SUPPLEMENTARY FILE

to the paper

Validation of digestion and element separation methods and a new data reduction program (IsotopeHf®) for Lu-Hf isotope dilution analysis by MC-ICP-MS

by

Reneé González-Guzmán, Bodo Weber, María D. Tazzo-Rangel, and Luigi Solari

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Package 'IsotopeHf'

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Type Package
Title Hf isotope reduction Toolkit for R: IsotopeHf
Description The main functionality of the package is to provide reliable Lu- Hf data from Neptune® MC ICP MS measurements.
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Author Reneé González Guzmán
Maintainer Reneé González Guzmán <rguzman@cicese.edu.mx></rguzman@cicese.edu.mx>
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IsotopeHf-package
BHVO.Hf
BHVO.Lu
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Description

This package imports mass spectrometer raw data files from Neptune® MC ICP MS, facilitates the visualization of temporal trends and scatter during measurements and implements point-by-point application of mass fractionation correction, isobaric correction, isotope dilution computation, and spike stripping. In addition, provides a set of tools that compute useful Hf isotope parameters.

Details

Package: IsotopeHf
Type: Package
Version: 0.6.9
Date: 2015-02-15
License: BSD-3

BHVO.Hf

BHVO-1 basalt (Hf)

Description

The 'BHVO.Hf' data frame has 80 rows and 11 columns, which are the measurements from a Hf BHVO-1 basalt aliquot.

Details

Variables:

- Time (POSIXct format)
- 172 Mass
- 174 Mass
- 175 Mass

BHVO.Lu 3

- 176 Mass
- 177 Mass
- 179 Mass
- 180 Mass
- 181 Mass
- 182 Mass
- Sample ID

See Also

```
BHVO.Lu, Standard.Hf, Standards.Lu_Re
```

Examples

```
## Not run:
#Preloaded data
data (BHVO.Hf)
head (BHVO.Hf)
## End(Not run)
```

BHV0.Lu

BHVO-1 basalt (Lu)

Description

The 'BHVO.Lu' data frame has 40 rows and 11 columns, which are the measurements from a Lu BHVO-1 basalt aliquot.

Details

Variables:

- Time (POSIXct format)
- 170 Mass
- 172 Mass
- 173 Mass
- 174 Mass
- 175 Mass
- 176 Mass
- 177 Mass
- 185 Mass
- 187 Mass
- Sample ID

4 errorutils

See Also

```
BHVO.Hf,Standard.Hf,Standards.Lu_Re
```

Examples

```
## Not run:
#Preloaded data
data (BHVO.Lu)
head (BHVO.Lu)
## End(Not run)
```

errorutils

A collection of some functions commonly used for setting uncertainties.

Description

General functions for the calculation of uncertainties and outlier detection.

Usage

```
outlier2sigma(data)
SD(data)
RSD(data)
SE(data)
```

Details

Variables:

- SD This function computes the standard deviation.
- **RSD** This function computes the coefficient of variation expressed as a percentage.
- SE This function computes the 2-standard error (2S.E.=1.96 $\sigma/\sqrt{(n)}$). Note: the real range of error is 95% confidence level.
- outlier2sigma If outliers are detected by the 2σ method, this function can remove it with 'NA' values.

```
## Not run:
data <- round(rnorm(80, mean = 0.283100, sd = 0.000018),6)
SD(data)
##
RSD(data)
##
SE(data)
##
outlier2sigma(data)
## End(Not run)</pre>
```

Hf180corr 5

Hf180corr

180Hf unspiked correct data

Description

Computes the corrected ¹⁸⁰Hf/¹⁷⁷Hf ratio and its uncertainty from standard solution or unspiked samples aliquot.

Usage

```
Hf180corr(data,outlier = TRUE)
```

Arguments

data a data file from the output of the function readHf.

outlier logical indicating whether you want outlier data rejection.

Details

This function calculates the mass bias factor (Rusell et al., 1978) and corrects the interferences on ¹⁸⁰Hf (from ¹⁸⁰Ta and ¹⁸⁰W) to compute a corrected ¹⁸⁰Hf/¹⁷⁷Hf ratio from unspiked sample data.

Value

A corrected ¹⁸⁰Hf/¹⁷⁷Hf value and its uncertainties are returned. This function has a graphic output.

