

# Package ‘ILS’

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stats

**Description** It performs interlaboratory studies (ILS) to detect those laboratories that provide non-consistent results when comparing to others. It permits to work simultaneously with various testing materials, from standard univariate, and functional data analysis (FDA) perspectives. The univariate approach based on ASTM E691-08 consist of estimating the Mandel's h and k statistics to identify those laboratories that provide more significant different results, testing also the presence of outliers by Cochran and Grubbs tests, Analysis of variance (ANOVA) techniques are provided (F and Tuckey tests) to test differences in means corresponding to different laboratories per each material. Taking into account the functional nature of data retrieved in analytical chemistry, applied physics and engineering (spectra, thermograms, etc.). ILS package provides a FDA approach for finding the Mandel's k and h statistics distribution by smoothing bootstrap resampling.

**License** GPL (>= 2)

**LazyData** yes

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---

|              |  |
|--------------|--|
| boot.sim.set | <i>Bootstrap samples of a functional statistic</i> |
|--------------|--|

---

**Description**

data.bootstrap provides bootstrap samples for functional data.

**Usage**

```
boot.sim.set(x, smo = 0.05)
```

**Arguments**

|     |  |
|-----|--|
| x   | A fdata object.                                    |
| smo | The smoothing parameter for the bootstrap samples. |

---

|              |   |
|--------------|---|
| cochran.test | <i>Function to compute the Grubbs test statistic.</i> |
|--------------|---|

---

## Description

Function to estimate the Cochran test statistic.

## Usage

```
cochran.test(x, ...)  
  
## Default S3 method:  
cochran.test(x, var.index = 1, replicate.index = 2,  
             material.index = 3, laboratory.index = 4, data.name = NULL,  
             alpha = 0.05, ...)  
  
## S3 method for class 'lab.qcdata'  
cochran.test(x, alpha = 0.05, ...)
```

## Arguments

|                  |   |
|------------------|---|
| x                | R object (used to select the method). See details.  |
| ...              | Arguments passed to or from methods.  |
| var.index        | Scalar with the column number corresponding to the observed variable (the critical to quality variable). Alternatively, a string with the name of a quality variable can be provided. |
| replicate.index  | Scalar with the column number corresponding to the index each replicate.  |
| material.index   | Scalar corresponding to the replicated number.  |
| laboratory.index | Scalar that defines the index number of each laboratory.  |
| data.name        | String specifying the name of the variable which appears on the plots. If name is not provided, it is retrieved from the object.  |
| alpha            | The significance level (0.05 by default)  |

## References

- Wilrich Peter-T. (2013), Critical values of mandel's h and k, the grubbs and the Cochran test statistic. *Asta-Advances in Statistical Analysis*, 97(1):1-10.
- ASTM E 691 (1999), Standard practice for conducting an interlaboratory study to determine the precision of a test method. American Society for Testing and Materials. West Conshohocken, PA, USA.

## Examples

```
library(ILS)
data(Glucose)
Glucose.qcdata <- lab.qcdata(Glucose)
str(Glucose.qcdata)
cochran.test(Glucose.qcdata)
```

---

DSC

*Differential Scanning Calorimetry curves*

---

## Description

90 samples of calcium oxalate were analyzed by differential scanning calorimetry technique (DSC), obtaining 90 DSC curves showing from an SDT device the difference of heat between a sample and an oxalate reference value depending on the temperature that the samples are heated at a constant temperature rate. The data set consists of 15 TG curves of 1000 observations from each of the 6 laboratories. Laboratory 2 to Laboratory 4 uses the same simultaneous SDT analyzer in similar conditions, Laboratory 6 uses a simultaneous SDT analyzer with an old calibration, and Laboratory 7 uses a simultaneous SDT analyzer with a calibration (2 degrees Celsius displaced from the zinc melting point).

## Format

5 x 1000 x 6 dimension array, where each matrix consists of the 15 DSC curves obtained by testing 15 different oxalate samples, and evaluated at 1000 different temperature values. These 15 curves were obtained for each of the 6 laboratories that performed the experiments.

## References

Naya, S., Tarrío-Saavedra, J., Lopez-Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". *Journal of Thermal Analysis and Calorimetry*, 118,1229-1243.

## Examples

```
library(ILS)
data(DSC)
summary(DSC)
```

---

func.ils.formula      *Descriptive measures for functional data.*

---

### Description

Central and dispersion measures for functional data.

