Package ‘bootcluster’

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Title Bootstrapping Estimates of Clustering Stability

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R topics documented:

  k.select ................................................................. 2
  k.select_ref ........................................................... 3
  network.stability ..................................................... 4
Estimate number of clusters by bootstrapping stability

Usage

k.select(x, range = 2:7, B = 20, r = 5, threshold = 0.8, scheme_2 = TRUE)

Arguments

x | a data.frame of the data set
range | a vector of integer values, of the possible numbers of clusters k
B | number of bootstrap re-samplings
r | number of runs of k-means
threshold | the threshold for determining k
scheme_2 | logical TRUE if scheme 2 is used, FALSE if scheme 1 is used

Details

This function estimates the number of clusters through a bootstrapping approach, and a measure Smin, which is based on an observation-wise similarity among clusterings. The number of clusters k is selected as the largest number of clusters, for which the Smin is greater than a threshold. The threshold is often selected between 0.8 ~ 0.9. Two schemes are provided. Scheme 1 uses the clustering of the original data as the reference for stability calculations. Scheme 2 searches across the clustering samples that gives the most stable clustering.

Value

profile | a vector of Smin measures for determining k
k | integer estimated number of clusters

Author(s)

Han Yu
References

Bootstrapping estimates of stability for clusters, observations and model selection. Han Yu, Brian Chapman, Arianna DiFlorio, Ellen Eischen, David Gotz, Matthews Jacob and Rachael Hageman Blair.

Examples

```r
set.seed(1)
data(wine)
x0 <- wine[,2:14]
x <- scale(x0)
k.select(x, range = 2:10, B=20, r=5, scheme_2 = TRUE)
```

<table>
<thead>
<tr>
<th>k.select_ref</th>
<th>Estimate number of clusters</th>
</tr>
</thead>
</table>

Description

Estimate number of clusters by bootstrapping stability

Usage

```r
k.select_ref(df, k_range = 2:7, n_ref = 5, B = 100, B_ref = 50, r = 5)
```

Arguments

- **df**: data.frame of the input dataset
- **k_range**: integer valued vector of the numbers of clusters k to be tested upon
- **n_ref**: number of reference distribution to be generated
- **B**: number of bootstrap re-samples
- **B_ref**: number of bootstrap resamples for the reference distributions
- **r**: number of runs of k-means

Details

This function uses the out-of-bag scheme to estimate the number of clusters in a dataset. The function calculate the Smin of the dataset and at the same time, generate a reference dataset with the same range as the original dataset in each dimension and calculate the Smin_ref. The differences between Smin and Smin_ref at each k, Smin_diff(k), is taken into consideration as well as the standard deviation of the differences. We choose the k to be the argmax of ( Smin_diff(k) - ( Smin_diff(k+1) + (Smin_diff(k+1))))). If Smin_diff(k) less than 0.1 for all k in k_range, we say k = 1.
network.stability

Value

profile vector of ( Smin_diff(k) - ( Smin_diff(k+1) + se(Smin_diff(k+1)) ) ) measures for researchers’s inspection

k estimated number of clusters

Author(s)

Tianmou Liu

References

Bootstrapping estimates of stability for clusters, observations and model selection. Han Yu, Brian Chapman, Arianna DiFlorio, Ellen Eischen, David Gotz, Matthews Jacob and Rachael Hageman Blair.

Examples

```r
set.seed(1)
data(iris)
df <- data.frame(iris[,1:4])
df <- scale(df)
k.select_ref(df, k_range = 2:7, n_ref = 5, B=500, B_ref = 500, r=5)
```

network.stability

Estimate of detect module stability

Description

Estimate of detect module stability

Usage

```r
network.stability(
data.input,
threshold,
B = 20,
cor.method,
large.size,
PermuNo,
scheme_2 = FALSE
)
```
Arguments

data.input  a data.frame of the data set where the rows are observations and columns are covariates
threshold  a numeric number of threshold for correlation matrix
B  number of bootstrap re-samplings
cor.method  the correlation method applied to the data set, three method are available: "pearson", "kendall", "spearman"
large.size  the smallest set of modules, the large.size=0 is recommended to use right now.
PermuNo  number of random graphs for null
scheme_2  logical TRUE if scheme 2 is used, FALSE if scheme 1 is used. Right now, only FALSE is recommended.

Details

This function estimates the modules' stability through bootstrapping approach for the given threshold. The approach to stability estimation is to compare the module composition of the reference correlation graph to the various bootstrapped correlation graphs, and to assess the stability at the (1) node-level, (2) module-level, and (3) overall.

Value

stability.result  a list of result for nodes-wise stability
modularity.result  list of modularity information with the given threshold
jaccard.result  list estimated unconditional observed stability and the estimates of expected stability under the null
original.info  list information for original data, igraph object and adjacency matrix constructed with the given threshold

Author(s)

Mingmei Tian

References


Examples

set.seed(1)
data(wine)
x0 <- wine[1:50,]

mytest<-network.stability(data.input=x0, threshold=0.7, B=20,
cor.method='pearson', large.size=0,
PermuNo = 10,
network.stability.output

Plot method for objects from threshold.select

Description

Plot method for objects from threshold.select

Usage

network.stability.output(input, optimal.only = FALSE)

Arguments

input a list of results from function threshold.select
optimal.only a logical value indicating whether only plot the network with optimal threshold or not. The default is False, generating all network figures with a large number of nodes could take some time.

Details

network.stability.output is used to generate a series of network plots based on the given threshold.seq, where the nodes are colored by the level of stability. The network with optimal threshold value selected by function threshold.select is colored as red.