References

Russel W.A, Papanastassiou D.A., Tombrello, T.A., 1978, Ca isotope fractionation on the earth and other solar system materials: Geochimica at Cosmochimica Acta, 42,1075-1090.

See Also

```
HfIDcorr, Hfcorr
```

```
## Not run:
#from preloaded data
data (Standard.Hf)
Hf180corr(Standard.Hf)
#############

#from .FIN2 file
##It is important set the work directory #setwd (path)
#getwd()To know the actual directory
data<-readHf()
Hf180corr(data)
## End(Not run)</pre>
```

6 Hfcorr

Hfcorr	Hf unspiked correct data	

Description

Computes the corrected ¹⁷⁶Hf/¹⁷⁷Hf ratio and its uncertainty from standard solution or unspiked sample aliquot.

Usage

```
Hfcorr(data, intLu.corr = FALSE, intYb.corr = FALSE, outlier = TRUE)
```

Arguments

data	a data file from the output of the function readHf.
intLu.corr	logical that carry into effect the interference correction of $^{176}\mathrm{Lu}$ on $^{176}\mathrm{Hf}$ from 175 Mass intensity.
intYb.corr	logical that carry into effect the interference correction of $^{176}{\rm Yb}$ on $^{176}{\rm Hf}$ from 172 Mass intensity.
outlier	logical indicating whether you want outlier values rejection.

Details

This function calculates the mass bias factor (Russel et al., 1978) and can correct the interferences on 176 Hf (from 176 Yb and 176 Lu) to perform corrected 176 Hf/ 177 Hf ratio from unspiked sample data. Additionally, computes isotope ratios measured in Hf aliquot to monitor the isobaric interference on 176 Hf, like 172 Yb/ 176 \sum and 175 Lu/ 176 \sum (Bast et al., 2015). Note: 176 \sum = 176 Yb + 176 Hf.

Value

A corrected 176 Hf/ 177 Hf value and its uncertainties are returned. To prove the data quality, the interference monitors are shown. This function has a graphic output.

References

Russel W.A, Papanastassiou D.A., Tombrello, T.A., 1978, Ca isotope fractionation on the earth and other solar system materials: Geochimica at Cosmochimica Acta, 42,1075-1090.

Bast, R., Scherer, E.E., Sprung, P., Fischer-Gödde, M., Stracke, A., Mezger, K., 2015, A rapid and efficient ion-exchange chromatography for Lu–Hf, Sm–Nd, and Rb–Sr geochronology and the routine isotope analysis of sub-ng amounts of Hf by MC-ICP-MS: Journal of Analytical Atomic Spectrometry, 30, 2323–2333.

See Also

HfIDcorr,Hf180corr

HfIDcorr 7

Examples

```
## Not run:
#from preloaded data
data (Standard.Hf)
Hfcorr(Standard.Hf, outlier = FALSE)
#############

#from .FIN2 file
##It is important set the work directory #setwd (path)
#getwd()To know the actual directory
data<-readHf()
Hfcorr(data, outlier = FALSE)
## End(Not run)</pre>
```

HfIDcorr

Hf spiked correct data

Description

Computes elemental concentration of Hf, corrected ¹⁷⁶Hf/¹⁷⁷Hf ratio and its uncertainty from spiked sample aliquot.

Usage

```
HfIDcorr(data,SampleWeight = NA, SpikeWeight = NA, intLu.corr = FALSE, intYb.corr = FALSE)
```

Arguments

data	a data file from the output of the function readHf.
SampleWeight	the sample weight (g). This value is used to calculate Hf concentration.
SpikeWeight	the spike added (g). This value is used to calculate Hf concentration.
intLu.corr	logical that carry into effect the interference correction of $^{176}\mathrm{Lu}$ on $^{176}\mathrm{Hf}$ from 175 Mass intensity.
intYb.corr	logical that carry into effect the interference correction of $^{176}{\rm Yb}$ on $^{176}{\rm Hf}$ from 172 Mass intensity.
outlier	logical indicating whether you want outlier values rejection.