### Usage

```
func.ils.formula(formula, data = NULL, drop = FALSE, func = func.mean)
```

### Arguments

|         |   |
|---------|---|
| formula | a formula, such as $y \sim \text{group}$ , where $y$ is a fdata object to be split into groups according to the grouping variable group (usually a factor). |
| data    | List that containing the variables in the formula. The item called "df" is a data frame with the grouping variable. The item called "y" is a fdata object.  |
| drop    | logical indicating if levels that do not occur should be dropped (if f is a factor or a list).  |
| func    | Measures for functional data.   |

---

Glucose      *Glucose in Serum*

---

### Description

Dataset corresponding to serum glucose (measurements of glucose concentration in blood used to control diabetes) testing. Eight laboratories conducted tests to five different blood samples tagged with different references, ranging them from low sugar content to very high. Three replicates were obtained for each sample. It is retrieved from ASTM E 691 standard.

### Format

A data frame with 120 observations composed of the following 4 variables:

**Glucose** Glucose content in Serum

**Replicate** Number of glucose measurement corresponding to each material

**Material** Level of glucose, ranging from low content of sugar to very high level of glucose in blood.

**Laboratory** Laboratories conducted tests

### References

ASTM E 691 (1999). Standard practice for conducting an interlaboratory study to determine the precision of a test method. American Society for Testing and Materials. West Conshohocken, PA, USA.

**Examples**

```
library(ILS)
data(Glucose)
summary(Glucose)
attach(Glucose)
str(Glucose)
table(Replicate,Material,Laboratory)
table(Laboratory,Material)
st <- with(Glucose, tapply(Glucose, list(Material,Laboratory), mean))
st
```

---

grubbs.test

*Function to compute the Grubbs test statistic.*


---

**Description**

Function to estimate the Grubbs test statistic.

**Usage**

```
grubbs.test(x, ...)
```

## Default S3 method:

```
grubbs.test(x, var.index = 1, replicate.index = 2,
  material.index = 3, laboratory.index = 4, data.name = NULL,
  alpha = 0.05, ...)
```

## S3 method for class 'lab.qcdata'

```
grubbs.test(x, alpha = 0.05, ...)
```

**Arguments**

|                  |   |
|------------------|---|
| x                | an R object (used to select the method). See details.   |
| ...              | arguments passed to or from methods.  |
| var.index        | Scalar with the column number corresponding to the observed variable (the critical to quality variable). Alternatively, a string with the name of a quality variable can be provided. |
| replicate.index  | Scalar with the column number corresponding to the index each replicate.  |
| material.index   | Scalar corresponding to the replicated number.  |
| laboratory.index | Scalar that defines the index number of each laboratory.  |
| data.name        | String specifying the name of the variable which appears on the plots. If name is not provided, it is retrieved from the object.  |
| alpha            | The significance level (0.05 for default)   |

## References

- Wilrich Peter-T. (2013), Critical values of Mandel's h and k, the Grubbs and the Cochran test statistic. *Asta-Advances in Statistical Analysis*, 97(1):1-10.
- ASTM E 691 (1999), Standard practice for conducting an interlaboratory study to determine the precision of a test method. American Society for Testing and Materials. West Conshohocken, PA, USA.

## Examples

```
library(ILS)
data(Glucose)
Glucose.qcdata<- lab.qcdata(Glucose)
str(Glucose.qcdata)
grubbs.test(Glucose.qcdata)
```

---

h.qcs

*Function to estimate the univariate Mandel's h statistic*

---

## Description

This function is used to compute the Mandel's h statistic.

## Usage

```
h.qcs(x, ...)

## Default S3 method:
h.qcs(x, var.index = 1, replicate.index = 2,
      material.index = 3, laboratory.index = 4, data.name = NULL,
      alpha = 0.05, ...)

## S3 method for class 'lab.qcdata'
h.qcs(x, alpha = 0.05, ...)
```

## Arguments

|                 |   |
|-----------------|---|
| x               | R object (used to select the method). See details.  |
| ...             | Arguments passed to or from methods.  |
| var.index       | Scalar with the column number corresponding to the observed variable (the critical to quality variable). Alternatively, a string with the name of a quality variable can be provided. |
| replicate.index | Scalar with the column number corresponding to the index each replicate.  |
| material.index  | Scalar corresponding to the replicated number.  |

|                  |  |
|------------------|--|
| laboratory.index | Scalar that defines the index number of each laboratory.   |
| data.name        | String specifying the name of the variable which appears on the plots. If name is not provided, it is retrieved from the object. |
| alpha            | The significance level (0.05 by default)   |

