

Package ‘FADPclust’

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Type Package

Title Functional Data Clustering Using Adaptive Density Peak Detection

Version 1.1.1

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Description An implementation of a clustering algorithm for functional data based on adaptive density peak detection technique, in which the density is estimated by functional k-nearest neighbor density estimation based on a proposed semi-metric between functions. The proposed functional data clustering algorithm is computationally fast since it does not need iterative process. (Alex Rodriguez and Alessandro Laio (2014) <[doi:10.1126/science.1242072](https://doi.org/10.1126/science.1242072)>; Xiaofeng Wang and Yifan Xu (2016) <[doi:10.1177/0962280215609948](https://doi.org/10.1177/0962280215609948)>).

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Description

Clustering of univariate or multivariate functional data by finding cluster centers from estimated density peaks. FADPclust is a non-iterative procedure that incorporates KNN density estimation algorithm. The number of clusters can also be selected by the user or selected automatically through an internal clustering criterion.

Usage

```
FADPclust(fdata, cluster = 2:10, method = "FADP1", proportion = NULL,
          f.cut = 0.15, pve = 0.9, stats = "Avg.silhouette")
```

Arguments

<code>fdata</code>	for univariate functional data clustering: a functional data object produced by <code>fd()</code> function of <code>fda</code> package; for multivariate functional data clustering: a list of functional data objects produced by <code>fd()</code> function of <code>fda</code> package.
<code>cluster</code>	integer, or a vector of integers specifying the pool of the number of clusters in automatic variation. The default is <code>2:10</code> .
<code>method</code>	character string specifying the method used to calculate the pseudo functional k-nearest neighbor density. Valid options are <code>'FADP1'</code> and <code>'FADP2'</code> (see details in references). The default is <code>'FADP1'</code> .
<code>proportion</code>	numeric, a number or numeric vector of numbers within the range <code>[0,1]</code> , specifying to automatically select the smoothing parameter <code>k</code> in density estimation (see details). The default is <code>0.1, 0.2, ..., 1</code> .
<code>f.cut</code>	numeric, a number within the range <code>[0,1]</code> , specified to automatically select cluster centroids from the decision plot. The default is <code>0.15</code> .
<code>pve</code>	numeric, a number within the range <code>[0,1]</code> , the proportion of variance explained: used to choose the number of functional principal components. The default is <code>0.9</code> . When the method is chosen to be <code>'FADP1'</code> , there is no need to specify parameter <code>'pve'</code> for univariate functional data clustering.
<code>stats</code>	character string specifying the distance based statistics for cluster validation and determining the number of clusters. Valid options are <code>'Avg.silhouette'</code> , <code>'Dunn'</code> , and <code>'CH'</code> (See the description document of the <code>cluster.stats</code> function in the <code>fpc</code> R package for more details about these statistics). The default is <code>"Avg.silhouette"</code> .

Details

Given `n` functional objects or curves, `FADPclust()` calculates $f(x)$ and $\delta(x)$ for each object based on the semi-metric distance (see details in references), where $f(x)$ is the local density calculated by the functional k-nearest neighbor density estimator of curve x , and $\delta(x)$ is the shortest semi-metric distance between sample curve x and y for all samples y such that $f(x) \leq f(y)$. Functional

objects or curves with large f and large δ values are labeled class centroids. In other words, they appear as isolated points in the upper right corner of the f vs δ plot (the decision plot, see details in FADPplot). After cluster centroids are determined, other objects are clustered according to their semi-metric distances to the closest centroids.

The smoothing parameter k in functional k -nearest neighbor density estimation must be explicitly provided. Following Lauter (1988)'s idea, suggest that the optimal size of k satisfies a certain proportion, $k = a \cdot n^{4/5}$, where a is a parameter about the optimal proportion to be determined. Here, users enter variable 'proportion' to specify the parameter a .

Value

An 'FADPclus' object that contains the list of the following items.

- nclus: number of clusters.
- para: smoothing parameter k selected automatically by KNN estimation.
- method: character string introducing the method used to calculate the smoothing parameter.
- clust: cluster assignments. A vector of the same length as the number of observations.
- density: final density vector $f(x)$.
- delta: final delta vector $\delta(x)$.
- center: indices of the clustering centers.
- Avg.silhouette: average silhouette score from the final clustering result.
- Dunn: Dunn statistics from the final clustering result.
- CH: CH statistics from the final clustering result.

References

- Lauter, H. (1988), "Silverman, B. W.: "Density Estimation for Statistics and Data Analysis.," Biometrical Journal, 30(7), 876-877.
- Wang, X. F., and Xu, Y. (2016), "Fast Clustering Using Adaptive Density Peak Detection," Statistical Methods in Medical Research.
- Rodriguez, A., and Laio, A. (2014), "Machine learning. Clustering by fast search and find of density peaks," Science, 344(6191), 1492.
- Liu Y, Ma Z, and Yu F. (2017), "Adaptive density peak clustering based on K-nearest neighbors with aggregating strategy," Knowledge-Based Systems, 133(oct.1), 208-220.

