

Package ‘OutlierDM’

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Title Outlier detection for replicated high-throughput data

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Version 1.0-0

Description Detecting outlying observations for replicated high-throughput high-dimensional data based on a MA plot with quantile regression approach.

Depends R (>= 3.0.0), quantreg, MatrixModels, outliers, pcaPP

Imports methods, graphics

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OutlierDM-package	<i>Functions for detecting outlying observations in (multiple) replicated high-throughput data</i>
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Description

This package provides outlier detection algorithms for multiple replicated high-throughput high-dimensional data, especially in the field of mass spectrometry.

Details

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References

- S-H Eo, D Pak, J Choi, H Cho (2012). Outlier Detection for Multiplicative High-throughput Data. *BMC Research Notes*, **5**, 1–6.
- Cho et al. (2008). OutlierD: an R package for outlier detection using quantile regression on mass spectrometry data. *Bioinformatics*, **24**(6), 882–884.
- Min et al. (2007). Ultrahigh-pressure dual online solid phase extraction/capillary reverse-phase liquid chromatography/tandem mass spectrometry (DO-SPE/cRPLC/MS/MS): A versatile separation platform for high-throughput and highly sensitive proteomic analyses. *ELECTROPHORESIS*, **28**, 1012–1021.

See Also

[odm](#), [odm.control](#), [quantreg](#)

Examples

```
## Not run:  
data(lcms3)
```

```
## Fit projection approaches
fit.proj.const <- odm(lcms3, method="constant")
fit.proj.linear <- odm(lcms3, method="linear")
fit.proj.nonlin <- odm(lcms3, method="nonlin")
fit.proj.nonpara <- odm(lcms3, method="nonpar", lbda = 1)

par(mfrow = c(2,2))
plot(fit.proj.const, main = "Constant")
plot(fit.proj.linear, main = "Linear")
plot(fit.proj.nonlin, main = "NonLinear")
plot(fit.proj.nonpara, main = "Nonparametric")

## Fit pairwise OutlierD algorithm
fit0 <- odm(lcms3, type = "pair")
plot(fit0)

## End(Not run)
```

lcms3

LC/MS/MS data

Description

This data set consists of three-replicated LC/MS/MS data obtained from laboratory of gaseous ion chemistry in department of chemistry, Korea University. The data is already converted into log-2 transformation.

Usage

```
data(lcms3)
```

Format

a matrix for LCMS data, rows=peptides, columns=samples

Source

Min et al. (2007). Ultrahigh-pressure dual online solid phase extraction/capillary reverse-phase liquid chromatography/tandem mass spectrometry (DO-SPE/cRPLC/MS/MS): A versatile separation platform for high-throughput and highly sensitive proteomic analyses. *ELECTROPHORESIS*, **28**, 1012–1021.

Examples

```
data(lcms3)
```

odm

*Outlier detection for (multiple) replicated high-throughput data***Description**

This function is designed to detect outlying observations, or peptides, based on a MA plot with boxplot by quantile regression methods.

Usage

```
odm(x, k=1.5, method= c("nonlin", "constant", "linear", "nonpar"),
    type = c("proj", "diff", "pair", "grubbs", "dixon"), ...)
```

Arguments

x	data vectors or matrices. These can be given as named arguments. If the number of predictors is 2, x1 describes one n-by-1 vector for data and x2 describes the other n-by-1 vector for data (n= number of peptides, proteins, or genes)
k	a tuning parameter used in $Q1-k*IQR$ and $Q3+k*IQR$, $IQR=Q3-Q1$. Default value is 1.5
method	Select a type of quantile regression methods used in an outlier detection algorithm. Use one of "constant", "linear", "nonlin", and "nonpar" which mean constant, linear, non-linear, and nonparametric quantile regression in order. For more details, see the <code>quantreg</code> package.
type	input parameter for multiplicative detection algorithm. Use one of "dixon", "grubbs", "pair", "diff", and "proj". dixon: Dixon's Q-test grubbs: Grubbs' test pair: OutlierD algorithm (Cho et al., 2008) diff: Difference approach for multiplicative highthroughput data proj: Outlier Detection using projections for multiplicative experiments (Eo et al., 2012)
...	tuning parameters used in <code>odm.control()</code> . See odm.control .

Details

Outlier detection is an important issue in high-throughput data generated from various biological or chemical experiments such as mass spectrometry and microarrays because elaborate pre-processing is essential for successful analysis and outlier detection is one of pre-processing steps. The heterogeneity of variability and low replication are often obstacles for successful analysis including outlier detection. An outlier detection algorithm using quantile regression was developed to solve the problems; however, it can be used only for duplicate experiments. We here proposed an outlier detection algorithm using projection for multiplicative experiments.