## References

- Wilrich Peter-T. (2013), Critical values of Mandel's h and k, the Grubbs and the Cochran test statistic. *Asta-Advances in Statistical Analysis*, 97(1):1-10.
- ASTM E 691 (1999), Standard practice for conducting an interlaboratory study to determine the precision of a test method. American Society for Testing and Materials. West Conshohocken, PA, USA.

## Examples

```
library(ILS)
data(Glucose)
Glucose.qcdata <- lab.qcdata(Glucose)
str(Glucose.qcdata)
h<- h.qcs(Glucose.qcdata, alpha = 0.005)
summary(h)
plot(h)
```

---

|     |  |
|-----|--|
| IDT | <i>Dataset composed of the initial decomposition temperature (IDT) of different samples of Calcium Oxalate, obtained by 7 different laboratories</i> |
|-----|--|

---

## Description

Initial decomposition temperature (IDT) is a parameter defined by temperature at which a material loss 5% of its weight when it is heated using a constant rate. One hundred and five calcium oxalate samples were tested by thermogravimetric analysis (TG), obtaining 105 TG curves from which the IDT is extracted. Summarizing, IDT dataset is composed of the IDT values of calcium oxalate obtained by 7 different laboratories that analyze 15 oxalate samples each one.: Laboratory 1 uses a simultaneous thermal analyzer (STA) with an old calibration program, Laboratory 2 to Laboratory 4 use a SDT simultaneous analyzer, Laboratory 6 utilizes a SDT simultaneous analyzer with an old calibration, and Laboratory 7 uses a SDT simultaneous analyzer with a biased calibration (2 degrees Celsius shifted from the zinc melting point).

## Format

Dataframe of dimension 105 x 44. The first column corresponds to IDT variable, the second (Sample) is the replicate number, the third is the tested material (Material), and fourth is the laboratory.



## References

Naya, S., Tarrío-Saavedra, J., Lopez- Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". *Journal of Thermal Analysis and Calorimetry*, 118,1229-1243.

## Examples

```
library(ILS)
data(IDT)
summary(IDT)
attach(IDT)
str(IDT)
table(Sample,Run,Laboratory)
table(Laboratory,Run)
st <- with(IDT, tapply(IDT, list(Run,Laboratory), mean))
st
```

---

ILS

*Interlaboratory Study*

---

## Description

Interlaboratory Study

## Details

It performs interlaboratory studies (ILS) to detect those laboratories that provide non-consistent results when comparing to others. It permits to work simultaneously with various testing materials, from standard univariate, and functional data analysis (FDA) perspectives. The univariate approach based on ASTM E691-08 consist of estimating the Mandel's h and k statistics to identify those laboratories that provide more significant different results, testing also the presence of outliers by Cochran and Grubbs tests, Analysis of variance (ANOVA) techniques are provided (F and Tuckey tests) to test differences in the testing variable means corresponding to test differences in means corresponding to different laboratories per each material. Taking into account the functional nature of data retrieved in analytical chemistry, applied physics and engineering (spectra, thermograms, etc.). ILS package provides a FDA approach for functional Mandel's k and h statistics by smoothing bootstrap resampling of distribution.

ils.fqcdata

*Functional Quality Control Data***Description**

It Creates an object of class 'ils.fqcd' to perform statistical quality control. This object is used to plot functional data.

**Usage**

```
ils.fqcdata(x, p = NULL, index.laboratory = NULL, argvals = NULL,
  rangeval = NULL, names = NULL)
```

**Arguments**

|                  |  |
|------------------|--|
| x                | A (n x m) matrix or data-frame. The m is the number of points observed in each curve, and n is the number of curves for each laboratory. |
| p                | the number of laboratories.  |
| index.laboratory | is the laboratory index. The index.laboratory length should be equal a p   |
| argvals          | Argvals, by default: 1:m.  |
| rangeval         | Range of discretization points, by default: range(argvals).  |
| names            | (optional) list with tree components: main an overall title, xlab title for x axis and ylab title for y axis.                            |

**References**

- Febrero-Bande, M. and Oviedo, M. (2012), "Statistical computing in functional data analysis: the R package fda.usc". Journal of Statistical Software 51 (4), 1-28.
- Naya, S., Tarrío-Saavedra, J., Lopez- Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". Journal of Thermal Analysis and Calorimetry, 118,1229-1243.