Value

Plot of network figures

Author(s)

Mingmei Tian

References

Examples

```r
set.seed(1)
data(wine)
x0 <- wine[1:50,]

mytest <- threshold.select(data.input = x0, threshold.seq = seq(0.1, 0.5, by = 0.05), B = 20, cor.method = "pearson", large.size = 0, PermuNo = 10, no_cores = 1, scheme_2 = FALSE)
network.stability.output(mytest)
```

---

**ob.stability**

*Estimate the stability of a clustering based on non-parametric bootstrap out-of-bag scheme, with option for subsampling scheme*

**Description**

Estimate the stability of a clustering based on non-parametric bootstrap out-of-bag scheme, with option for subsampling scheme.

**Usage**

```r
ob.stability(x, k, B = 500, r = 5, subsample = FALSE, cut_ratio = 0.5)
```

**Arguments**

- **x**
  - data.frame of the data set where the rows as observations and columns as dimensions of features
- **k**
  - number of clusters for which to estimate the stability
- **B**
  - number of bootstrap re-samples
- **r**
  - integer parameter in the kmeansCBI() function
- **subsample**
  - logical parameter to use the subsampling scheme option in the resampling process (instead of bootstrap)
- **cut_ratio**
  - numeric parameter between 0 and 1 for subsampling scheme training set ratio

**Details**

This function estimates the stability through out-of-bag observations. It estimate the stability at the (1) observation level, (2) cluster level, and (3) overall.
Value

- membership vector of membership for each observation from the reference clustering
- obs_wise vector of estimated observation-wise stability
- clust_wise vector of estimated cluster-wise stability
- overall numeric estimated overall stability
- Smin numeric estimated Smin through out-of-bag scheme

Author(s)

Tianmou Liu

References

Bootstrapping estimates of stability for clusters, observations and model selection. Han Yu, Brian Chapman, Arianna DiFlorio, Ellen Eischen, David Gotz, Matthews Jacob and Rachael Hageman Blair.

Examples

```r
set.seed(123)
data(iris)
df <- data.frame(iris[,1:4])
# You can choose to scale df before clustering by
# df <- scale(df)
ob.stability(df, k = 2, B=500, r=5)
```

---

**stability**  
*Estimate clustering stability of k-means*

Description

Estimate of k-means bootstrapping stability

Usage

```
stability(x, k, B = 20, r = 5, scheme_2 = TRUE)
```

Arguments

- `x` a data.frame of the data set
- `k` a integer number of clusters
- `B` number of bootstrap re-samplings
- `r` number of runs of k-means
- `scheme_2` logical TRUE if scheme 2 is used, FALSE if scheme 1 is used
Details

This function estimates the clustering stability through bootstrapping approach. Two schemes are provided. Scheme 1 uses the clustering of the original data as the reference for stability calculations. Scheme 2 searches across the clustering samples that gives the most stable clustering.

Value

- `membership`: a vector of membership for each observation from the reference clustering
- `obs_wise`: vector of estimated observation-wise stability
- `overall`: numeric, estimated overall stability

Author(s)

Han Yu

References

Bootstrapping estimates of stability for clusters, observations and model selection. Han Yu, Brian Chapman, Arianna DiFlorio, Ellen Eischen, David Gotz, Matthews Jacob and Rachael Hageman Blair.

Examples

```r
set.seed(1)
data(wine)
x0 <- wine[,2:14]
x <- scale(x0)
stability(x, k = 3, B=20, r=5, scheme_2 = TRUE)
```

---

**threshold.select**  
Estimate of the overall Jaccard stability

Description

Estimate of the overall Jaccard stability

Arguments

data.input: a data.frame of the data set where the rows are observations and columns are covariates
threshold.seq: a numeric sequence of candidate threshold
B: number of bootstrap re-samplings
.cor.method: the correlation method applied to the data set, three methods are available: "pearson", "kendall", "spearman"
threshold.select

large.size the smallest set of modules, the large.size=0 is recommended to use right now.
PermuNo number of random graphs for the estimation of expected stability
no_cores a integer number of CPU cores on the current host (This function can’t not be used yet).

Details

threshold.select is used to estimate of the overall Jaccard stability from a sequence of given threshold candidates, threshold.seq.

Value

stabilityresult a list of result for nodes-wise stability
modularityresult a list of modularity information with each candidate threshold
jaccardresult a list estimated unconditional observed stability and the estimates of expected stability under the nul
originalinformation a list information for original data, igraph object and adjacency matrix constructed with each candidate threshold
threshold.seq a list of candidate threshold given to the function

Author(s)

Mingmei Tian

References


Examples

```r
set.seed(1)
data(wine)
x0 <- wine[1:50,]
mytest<-threshold.select(data.input=x0,threshold.seq=seq(0.5,0.8,by=0.05), B=20,
cor.method='pearson',large.size=0,
PermuNo = 10,
no_cores=1,
scheme_2 = FALSE)
```
Wine Data Set

Description

These data are the results of a chemical analysis of wines grown in the same region in Italy but derived from three different cultivars. The analysis determined the quantities of 13 constituents found in each of the three types of wines.

Usage

data(wine)

Format

The data set wine contains a data.frame of 14 variables. The first variable is the types of wines. The other 13 variables are quantities of the constituents.

References

https://archive.ics.uci.edu/ml/datasets/wine
Index

k.select, 2
k.select_ref, 3

network.stability, 4
network.stability.output, 6

ob.stability, 7
stability, 8
threshold.select, 9
wine, 11