluster analysis using
stability methods.

BugReports <https://github.com/jmbh/cstab/issues>

License GPL (>= 2)

Depends R (>= 3.1.0), Rcpp (>= 0.11.4)

Imports cluster, fastcluster,

LinkingTo Rcpp

RoxygenNote 6.0.1

NeedsCompilation yes

Repository CRAN

Date/Publication 2018-06-19 16:59:43 UTC

R topics documented:

cDistance	2
cluster_example	3
cstab	3
cStability	4
lookup	5
plot.cstab	6
print.cstab	6
Index	7

cDistance

Selection of number of clusters via distance-based measures

Description

Selection of number of clusters via *gap statistic*, *jump statistic*, and *slope statistic*

Usage

```
cDistance(data, kseq, method = "kmeans", linkage = "complete",
          kmIter = 10, gapIter = 10)
```

Arguments

data	a n x p data matrix of type numeric.
kseq	a vector with considered numbers clusters $k > 1$
method	character string indicating the clustering algorithm. 'kmeans' for the k-means algorithm, 'hierarchical' for hierarchical clustering.
linkage	character specifying the linkage criterion, in case type='hierarchical'. The available options are "single", "complete", "average", "mcquitty", "ward.D", "ward.D2", "centroid" or "median". See hclust .
kmIter	integer specifying the the number of restarts of the k-means algorithm in order to avoid local minima.
gapIter	integer specifying the number of simulated datasets to compute the <i>gap statistic</i> (see Tibshirani et al., 2001).

Value

a list with the optimal numbers of cluster determined by the *gap statistic* (Tibshirani et al., 2001), the *jump Statistic* (Sugar & James, 2011) and the *slope statistic* (Fujita et al., 2014). Along the function returns the *gap*, *jump* and *slope* for each k in kseq.

Author(s)

Dirk U. Wulff <dirk.wulff@gmail.com> Jonas M. B. Haslbeck <jonas.haslbeck@gmail.com>

References

- Tibshirani, R., Walther, G., & Hastie, T. (2001). Estimating the number of clusters in a data set via the gap statistic. *Journal of the Royal Statistical Society: Series B (Statistical Methodology)*, 63(2), 411-423.
- Sugar, C. A., & James, G. M. (2011). Finding the number of clusters in a dataset. *Journal of the American Statistical Association*, 98(463), 750-763,
- Fujita, A., Takahashi, D. Y., & Patriota, A. G. (2014). A non-parametric method to estimate the number of clusters. *Computational Statistics & Data Analysis*, 73, 27-39.

Examples

```
## Not run:
# Generate Data from Gaussian Mixture
s <- .1
n <- 50
data <- rbind(cbind(rnorm(n, 0, s), rnorm(n, 0, s)),
              cbind(rnorm(n, 1, s), rnorm(n, 1, s)),
              cbind(rnorm(n, 0, s), rnorm(n, 1, s)),
              cbind(rnorm(n, 1, s), rnorm(n, 0, s)))
plot(data)

# Selection of Number of Clusters using Distance-based Measures
cDistance(data, kseq=2:10)

## End(Not run)
```

cluster_example	<i>Cluster example</i>
-----------------	------------------------

Description

An example, 2-dimensional dataset containing the 100 points for each of five bivariate normal distributions arranged equidistant along the outline of a circle.

Usage

```
cluster_example
```

Format

An object of class `matrix` with 500 rows and 2 columns.

Details

To inspect execute `plot(cluster_example)`.

cstab	<i>cstab: Selection of number of clusters via normalized clustering instability</i>
-------	---

Description

Selection of the number of clusters in cluster analysis using stability methods.

Details

Package: cstab
 Type: Package
 Version: 0.01
 Date: 2016-07-26
 License: GPL (>= 2)

Author(s)

Dirk U. Wulff <dirk.wulff@gmail.com> Jonas M. B. Haslbeck <jonas.haslbeck@gmail.com>

cStability

Selection of number of clusters via clustering instability

Description

Selection of number of clusters via *model-based* or *model-free*, *normalized* or *unnormalized* clustering instability.