**Examples**

```
library(ILS)
data(TG)
delta <- seq(from = 40 ,to = 850 ,length.out = 1000 )
fqcddata <- ils.fqcdata(TG, p = 7, argvals = delta)
xlab <- "Temperature (C)"
ylab <- "Mass (%)"
main <- "TG curves obtained from calcium oxalate"
plot(x = fqcddata, main = main, xlab=xlab, ylab=xlab,col = 1:7,legend = TRUE)
```

---

`ils.fqcs`*It develops an object of class 'ils.fqcs'*

---

## Description

Create an object of class 'ils.fqcs' to perform statistical quality control. This function is used to compute requested FDA.

## Usage

```
ils.fqcs(x, ...)  
  
## Default S3 method:  
ils.fqcs(x, argvals = NULL, rangeval = NULL, ...)  
  
## S3 method for class 'ils.fqcdata'  
ils.fqcs(x, ...)  
  
## S3 method for class 'ils.fqcs'  
print(x, ...)  
  
## S3 method for class 'ils.fqcs'  
summary(object, ...)
```

## Arguments

|                       |   |
|-----------------------|---|
| <code>x</code>        | A <code>ils.fqcs</code> object for which a print is desired.              |
| <code>...</code>      | Arguments passed to or from methods.                                      |
| <code>argvals</code>  | Argvals, by default: 1:m.   |
| <code>rangeval</code> | Range of discretization points, by default: <code>range(argvals)</code> . |
| <code>object</code>   | A <code>ils.fqcs</code> object for which a summary is desired.            |

## References

- Febrero-Bande, M. and Oviedo, M. (2012), "Statistical computing in functional data analysis: the R package `fda.usc`". *Journal of Statistical Software* 51 (4), 1-28.
- Cuevas A., Febrero-Bande, M. and Fraiman, R. (2006), "On the use of the bootstrap for estimating functions with functional data". *Computational Statistics & Data Analysis* 51, 2, 1063-1074.
- Naya, S., Tarrío-Saavedra, J., Lopez- Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiaga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". *Journal of Thermal Analysis and Calorimetry*, 118,1229-1243.

**Examples**

```
library(ILS)
data(TG)
delta <- seq(from = 40 ,to = 850 ,length.out = 1000 )
fqcdata <- ils.fqcdata(TG, p = 7, argvals = delta)
xlab <- "Temperature/ C"
ylab <- "Mass/ %"
fqcstat <- ils.fqcs(fqcdata)
plot(fqcstat, xlab = xlab, ylab = ylab, legend = TRUE)
```

---

k.qcs

---

*Function to calcute the Mandel's k statistic*


---

**Description**

This function is used to compute the statistic k of Mandel.

**Usage**

```
k.qcs(x, ...)
```

```
## Default S3 method:
k.qcs(x, var.index = 1, replicate.index = 2,
      material.index = 3, laboratory.index = 4, data.name = NULL,
      alpha = 0.05, ...)
```

```
## S3 method for class 'lab.qcdata'
k.qcs(x, alpha = 0.05, ...)
```

**Arguments**

|                  |   |
|------------------|---|
| x                | an R object (used to select the method). See details.   |
| ...              | arguments passed to or from methods.  |
| var.index        | Scalar with the column number corresponding to the observed variable (the critical to quality variable). Alternatively, a string with the name of a quality variable can be provided. |
| replicate.index  | Scalar with the column number corresponding to the index each replicate.  |
| material.index   | Scalar corresponding to the replicated number.  |
| laboratory.index | Scalar that defines the index number of each laboratory.  |
| data.name        | String specifying the name of the variable which appears on the plots. If name is not provided, it is retrieved from the object.  |
| alpha            | The significance level (0.05 by default)  |

## References

Wilrich Peter-T. (2013), Critical values of Mandel's h and k, the Grubbs and the Cochran test statistic. *Asta-Advances in Statistical Analysis*, 97(1):1-10.

ASTM E 691 (1999), Standard practice for conducting an interlaboratory study to determine the precision of a test method. American Society for Testing and Materials. West Conshohocken, PA, USA.