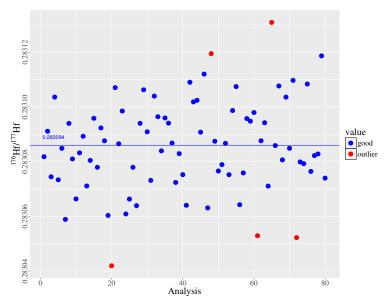
Details

This function calculates the Hf concentration from isotope dilution equations (Gopalan, 2002). It is important to note that values from WR-1 spike (University of Münster) are used by default. This function calculates all necessary corrections for spiked samples (mass bias (Russel et al., 1978), interference and spike (Chu et al., 2011) corrections). Additionally, computes isotope ratios measured in Hf aliquot to monitor the isobaric interference on $^{176} \rm Hf$ like $^{172} \rm Yb/^{176} \sum$ and $^{175} \rm Lu/^{176} \sum$ (Bast et al., 2015). Note: $^{176} \sum = ^{176} \rm Yb + ^{176} \rm Lu + ^{176} \rm Hf$.

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Value

A corrected ¹⁷⁶Hf/¹⁷⁷Hf value and its uncertainty are returned. It is critical for calculating the concentration add the weights of both the sample and the spike into the command. To prove high data quality, the observed interference monitors are shown. This function has a graphic output. See below.



References

Russel W.A, Papanastassiou D.A., Tombrello, T.A., 1978, Ca isotope fractionation on the earth and other solar system materials: Geochimica at Cosmochimica Acta, 42,1075-1090.

Gopalan, K., 2002, Direct correction for mass fractionation of spiked isotopic ratios from dynamic multicollector measurements: Geochemical journal 36(1), 83 - 89.

Chu, Z.Y., Yang, Y.H., Jinghui, G., Qiao, G.S., 2011, Calculation methods for direct internal mass fractionation correction of spiked isotopic ratios from Multi-collector mass spectrometric measurements: International Journal of Mass Spectrometry, 299 (2), 87-93.

Bast, R., Scherer, E.E., Sprung, P., Fischer-Gödde, M., Stracke, A., Mezger, K., 2015, A rapid and efficient ion-exchange chromatography for Lu–Hf, Sm–Nd, and Rb–Sr geochronology and the routine isotope analysis of sub-ng amounts of Hf by MC-ICP-MS: Journal of Analytical Atomic Spectrometry, 30, 2323–2333.

See Also

Hfcorr,Hf180corr

Examples

Not run:
#from preloaded data
data (BHVO.Hf)

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Hftools

Hftools

Description

Computes useful parameters of Lu-Hf pair.

Usage

```
Hfi (ratio, lu = NA, hf = NA, age = NA)
Eps (ratio)
EpsT (ratio, lu = NA, hf = NA, age = NA)
TDm (ratio, lu = NA, hf = NA)
TDm2 (ratio, lu = NA, hf = NA, age = NA)
allHf (ratio, lu = NA, hf = NA, age = NA)
```

Arguments

```
ratio ^{176} \mathrm{Hf/^{177} Hf} ratio. lu Lu concentration (ppm). hf Hf concentration (ppm). age age in Ma.
```

Details

Functions:

- **Hfi** This function computes initial ¹⁷⁶Hf/¹⁷⁷Hf ratios.
- **Eps** This function computes ε Hf values.
- **EpsT** This function computes initial ε Hf values.
- TDm This function computes one-stage model ages.
- TDm2 This function computes two-stage model ages.
- allHf This function runs all the parameters.

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Initial isotope values are calculated at t = age in Ma, using $\lambda^{176}Lu = 1.867 \times 10^{11}$ (Scherer et al. 2001; Scherer et al., 2003; Söderlund et al., 2004). The ε Hfi is calculated following CHUR (CHondrite Uniform Reservoir) values: 176 Hf/ 177 Hf = 0.282785 and 176 Lu/ 177 Hf = 0.0336 (Bouvier et al., 2008). The Depleted Mantle (DM) values used are after Vervoort (2011): 176 Hf/ 177 Hf = 0.28321 and 176 Lu/ 177 Hf = 0.0399. Two-stage crustal residence model ages are calculated assuming (176 Lu/ 177 Hf)avg. crust = 0.015 (Condie et al., 2005).

