Package ‘HighDimOut’

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Description Three high-dimensional outlier detection algorithms and an outlier unification scheme are implemented in this package. The angle-based outlier detection (ABOD) algorithm is based on the work of Kriegel, Schubert, and Zimek [2008]. The subspace outlier detection (SOD) algorithm is based on the work of Kriegel, Kroger, Schubert, and Zimek [2009]. The feature bagging-based outlier detection (FBOD) algorithm is based on the work of Lazarevic and Kumar [2005]. The outlier unification scheme is based on the work of Kriegel, Kroger, Schubert, and Zimek [2011].

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Description

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Details

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References


Description

This function performs the basic and approximated version of angle-based outlier detection algorithm. The ABOD method is especially useful for high-dimensional data, as angle is a more robust measure than distance in high-dimensional space. The basic version calculate the angle variance based on the whole data. The results obtained are more reliable. However, the speed can be very slow. The approximated version calculate the angle variance based on a subset of data and thereby, increasing the calculation speed. This function is based on the work of Kriegel, H.P., Schubert, M., Zimek, A., Angle-based outlier detection in high dimensional data, 2008.

Usage

Func.ABOD(data, basic = FALSE, perc)

Arguments

data
is the data frame containing the observations. Each row represents an observation and each variable is stored in one column.

basic
is a logical value, indicating whether the basic method is used. The speed of basic version can be very slow if the data size is large.

perc
defines the percentage of data to use when calculating the angle variance. It is only needed when basic=F.

Value

The function returns the vector containing the angle variance for each observation

Examples

library(ggplot2)
res.ABOD <- Func.ABOD(data=TestData[,1:2], basic=FALSE, perc=0.2)
data.temp <- TestData[,1:2]
data.temp$Ind <- NA
data.temp[order(res.ABOD, decreasing = FALSE)[1:10],"Ind"] <- "Outlier"
data.temp[is.na(data.temp$Ind),"Ind"] <- "Inlier"
data.temp$Ind <- factor(data.temp$Ind)
ggplot(data = data.temp) + geom_point(aes(x = x, y = y, color=Ind, shape=Ind))
Feature-bagging outlier detection (FBOD) algorithm

Description

This function performs feature-bagging based outlier detection algorithm. The implemented method is based on the work of "Lazarevic, A., Kumar, V., Feature bagging for outlier detection, 2005". This method can be regarded as an ensemble method, which is based on the results of local outlier factor (LOF). During each iteration, a random subset of variables, whose size is randomly chosen between \(d/2\) to \(d\) (where \(d\) is the dimensionality of the input data), is selected. The LOF method is applied to calculate the LOF scores based on the selected data subset. The final score of FBOD is the cumulative sum of each iteration.

Usage

\[
\text{Func.FBOD}(\text{data, iter, k.nn})
\]

Arguments

- `data` is the data frame containing the observations. Each row represents an observation and each variable is stored in one column.
- `iter` is the iteration used.
- `k.nn` is the value used for calculating the LOF score.

Value

The function returns a vector containing the FBOD outlier scores for each observation.

Examples

```r
library(ggplot2)
res.FBOD <- Func.FBOD(data = TestData[,1:2], iter=10, k.nn=5)
data.temp <- TestData[,1:2]
data.temp$Ind <- NA
data.temp[order(res.FBOD, decreasing = TRUE)[1:10],"Ind"] <- "Outlier"
data.temp[is.na(data.temp$Ind),"Ind"] <- "Inlier"
data.temp$Ind <- factor(data.temp$Ind)
ggplot(data = data.temp) + geom_point(aes(x = x, y = y, color=Ind, shape=Ind))
```
**Func.SNN**

*A function to calculate the shared nearest neighbors (SNN)*

**Description**

This function calculate the shared nearest neighbors (SNN). SNN is reported to be more robust than k nearest neighbors. Firstly, the k nearest neighbor distances for each observation is calculated. Then, the shared nearest neighbor similarity is calculated based on the result of k nearest neighbor. Note that k.nn should be greater than k.sel.