Usage

```
cStability(data, kseq = 2:20, nB = 10, norm = TRUE, predict = TRUE,
  method = "kmeans", linkage = "complete", kmIter = 5, pbar = TRUE)
```

Arguments

data	a n x p data matrix of type numeric.
kseq	a vector with considered numbers clusters $k > 1$
nB	an integer specifying the number of bootstrap comparisons.
norm	logical specifying whether the instability path should be normalized. If TRUE, the instability path is normalized, accounting for a trivial decrease in instability due to a increasing k (see Haslbeck & Wulff, 2016).
predict	boolean specifying whether the model-based or the model-free variant should be used (see Haslbeck & Wulff, 2016).
method	character string specifying the clustering algorithm. 'kmeans' for the k-means algorithm, 'hierarchical' for hierarchical clustering.
linkage	character specifying the linkage criterion, in case type='hierarchical'. The available options are "single", "complete", "average", "mcquitty", "ward.D", "ward.D2", "centroid" or "median". See hclust .
kmIter	integer specifying the the number of restarts of the k-means algorithm in order to avoid local minima.
pbar	logical

Value

a list that contains the optimal k selected by the unnormalized and normalized instability method. It also includes a vector containing the averaged instability path (over bootstrap samples) and a matrix containing the instability path of each bootstrap sample for both the normalized and the unnormalized method.

Author(s)

Dirk U. Wulff <dirk.wulff@gmail.com> Jonas M. B. Haslbeck <jonas.haslbeck@gmail.com>

References

Ben-Hur, A., Elisseeff, A., & Guyon, I. (2001). A stability based method for discovering structure in clustered data. *Pacific symposium on biocomputing*, 7, 6-17.

Tibshirani, R., & Walther, G. (2005). Cluster validation by prediction strength. *Journal of Computational and Graphical Statistics*, 14(3), 511-528.

Examples

```
## Not run:
# Generate Data from Gaussian Mixture
s <- .1
n <- 50
data <- rbind(cbind(rnorm(n, 0, s), rnorm(n, 0, s)),
             cbind(rnorm(n, 1, s), rnorm(n, 1, s)),
             cbind(rnorm(n, 0, s), rnorm(n, 1, s)),
             cbind(rnorm(n, 1, s), rnorm(n, 0, s)))
plot(data)

# Selection of Number of Clusters using Instability-based Measures
stab_obj <- cStability(data, kseq=2:10)
print(stab_obj)

## End(Not run)
```

lookup

Create lookup table

Description

Create lookup table for faculties

Usage

```
lookup(n = 10000L, root = 200)
```

Arguments

n integer specifying the number of
 root numeric specifying the root used to avoid machine limit.

plot.cstab *Plot method for cstab objects*

Description

plot.cstab plots *instability* path.

Usage

```
## S3 method for class 'cstab'
plot(x, ...)
```

Arguments

x a cstab object (output of functions cStability).
 ... additional arguments passed to print.

Author(s)

Jonas M. B. Haslbeck <jonas.haslbeck@gmail.com> Dirk U. Wulff <dirk.wulff@gmail.com>

print.cstab *Print method for cstab objects*

Description

print.cstab prints key variables of cstab objects.

Usage

```
## S3 method for class 'cstab'
print(x, ...)
```

Arguments

x a cstab object (output of functions cStability).
 ... additional arguments passed to print.

Author(s)

Jonas M. B. Haslbeck <jonas.haslbeck@gmail.com> Dirk U. Wulff <dirk.wulff@gmail.com>

Index

* datasets

- cluster_example, 3

- cDistance, 2
- cluster_example, 3
- cstab, 3
- cstab-package (cstab), 3
- cStability, 4

- hclust, 2, 4

- lookup, 5

- plot.cstab, 6
- print.cstab, 6