Package ‘streamMOA’

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Title Interface for MOA Stream Clustering Algorithms
Description Interface for data stream clustering algorithms implemented in the MOA (Massive Online Analysis) framework.
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Imports graphics, stats, methods
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BugReports https://github.com/mhahsler/streamMOA
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R topics documented:

DSC_CluStream .................................................. 2
DSC_ClusTree .................................................... 3
DSC_DenStream ................................................... 5
DSC_MOA ....................................................... 6
DSD_RandomRBFGeneratorEvents .......................... 7
Index .......................................................... 9
**DSC_CluStream**

**CluStream Data Stream Clusterer**

**Description**

Class implements the CluStream cluster algorithm for data streams.

**Usage**

```
DSC_CluStream(m = 100, horizon = 1000, t = 2, k=NULL)
```

**Arguments**

- **m**
  - Defines the maximum number of micro-clusters used in CluStream.
- **horizon**
  - Defines the time window to be used in CluStream.
- **t**
  - Maximal boundary factor (=Kernel radius factor). When deciding to add a new data point to a micro-cluster, the maximum boundary is defined as a factor of \( t \) of the RMS deviation of the data points in the micro-cluster from the centroid.
- **k**
  - Number of macro-clusters to produce using weighted k-means. NULL disables automatic reclustering.

**Details**

This is an interface to the MOA implementation of CluStream.

CluStream applies a weighted k-means algorithm for reclustering (see Examples section below).

**Value**

An object of class DSC_CluStream (subclass of DSC_Micro, DSC_MOA and DSC), or, if \( k \) is not NULL then an object of DSC_TwoStage.

**Author(s)**

Michael Hahsler and John Forrest

**References**


**See Also**

DSC, DSC_Micro, DSC_MOA
**DSC_ClusTree**

**ClusTree Data Stream Clusterer**

### Description

Class implements the ClusTree cluster algorithm for data streams.

### Usage

```
DSC_ClusTree(horizon = 1000, maxHeight = 8, lambda = NULL)
```

### Arguments

- **horizon**: Range of the (time) window.
- **maxHeight**: The maximum height of the tree.
- **lambda**: number used to override computed lambda (decay).

### Details

This is an interface to the MOA implementation of ClusTree.
Value

An object of class DSC_ClusTree (subclass of DSC, DSC_MOA, DSC_Micro).

Author(s)

Michael Hahsler and John Forrest

References


See Also

DSC, DSC_Micro, DSC_MOA

Examples

# data with 3 clusters and 5% noise
stream <- DSD_Gaussians(k=3, d=2, noise=0.05)

clustree <- DSC_ClusTree(maxHeight=3)
update(clustree, stream, 500)
clustree

# plot micro-clusters
plot(clustree, stream)

# recluster with k-means
kmeans <- DSC_Kmeans(k=3)
recluster(kmeans, clustree)
plot(kmeans, stream, type="both")

# create a two stage clusering using ClusTree and reachability reclustering
CTXReach <- DSC_Twostage(
    micro=DSC_ClusTree(maxHeight=3),
    macro=DSC_Reachability(epsilon = .15)
)
CTXReach
update(CTXReach, stream, 500)
plot(CTXReach, stream, type="both")
Description

Class implements the DenStream cluster algorithm for data streams.

Usage

```
DSC_DenStream(epsilon, mu = 1, beta = 0.2, lambda = 0.001,
    initPoints = 100, offline = 2, processingSpeed=1, recluster = TRUE, k=NULL)
```

Arguments

- **epsilon**: defines the epsilon neighbourhood which is the maximal radius of micro-clusters (r<=epsilon). Range: 0 to 1.
- **mu**: minpoints as the weight w a core-micro-clusters needs to be created (w>=mu). Range: 0 to max(int).
- **beta**: multiplier for mu to detect outlier micro-clusters given their weight w (w<beta x mu). Range: 0 to 1
- **lambda**: decay constant.
- **initPoints**: number of points to use for initialization via DBSCAN.
- **offline**: offline multiplier for epsilon. Range: between 2 and 20). Used for reachability reclustering
- **processingSpeed**: Number of incoming points per time unit (important for decay). Range: between 1 and 1000.
- **recluster**: logical; should the offline DBSCAN-based (i.e., reachability at a distance of epsilon) be performed?
- **k**: integer; tries to automatically chooses offline to find k macro-clusters.