Value

call:	evaluated function call
raw.data:	data to be used in the fitted model
res:	a data.frame including the information about the fitted model. It consists of several columns including outlier, M, A, Q3, Q1, UB and LB.
x.pair:	Object of class "list" ~~~
k:	a scalar parameter for constructing boxplot used in the fitted models
n.outliers:	a scalar value that denotes the number of outliers to be detected by the fitted model.
method:	the type of method used in the fitted model
type:	the type of algorithm used in the fitted model
contrl.para:	a list including information about tuning parameters

References

S-H Eo, D Pak, J Choi, H Cho (2012). Outlier Detection for Multiplicative High-throughput Data. *BMC Research Notes*, **5**, 1–6.

See Also

[OutlierDM-package](#) to show general information about the package
[OutlierDM-class](#) to provide information about the "OutlierDM" class
[odm.control](#) to control tuning parameters

Examples

```
## Not run:
data(lcms3)

## Fit projection approaches
fit.proj.const <- odm(lcms3, method="constant")
fit.proj.linear <- odm(lcms3, method="linear")
fit.proj.nonlin <- odm(lcms3, method="nonlin")
fit.proj.nonpara <- odm(lcms3, method="nonpar", lbda = 1)

par(mfrow = c(2,2))
plot(fit.proj.const, main = "Constant")
plot(fit.proj.linear, main = "Linear")
plot(fit.proj.nonlin, main = "NonLinear")
plot(fit.proj.nonpara, main = "Nonparametric")

## Fit pairwise OutlierD algorithm
fit0 <- odm(lcms3, type = "pair")
plot(fit0)

## End(Not run)
```

odm.control

*Control tuning parameters for "OutlierDM" object***Description**

various parameters that control aspects of the "OutlierDM" object

Usage

```
odm.control(pair.cre = 1, dist.mthd = "median",
            Lower = .25, Upper = .75, trans = "log2",
            centering = TRUE, projection.type = "PCA", lbda = 1,
            nonlin.method = "L-BFGS-B", nonlin.SS = "AsymOff",
            nonlin.Frank = c(2, -8, 0, 1), ncl = 2 )
```

Arguments

pair.cre	a scalar parameter to specify the minimum number of pairs, used in "type = pair".
dist.mthd	a distance parameter used in "type = diff". you can choose one of "median", "mean" or and so on.
Lower	a criterion for lower quantile value used to construct boxplot
Upper	a criterion for upper quantile value used to construct boxplot
trans	a parameter for a logarithm and exponential transformation. If a log 2 transformation is needed, set "trans = log2". If no transformation is needed, set "trans = FALSE".
centering	a logical parameter for the status of centering. If "centering = TRUE", data are centered by its column means.
projection.type	a parameter to determine a type of projection methods. Choose one of "naive", "pca", and "robust".
lbda	a criterion about lambda used for nonlinear quantile regression.
nonlin.method	a parameter to determine a type of methods used for nonlinear quantile regression. choose one of "L-BFGS-B" and "BFGS". Default is "L-BFGS-B".
nonlin.SS	a parameter to determine a type of structure used for nonlinear quantile regression. choose one of "Frank", "Self", "Asym" and "AsymOff". Default is "AsymOff", Asymptotic Regression Model with an Offset.
nonlin.Frank	a structure parameter used for Frank copula model. Gain c(df, delta, mu, sigma) in the Frank copula formula
ncl	A parameter to determine the number of cores used in parallel computing. A default value is 2.

See Also

[odm](#)

OutlierDM-class	Class "OutlierDM"
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Description

A class for outlier detection algorithms for high-throughput high-dimensional data.

Objects from the Class

Objects can be created by calls of the form `new("OutlierDM", ...)`. See following information about slots.

Slots

`call`: evaluated function call

`raw.data`: data to be used in the fitted model

`res`: a data.frame including the information about the fitted model. It consists of several columns including `outlier`, `M`, `A`, `Q3`, `Q1`, `UB` and `LB`.

`x.pair`: Object of class "list" ~~

`k`: a scalar parameter for constructing boxplot used in the fitted models

`n.outliers`: a scalar value that denotes the number of outliers to be detected by the fitted model.

`method`: the type of method used in the fitted model

`type`: the type of algorithm used in the fitted model

`contrl.para`: a list including information about tuning parameters

Methods

show signature(object = "OutlierDM"): Same as the show method without the optional arguments

plot signature(x = "OutlierDM", y = "missing"): Plot an object.

See Also

[odm](#)

Examples

```
showClass("OutlierDM")
```

plot *a plot-method for a "OutlierDM" object*

Description

This function provides a MA scatter plot with quantile regression based boxlplot.

Usage

```
## S4 method for signature 'OutlierDM'  
plot(x, y = NA, pch = 20, cex = 0.5, xlab = "A", ylab = "M",  
legend.use = TRUE, ...)
```

Arguments

x	fitted model object of class odm .
y	the "y" argument is not used in the plot-method for "OutlierDM" object.
pch	a vector of plotting characters or symbols: see plot.default .
cex	See plot.default .
xlab	label X
ylab	label Y
legend.use	logical option for using legend box
...	plot.default arguments

Details

This function is a method for the generic function `plot` for the S4 class `OutlierDM`. It can be invoked by calling `print` for an object of the appropriate class, or directly by calling `plot.OutlierDM` regardless of the class of the object.

See Also

[odm](#)

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