## Examples

```
library(ILS)
data(Glucose)
Glucose.qcdata <- lab.qcdata(Glucose)
str(Glucose.qcdata)
k<- k.qcs(Glucose.qcdata, alpha = 0.005)
summary(k)
plot(k)
```

---

|         |                                    |
|---------|------------------------------------|
| lab.aov | <i>Function to compute the AOV</i> |
|---------|------------------------------------|

---

## Description

Function to compute the analysis of variance of ILS data, taking into account the laboratories and material factors.

## Usage

```
lab.aov(x, ...)

## Default S3 method:
lab.aov(x, var.index = 1, replicate.index = 2,
        material.index = 3, laboratory.index = 4, data.name = NULL,
        level = 0.95, plot = FALSE, pages = 0, ...)

## S3 method for class 'lab.qcdata'
lab.aov(x, level = 0.95, plot = FALSE, pages = 0,
        ...)
```

## Arguments

|           |   |
|-----------|---|
| x         | Object lab.qcdata.  |
| ...       | Arguments passed to or from methods.  |
| var.index | Scalar with the column number corresponding to the observed variable (the critical to quality variable). Alternatively, a string with the name of a quality variable can be provided. |

|                  |   |
|------------------|---|
| replicate.index  | Scalar with the column number corresponding to the index each replicate.  |
| material.index   | Scalar corresponding to the replicated number.  |
| laboratory.index | Scalar that defines the index number of each laboratory.  |
| data.name        | String specifying the name of the variable which appears on the plots. If name is not provided, it is retrieved from the object.  |
| level            | Requested confidence level (0.95 by default)  |
| plot             | If TRUE, confidence intervals are plot.   |
| pages            | By default 0, it indicates the number of pages over which to spread the output. For example, if pages=1, all terms will be plotted on one page with the layout performed automatically. If pages=0, one plot will be displayed by each tested material. |

## References

- WHothorn T., Bretz, F., and Westfall, P. (2008), Simultaneous inference in general parametric models. *Biometrical Journal*, 50(3):346-363.
- Heyden, Y., Smeyers-Verbeke, J. (2007), Set-up and evaluation of interlaboratory studies. *J. Chromatogr. A*, 1158:158-167.

## Examples

```
## Not run:
library(ILS)
data(Glucose)
Glucose.qcdata <- lab.qcdata(Glucose)
str(Glucose.qcdata)
lab.aov(Glucose.qcdata, level = 0.95, plot = TRUE, pages = 1)

## End(Not run)
```

---

lab.qcdata

*Quality Control Data*

---

## Description

It creates a 'lab.qcdata' class object to perform the interlaboratory study. This object is used to plot ILS data and more.

## Usage

```
lab.qcdata(data, var.index = 1, replicate.index = 2, material.index = 3,
  laboratory.index = 4, data.name = NULL)
```

**Arguments**

|                  |   |
|------------------|---|
| data             | Matrix or data-frame that contains the data, replicate index, type of material, and the laboratory.   |
| var.index        | Scalar with the column number corresponding to the observed variable (the critical to quality variable). Alternatively, a string with the name of a quality variable can be provided. |
| replicate.index  | Scalar with the column number corresponding to the index each replicate.  |
| material.index   | Scalar corresponding to the replicated number.  |
| laboratory.index | Scalar that defines the index number of each laboratory.  |
| data.name        | String specifying the name of the variable which appears on the plots. If name is not provided, it is retrieved from the object.  |

**Examples**

```
library(ILS)
data(Glucose)
Glucose.qcdata <- lab.qcdata(Glucose)
str(Glucose.qcdata)
summary(Glucose.qcdata)
```

---

|         |   |
|---------|---|
| lab.qcs | <i>Create an object of class 'lab.qcs' to perform statistical quality control. This function is used to compute statistics required for plotting Statistics</i> |
|---------|---|

---

**Description**

It develops an object of `lab.qcs-codelinkclass` to perform statistical quality control. This function is used to compute the requested statistics to be summarized and plotted.