Value

Numeric: ${}^{176}\text{Hf}/{}^{177}\text{Hf}_i$, εHf , εHf_i , one- and two-stage model ages

References

Scherer, E.E., Munker, C., Mezger, K., 2001, Calibration of the Lutetium-Hafnium Clock: Science 293, 683–687.

Scherer, E.E., Mezger, K., Münker, C., 2003, The ¹⁷⁶Lu decay constant discrepancy: terrestrial samples vs. meteorites: Meteoritics and Planetary Science Suppl 38, 5263.

Söderlund, U., Patchett, P.J., Vervoort, J.D., Isachsen, C.E., 2004, The ¹⁷⁶Lu decay constant determined by Lu-Hf and U-Pb isotope systematics of Precambrian mafic intrusions: Earth and Planetary Science Letters 219, 311–324.

Condie, K.C., Beyer, E., Belousova, E., Griffin, W.L., O'Reilly, S.Y., 2005, U–Pb isotopic ages and Hf isotopic composition of single zircons: The search for juvenile Precambrian continental crust: Precambrian Research 139, 42-100.

Bouvier, A., Vervoort, J.D., and Patchett, P.J., 2008, The Lu-Hf and Sm-Nd isotopic composition of CHUR: Constraints from unequilibrated chondrites and implications for the bulk composition of terrestrial planets, Earth and Planetary Science Letters, 273, 48-57.

Vervoort, J.D., Plank, T., Prytulak, J., 2011, The Hf-Nd isotopic composition of marine sediments: Geochimica at Cosmochimica Acta, 75, 5903-5926.

```
## Not run:
sample<-c("03-1b-1","unknown","03-2a-1","03-2a-2","zircon-1","zircon-2")
ratio<-c(0.282261,0.282432,0.282254,0.282261,0.282191,0.282200)
lu<-c(0.62,0.52,0.59,0.53,26.8,46.7)
hf<-c(25.64,19.90,28.72,14.79,6990,9780)
age<-rep(1000,6)
table<-allHf(ratio,lu,hf,age)
rownames(table) <- sample
print(table)
## End(Not run)</pre>
```

LuIDcorr 11

LuIDcorr	Lu spiked correct data
	_

Description

Computes elemental concentration of Lu as well as 175 Lus/ 176 Lun ratio and its uncertainties from spiked sample aliquot.

Usage

```
LuIDcorr(data,SampleWeight = NA, SpikeWeight = NA, intHf.corr = FALSE, Slope=NA, outlier = TRUE)
```

Arguments

data	a data file from the output of the function readLu.
SampleWeight	the sample weight (g). This value is used to calculate Lu concentration.
SpikeWeight	the spike added (g). This value is used to calculate Lu concentration.
intLu.corr	carry into effect the interference correction of $^{176}{\rm Hf}$ on $^{176}{\rm Lu}$ from the measured intensity of $^{177}{\rm Hf}$.
Slope	the slope value of the log-log plot of the standard (Lu-Re) in a given work session. See SlopeReLu.
outlier	a logical indicating whether you want outlier data rejection.

Details

This function computes the Re-Lu doping method (Scherer et al., 1999) in order to return Lu data. It is important to note that values from WR-1 spike (University of Münster) are used by default.

Value

A corrected ¹⁷⁵Lus/¹⁷⁶Lun value and its uncertainty are returned. It is critical for calculating the concentration add the weights of the sample and the spike as well as the slope value (log-log plot Re-Lu) into the command. The mass-bias factors of Yb, Lu, and Re are also returned. This function has a graphic output.

References

Scherer, E, Munker, C, Rehkamper, M, Mezger, K., 1999, Improved precision of Lu isotope dilution measurements by MC-ICP-MS and application to Lu-Hf geochronology: Eos Suppl 80(46), 1118.

See Also

Hfcorr,HfIDcorr,SlopeReLu

Pearson Pearson

Examples

```
## Not run:
#from preloaded data
data (BHVO.Lu)
LuIDcorr(BHVO.Lu, outlier = FALSE, SampleWeight = 0.1198, SpikeWeight =0.0334, S = 0.67202)
################

#from .FIN2 file
##It is important set the work directory #setwd (path)
#getwd()To know the actual directory
data<-readLu()
LuIDcorr(data, outlier = TRUE, SampleWeight = 0.1198, SpikeWeight =0.0334, S = 0.67202)
## End(Not run)</pre>
```

Pearson

Pearson

Description

The 'Pearson' data frame has 10 rows and 4 columns, which are Pearsson's x-y data used in York's function example.