**Usage**

```r
Func.SNN(data, k.nn, k.sel)
```

**Arguments**

- `data` is the data frame containing the observations (should be numeric data). Each row represents an observation and each variable is stored in one column.
- `k.nn` specifies the value used for calculating the shared nearest neighbors.
- `k.sel` specifies the number of shared nearest neighbors

**Value**

The function returns the matrix containing the indices of top k shared nearest neighbors for each observation.

**Examples**

```r
Func.SNN(data=Test[,1:2], k.nn=5, k.sel=3)
```

---

**Func.SOD**

*Subspace outlier detection (SOD) algorithm*

**Description**

This function performs suspace outlier detection algorithm. The implemented method is based on the work of Krigel, H.P., Kroger, P., Schubert, E., Zimek, A., Outlier detection in axis-parallel subspaces of high dimensional data, 2009.

**Usage**

```r
Func.SOD(data, k.nn, k.sel, alpha = 0.8)
```
Arguments

data is the data frame containing the observations. Each row represents an observation and each variable is stored in one column.

k.nn specifies the value used for calculating the shared nearest neighbors. Note that k.nn should be greater than k.sel.

k.sel specifies the number shared nearest neighbors. It can be interpreted as the number of reference set for constructing the subspace hyperplane.

alpha specifies the lower limit for selecting subspace. 0.8 is set as default as suggested in the original paper.

Value

The function returns a vector containing the SOD outlier scores for each observation.

Examples

```r
library(ggplot2)
res.SOD <- Func.SOD(data = TestData[,1:2], k.nn = 10, k.sel = 5, alpha = 0.8)
data.temp <- TestData[,1:2]
data.temp$Ind <- NA
data.temp[order(res.SOD, decreasing = TRUE)[1:10],"Ind"] <- "Outlier"
data.temp[is.na(data.temp$Ind),"Ind"] <- "Inlier"
data.temp$Ind <- factor(data.temp$Ind)
ggplot(data = data.temp) + geom_point(aes(x = x, y = y, color=Ind, shape=Ind))
```

---

**Func.trans**  
*Outlier score transformation*

Description

This function calculate the transformed outlier scores, with the aim of unifying the results from different methods. The method is based on the work of Kriegel, H.P., Kroger, P., Schubert, E., Zimek, A., Interpreting and unifying outlier scores, 2011. It consists of two steps, regularization and normalization. For the ABOD scores, logarithmic inversion is used for regularization For the SOD scores, no action is taken to perform regularization For the FBOD method, the basic regularization, i.e., score-1, is used for regularization For the normalization step, the gaussian scaling method is used. The final output can be interpreted as the outlier probability, ranging from 0 to 1.

Usage

```
Func.trans(raw.score, method)
```

Arguments

- raw.score is the scores returned by each method
- method should be a character specifying the method used to generate the raw score. It has 3 possible values, "ABOD", "SOD", and "FBOD".

Description
The data contains the statistics of the players in one NBA team, i.e., Golden States Warriors, during the season 2013-2014. It can be obtained from the following website: http://www.basketball-reference.com. The data has 18 rows since there were 18 players shown in the lineups for the Golden States Warriors during that season. The data has 27 columns, including the player names, age, games played, games started, minutes played, field goal made, field goal attempts, field goal percentage, 3-pointers made, 3-pointer attempts, 3-pointer percentage, 2-pointers made, 2-pointer attempts, 2-pointer percentages, effective field goal percentage, free throws made, free throw attempts, free throw percentage, offensive rebounds, defensive rebounds, total rebounds, assists, steals, blocks, turnovers, personal fouls, and total points.

Usage
data(TestData)

Format
A data frame with 18 rows and 27 variables:
Examples

```r
library(ggplot2)
data(TestData)
ggplot(data = TestData, aes(x = x, y = y, shape=Lab, color=Lab)) + geom_point()
```
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