**Usage**

```
lab.qcs(x, ...)

## S3 method for class 'lab.qcs'
print(x, ...)

## S3 method for class 'lab.qcs'
summary(object, ...)
```

**Arguments**

|        |  |
|--------|--|
| x      | Object <code>lab.qcdata</code> (Functional Quality Control Data) |
| ...    | Arguments passed to or from methods.                             |
| object | A <code>lab.qcs</code> object for which a summary is desired.    |

**Examples**

```
library(ILS)
data(Glucose)
Glucose.qcdata <- lab.qcdata(Glucose)
str(Glucose.qcdata)
Glucose.qcs <- lab.qcs(Glucose.qcdata)
str(Glucose.qcs)
summary(Glucose.qcs)
```

---

mandel.fqcs

*This function is used to compute the FDA Mandel's h and k statistic.*


---

**Description**

It develops an object of 'mandel.fqcs' class to perform statistical quality control analysis. This function is used to compute the functional approach of Mandel's h and k statistic. It is specifically designed to deal with experimental data results defined by curves such as thermograms and spectra.

**Usage**

```
mandel.fqcs(x, ...)
```

## Default S3 method:

```
mandel.fqcs(x, p = NULL, index.laboratory = NULL,
  argvals = NULL, rangeval = NULL, names = NULL, ...)
```

## S3 method for class 'ils.fqcsdata'

```
mandel.fqcs(x, fdep = depth.mode, outlier = TRUE,
  trim = 0.01, alpha = 0.01, nb = 200, smo = 0.05, ...)
```

**Arguments**

|                  |  |
|------------------|--|
| x                | A (n x m) matrix or data-frame. The m is the number of points observed in each curve, and n is the number of curves for each laboratory. |
| ...              | Further arguments passed to or from other methods.   |
| p                | the number of laboratories.  |
| index.laboratory | is the laboratory index. The index.laboratory length should be equal a p   |
| argvals          | Argvals, by default: 1:m.  |
| rangeval         | Range of discretization points, by default: range(argvals).  |
| names            | (optional) list with tree components: main an overall title, xlab title for x axis and ylab title for y axis.                            |
| fdep             | Type of depth measure, by default depth.mode.  |
| outlier          | = TRUE   |



|       |  |
|-------|--|
| trim  | The alpha of the trimming.                         |
| alpha | Significance level, by default 1%.                 |
| nb    | The number of bootstrap samples.                   |
| sno   | The smoothing parameter for the bootstrap samples. |

## References

- Febrero-Bande, M. and Oviedo, M. (2012), "Statistical computing in functional data analysis: the R package fda.usc". *Journal of Statistical Software* 51 (4), 1-28.
- Cuevas A., Febrero-Bande, M. and Fraiman, R. (2006), "On the use of the bootstrap for estimating functions with functional data". *Computational Statistics & Data Analysis* 51, 2, 1063-1074.
- Naya, S., Tarrío-Saavedra, J., Lopez- Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". *Journal of Thermal Analysis and Calorimetry*, 118,1229-1243.

## Examples

```
## Not run:
library(ILS)
data(TG)
delta <- seq(from = 40 ,to = 850 ,length.out = 1000 )
fqcddata <- ils.fqcddata(TG, p = 7, argvals = delta)
mandel.tg <- mandel.fqcs(fqcddata.tg,nb = 200)
plot(mandel.tg,legend = F,col=c(rep(3,5),1,1))

## End(Not run)
```

---

outliers.ils

*Detecting outliers for functional dataset*

---

## Description

Procedure for detecting functional outliers.

## Usage

```
outliers.ils(x, fdep = depth.FM, trim = 0.01)
```

## Arguments

|      |  |
|------|--|
| x    | A fdata  |
| fdep | Type of depth measure, by default depth.mode.    |
| trim | The percentaje of the trimming, by default is 1% |

---

plot.ils.fqcdata      *Plotting method for 'ils.fqcdata' objects*

---

### Description

Generic function to plot objects of 'ils.fqcdata' class

### Usage

```
## S3 method for class 'ils.fqcdata'
plot(x, type = "l", main = NULL, xlab = NULL,
     ylab = NULL, ylim = NULL, x.co = NULL, y.co = NULL, legend = TRUE,
     col = NULL, ...)
```