Details

Interface to the DenStream implementation in MOA.

DenStream applies reachability (from DBSCAN) between micro-clusters for reclustering using epsilon x offline (defaults to 2) as the reachability threshold.

If k is specified it automatically chooses the reachability threshold to find k clusters. This is achieved using single-link hierarchical clustering.

Value

An object of class DSC_DenStream (subclass of DSC, DSC_MOA, DSC_Micro) or, for recluster=TRUE, an object of class DSC_TwoStage.
Author(s)

Michael Hahsler and John Forrest

References


See Also

DSC, DSC_Micro, DSC_MOA

Examples

# data with 3 clusters and 5% noise
stream <- DSD_Gaussians(k = 3, d = 2, noise = 0.05)

# use Den-Stream with reachability reclustering
denstream <- DSC_DenStream(epsilon = .05)
update(denstream, stream, 500)
denstream

# plot macro-clusters
plot(denstream, stream)

# plot micro-cluster
plot(denstream, stream, type = "micro")

# show micro and macro-clusters
plot(denstream, stream, type = "both")

# reclustering. Choose reclustering reachability threshold automatically to find 3 clusters
denstream2 <- DSC_DenStream(epsilon = .05, k = 3)
update(denstream2, stream, 500)
plot(denstream2, stream, type = "both")

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DSC_MOA

DSC_MOA Class

Description

An abstract class that inherits from the base class DSC and provides the common functions needed to interface MOA clusterers.
Details

DSC_MOA classes operate in a different way in that the centers of the micro-clusters have to be extracted from the underlying Java object. This is done by using rJava to perform method calls directly in the JRI and converting the multi-dimensional Java array into a local R data type.

Author(s)

Michael Hahsler and John Forrest

References


See Also

DSC

Description

A class that generates random data based on RandomRBFGeneratorEvents implemented in MOA.

Usage

DSD_RandomRBFGeneratorEvents(k = 3, d = 2, numClusterRange = 3L, kernelRadius = 0.07, kernelRadiusRange = 0, densityRange = 0, speed =100L, speedRange = 0L, noiseLevel = 0.1, noiseInCluster = FALSE, eventFrequency = 3000L, eventMergeSplitOption = FALSE, eventDeleteCreate = FALSE, modelSeed = NULL, instanceSeed = NULL)

Arguments

k The average number of centroids in the model.
d The dimensionality of the data.
numClusterRange Range for number of clusters.
kernelRadius The average radius of the micro-clusters.
kernelRadiusRange Deviation of the number of centroids in the model.
densityRange Density range.
speed Kernels move a predefined distance of 0.01 every X points.
speedRange Speed/Velocity point offset.
noiseLevel Noise level.
noiseInCluster Allow noise to be placed within a cluster.
eventFrequency Frequency of events.
eventMergeSplitOption Merge and split?
eventDeleteCreate Delete and create?
modelSeed Random seed for the model.
instanceSeed Random seed for the instances.

Details

There are an assortment of parameters available for the underlying MOA data structure, however, we have currently limited the available parameters to the arguments above. Currently the modelSeed and instanceSeed are set to default values every time a DSD_MOA is created, therefore the generated data will be the same. Because of this, it is important to set the seed manually when different data is needed.

The default behavior is to create a data stream with 3 clusters and concept drift. The locations of the clusters will change slightly, and they will merge with one another as time progresses.

Value

An object of class DSD_RandomRBFGeneratorEvent (subclass of DSD_MOA, DSD).

Author(s)

Michael Hahsler and John Forrest

References


See Also

DSD

Examples

stream <- DSD_RandomRBFGeneratorEvents()
get_points(stream, 10, class=TRUE)

## Not run:
animate_data(stream, n=5000, pointInterval=100, xlim=c(0,1), ylim=c(0,1))

## End(Not run)
Index

CluStream (DSC_CluStream), 2
clustream (DSC_CluStream), 2
ClusTree (DSC_ClusTree), 3
clustree (DSC_ClusTree), 3

DenStream (DSC_DenStream), 5
denstream (DSC_DenStream), 5
DSC, 2, 4, 6, 7
DSC_CluStream, 2
DSC_ClusTree, 3
DSC_DenStream, 5
DSC_Micro, 2, 4, 6
DSC_MOA, 2, 4, 6, 6
DSD, 8
DSD_RandomRBFGGeneratorEvents, 7