See Also

[FADPsummary](#), [FADPplot](#).

Examples

```
###univariate functional data
data("simData1")
plot(simData1, xlab = "x", ylab = "y")
FADP1.sil.ans <- FADPclus(fdata = simData1, cluster = 2:5, method = "FADP1",
                        proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
```

```

                                stats = "Avg.silhouette")
FADPsummary(FADP1.sil.ans); FADPplot(FADP1.sil.ans)

FADP1.dunn.ans <- FADPclust(fdata = simData1, cluster = 2:5, method = "FADP1",
                          proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                          stats = "Dunn")
FADPsummary(FADP1.dunn.ans); FADPplot(FADP1.dunn.ans)

FADP1.ch.ans <- FADPclust(fdata = simData1, cluster = 2:5, method = "FADP1",
                         proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                         stats = "CH")
FADPsummary(FADP1.ch.ans); FADPplot(FADP1.ch.ans)

FADP2.ans <- FADPclust(fdata = simData1, cluster = 2:5, method = "FADP2",
                      proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                      pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP2.ans); FADPplot(FADP2.ans)

###multivariate functional data
data("simData2")
FADP1.ans <- FADPclust(fdata = simData2, cluster = 2:5, method = "FADP1",
                      proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                      pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP1.ans); FADPplot(FADP1.ans)

FADP2.ans <- FADPclust(fdata = simData2, cluster = 2:5, method = "FADP2",
                      proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                      pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP2.ans); FADPplot(FADP2.ans)

```

FADPplot

Visualize the result of FADPclust

Description

Plot the f vs delta plot with selected centroids.

Usage

```
FADPplot(object, cols = "default")
```

Arguments

object	object of class 'FADPclust' that is returned from FADPclust().
cols	vector of colors used to distinguish different clusters. Ten default colors are given.

See Also

[FADPclust](#), [FADPsummary](#).

Examples

```
###univariate functional data
data("simData1")
plot(simData1, xlab = "x", ylab = "y")
FADP1.ans <- FADPclust(fdata = simData1, cluster = 2:5, method = "FADP1",
                    proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                    stats = "Avg.silhouette")
FADPsummary(FADP1.ans); FADPplot(FADP1.ans)

FADP2.ans <- FADPclust(fdata = simData1, cluster = 2:5, method = "FADP2",
                    proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                    pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP2.ans); FADPplot(FADP2.ans)

###multivariate functional data
data("simData2")
FADP1.ans <- FADPclust(fdata = simData2, cluster = 2:5, method = "FADP1",
                    proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                    pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP1.ans); FADPplot(FADP1.ans)

FADP2.ans <- FADPclust(fdata = simData2, cluster = 2:5, method = "FADP2",
                    proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
                    pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP2.ans); FADPplot(FADP2.ans)
```

FADPsummary

Summary of FADPclust

Description

Summarize the result obtained from the `FADPclust()` function.

Usage

```
FADPsummary(object)
```

Arguments

object object of class 'FADPclust' that is returned from `FADPclust()`.

See Also

[FADPclust](#), [FADPplot](#).

Examples

```

###univariate functional data
data("simData1")
plot(simData1, xlab = "x", ylab = "y")
FADP1.ans <- FADPclust(fdata = simData1, cluster = 2:5, method = "FADP1",
  proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
  stats = "Avg.silhouette")
FADPsummary(FADP1.ans); FADPplot(FADP1.ans)

FADP2.ans <- FADPclust(fdata = simData1, cluster = 2:5, method = "FADP2",
  proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
  pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP2.ans); FADPplot(FADP2.ans)

###multivariate functional data
data("simData2")
FADP1.ans <- FADPclust(fdata = simData2, cluster = 2:5, method = "FADP1",
  proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
  pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP1.ans); FADPplot(FADP1.ans)

FADP2.ans <- FADPclust(fdata = simData2, cluster = 2:5, method = "FADP2",
  proportion = seq(0.02, 0.2, 0.02), f.cut = 0.15,
  pve = 0.9, stats = "Avg.silhouette")
FADPsummary(FADP2.ans); FADPplot(FADP2.ans)

```

simData1

Simulated univariate functional data for method FADPclust

Description

Simulated univariate functional data, with 2 clusters each containing 100 sample curves, were for users to apply the method FADPclust.

Format

fd, see FDA R package for details.

simData2

Simulated multivariate functional data for method FADPclust

Description

Simulated three-dimensional multivariate functional data, with 2 clusters each containing 100 sample curves, were for users to apply the method FADPclust.

Format

fd, see FDA R package for details.

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