### Arguments

|        |  |
|--------|--|
| x      | Object ils.fqcdata (Functional Quality Control Data)   |
| type   | 1-character string giving the type of plot desired. The following values are possible for fdata class object: "l" for lines (by default), "p" for points, "o" for overplotted points and lines, "b", "c" for (empty if "c") points joined by lines, "s" and "S" for stair steps and "h" for histogram-like vertical lines. Finally, "n" does not produce any points or lines. The following values are possible for fdata2d class object: "image.contour" (by default) to display three-dimensional data and add the contour lines, "image" to display three-dimensional data, "contour" to display a contour plot, "persp" to display a perspective plots of a surface over the x-y plane and "filled.contour" to display a contour plot with the areas between the contours filled in solid color. |
| main   | Main title for the plot  |
| xlab   | Title for the x axis   |
| ylab   | Title for the y axis   |
| ylim   | The y limits of the plot   |
| x.co   | It specifies the x co-ordinates to be used to place a legend.  |
| y.co   | It specifies the y co-ordinates to be used to place a legend.  |
| legend | Logical argument. Default is TRUE then The legend default is used.   |
| col    | Color specifications   |
| ...    | Further arguments passed to matplot function (for fdata class).  |

### References

- Febrero-Bande, M. and Oviedo, M. (2012), "Statistical computing in functional data analysis: the R package fda.usc". *Journal of Statistical Software* 51 (4), 1-28.
- Naya, S., Tarrío-Saavedra, J., Lopez- Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". *Journal of Thermal Analysis and Calorimetry*, 118,1229-1243.

**Examples**

```
## Not run:
library(ILS)
data(TG)
delta <- seq(from = 40 ,to = 850 ,length.out = 1000 )
fqcdata <- ils.fqcdata(TG, p = 7, argvals = delta)
windows()
xlab <- "Temperature (C)"
ylab <- "Mass (%)"
main <- "TG curves obtained from calcium oxalate"
plot(x = fqcdata, main = main, xlab=xlab, ylab=ylab,legend = TRUE)
## End(Not run)
```

plot.ils.fqcs

*Plotting method for 'ils.fqcs' objects***Description**

Generic function to plot objects of 'ils.fqcs' class. Results of functional ILS studies are graphically shown.

**Usage**

```
## S3 method for class 'ils.fqcs'
plot(x, type = "l", xlab = NULL, ylab = NULL,
     legend = TRUE, col = NULL, ...)
```

**Arguments**

|        |  |
|--------|--|
| x      | Object functional data or a list with objects of functional data type  |
| type   | 1-character string giving the type of plot desired. The following values are possible for fdata class object: "l" for lines (by default), "p" for points, "o" for overplotted points and lines, "b", "c" for (empty if "c") points joined by lines, "s" and "S" for stair steps and "h" for histogram-like vertical lines. Finally, "n" does not produce any points or lines. The following values are possible for fdata2d class object: "image.contour" (by default) to display three-dimensional data and add the contour lines, "image" to display three-dimensional data, "contour" to display a contour plot, "persp" to display a perspective plots of a surface over the x-y plane and "filled.contour" to display a contour plot with the areas between the contours filled in solid color. |
| xlab   | Title for the x axis   |
| ylab   | Title for the y axis   |
| legend | Logical argument. Default is TRUE then The legend default is used.   |
| col    | Color specifications   |
| ...    | arguments to be passed to or from methods  |

## References

- Febrero-Bande, M. and Oviedo, M. (2012), "Statistical computing in functional data analysis: the R package fda.usc". Journal of Statistical Software 51 (4), 1-28.
- Naya, S., Tarrío-Saavedra, J., Lopez- Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". Journal of Thermal Analysis and Calorimetry, 118,1229-1243.

## Examples

```
library(ILS)
data(TG)
delta <- seq(from = 40 ,to = 850 ,length.out = 1000 )
fqcdata <- ils.fqcdata(TG, p = 7, argvals = delta)
xlab <- "Temperature/ C"
ylab <- "Mass/ %"
fqcstat <- ils.fqcs(fqcdata)
plot(fqcstat, xlab = xlab, ylab = ylab, legend = TRUE)
```

---

plot.lab.qcdata      *Plot method for 'lab.qcdata' objects*

---

## Description

Generic function for plotting objects of 'lab.qcdata' class. Results of univariate ILS studies are graphically shown.

## Usage

```
## S3 method for class 'lab.qcdata'
plot(x, xlab = NULL, ylab = NULL, col = "blue", ...)
```

## Arguments

|      |  |
|------|--|
| x    | Functional data object or a list with functional objects |
| xlab | Title for the x axis                                     |
| ylab | Title for the y axis                                     |
| col  | Color of type material, when there only one              |
| ...  | arguments to be passed to or from methods.               |

## Examples

```
library(ILS)
data(Glucose)
Glucose.qcdata <- lab.qcdata(Glucose)
str(Glucose.qcdata)
plot(Glucose.qcdata)
```

---

|              |  |
|--------------|--|
| plot.lab.qcs | <i>Plot method for 'lab.qcs' objects</i> |
|--------------|--|

---

**Description**

Generic function for plotting objects of 'lab.qcs' class. Results of univariate ILS studies are graphically shown.

**Usage**

```
## S3 method for class 'lab.qcs'
plot(x, title = NULL, xlab = NULL, ylab = NULL,
     col = NULL, ylim = NULL, ...)
```

**Arguments**

|       |  |
|-------|--|
| x     | Functional data object or a list with functional objects |
| title | Main title for the plot                                  |
| xlab  | Title for the x axis                                     |
| ylab  | Title for the y axis                                     |
| col   | Color specifications                                     |
| ylim  | Numeric vectors of length 2 (coordinates ranges)         |
| ...   | arguments to be passed to or from methods.               |

---

|                  |  |
|------------------|--|
| plot.mandel.fqcs | <i>Plotting method for 'mandel.fqcs' objects</i> |
|------------------|--|

---

**Description**

Generic function to plot objects of 'mandel.fqcs' class. Results of functional ILS studies are graphically shown.

**Usage**

```
## S3 method for class 'mandel.fqcs'
plot(x, xlab = NULL, ylab = NULL, x.co = NULL,
     y.co = NULL, legend = TRUE, col = NULL, ...)
```

**Arguments**

|        |  |
|--------|--|
| x      | A mandel.fqcs object   |
| xlab   | Title for the x axis   |
| ylab   | Title for the y axis   |
| x.co   | It specifies the x co-ordinates to be used to place a legend.      |
| y.co   | It specifies the y co-ordinates to be used to place a legend.      |
| legend | Logical argument. Default is TRUE then The legend default is used. |
| col    | Color specifications   |
| ...    | arguments to be passed to or from methods                          |

**References**

- Febrero-Bande, M. and Oviedo, M. (2012), "Statistical computing in functional data analysis: the R package fda.usc". *Journal of Statistical Software* 51 (4), 1-28.
- Naya, S., Tarrío-Saavedra, J., Lopez-Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". *Journal of Thermal Analysis and Calorimetry*, 118,1229-1243.

**Examples**

```
## Not run:
library(ILS)
data(TG)
delta <- seq(from = 40 ,to = 850 ,length.out = 1000 )
fqcdata <- ils.fqcdata(TG, p = 7, argvals = delta)
mandel.tg <- mandel.fqcs(fqcdata.tg,nb = 200)
plot(mandel.tg,legend = F,col=c(rep(3,5),1,1))

## End(Not run)
# $H(t)$ y $K(t)$
```

---

 TG

*Thermogravimetry curves*


---

**Description**

One hundred and five Calcium oxalate samples were tested by thermogravimetric (TG) analysis, obtaining 105 TG curves that shows the mass loss of oxalate depending on time when samples are heated at a constant temperature rate. Dataset is composed by fifteen TG curves of 1000 observations each of overall 7 different laboratories. Laboratory 1 uses a simultaneous thermal analyzer (STA) with an old calibration program, Laboratory 2 to Laboratory 4 use a SDT simultaneous analyzer, Laboratory 6 utilizes a SDT simultaneous analyzer with an old calibration, and Laboratory 7 uses a SDT simultaneous analyzer with a biased calibration (2 degrees Celsius shifted from the zinc melting point).

**Format**

A  $15 \times 1000 \times 7$  dimension array, where each matrix consists of the 15 TG curves obtained testing 15 different oxalate samples, and evaluated in 1000 different values of temperature. These 15 curves were obtained for each of the overall 7 laboratories that have performed the experiments.

**References**

Naya, S., Tarrío-Saavedra, J., Lopez-Beceiro, J., Francisco Fernandez, M., Flores, M. and Artiga, R. (2014), "Statistical functional approach for interlaboratory studies with thermal data". *Journal of Thermal Analysis and Calorimetry*, 118,1229-1243.

**Examples**

```
library(ILS)
data(TG)
summary(TG)
```

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