Details

Variables:

- x (x-value)
- sx (absolute error)
- y (y-value)
- sy (absolute error)

References

Pearson, K., 1901, On Lines and planes of closest fit to systems of points in space: Philosophical Magazzine, 2(6), 559-572.

See Also

York

```
## Not run:
#Preloaded data
data (Pearson)
## End(Not run)
```

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readHf

Read file Hf.FIN2

Description

Reads the selected file with .FIN2 extension.

Usage

readHf(data)

Arguments

data

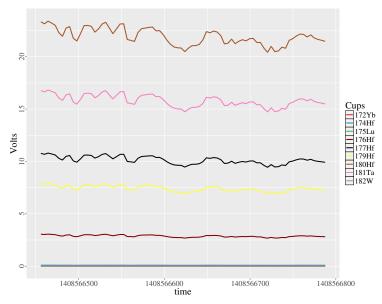
path to the .FIN2 file (.FIN2 is the output extension from Thermo Neptune® MC-ICP-MS, same as .csv extension).

Details

This function reads the indicated file. The argument must to be the complete path to a file. If it does not contain an absolute path, the file name is relative to the current working directory (getwd()). This function creates a data frame with the measured voltages of the Mass spectrometer to determine Hf concentrations. The time is changed to POSIXct format.

Value

If a .FIN2 file is selected, then a data frame is returned. This function has a graphic output, see below.



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See Also

readLu

Examples

```
## Not run:
#It is important set the work directory
#getwd()#To know the actual directory
#setwd(path)#To change directory
data<-readHf()
## End(Not run)</pre>
```

readLu

Read file Lu.FIN2

Description

Reads the selected file(s) with .FIN2 extension.

Usage

readLu(data)

Arguments

data

path to the .FIN2 file (.FIN2 is the output extension from Thermo Neptune® MC-ICP-MS, same as .csv extension).

Details

This function read(s) the indicated file(s). The argument must to be the complete path to a file(s). If it does not contain an absolute path, the file name is relative to the current working directory (getwd()). This function creates a data frame with the measured voltages of the Mass spectrometer to determine Lu concentrations. The time is changed to POSIXct format.

Value

If a .FIN2 file is selected, then a data frame is returned. This function has a graphic output.

See Also

readHf

SlopeReLu 15

Examples

```
## Not run:
#It is important set the work directory
#setwd(path)#To change directory
#getwd()#To know the actual directory
data<-readLu(getwd())
## End(Not run)</pre>
```

SlopeReLu

Slope of Re-Lu (log-log) plot

Description

Computes the slope, intercept and other parameters from a linear array in a log-log diagram.

Usage

SlopeReLu(data)

Arguments

data

path to the .FIN2 files (.FIN2 is the output extension from Thermo Neptune® MC-ICP-MS, same as .csv extension).

Details

Computes a weighted regression following the approach of York et al. (2004) (see York). It is calculated from a standard solution containing both Re and Lu, which is measured interspersed with unknown samples in a given analytical session. All files of these measurements must be in the same folder.

Value

It is returned the regression coefficients from the $\ln(^{187}\text{Re}/^{185}\text{Re})$ vs. $\ln(^{176}\text{Lu}/^{175}\text{Lu})$ diagram and its uncertainties. This function has a graphic output.

References

Scherer, E, Munker, C, Rehkamper, M, Mezger, K., 1999, Improved precision of lu isotope dilution measurements by MC-ICP-MS and application to Lu-Hf geochronology: Eos Suppl 80(46), 1118.

York, D., Evensen, N.M., López-Martínez, M., De Basabe-Delgado, J., 2004, Unified equations for the slope, intercept, and standard errors of the best straight line: American Journal of Physics, 72, 367-375.

See Also

LuIDcorr, York

Standard.Hf

Examples

```
## Not run:
#It is important set the work directory
#setwd(path)#To change directory
#getwd()#To know the actual directory
SlopeReLu()
## End(Not run)
```

Standard.Hf

Standard.Hf

Description

The 'Standard.Hf' data frame has 80 rows and 11 columns, which are the measurements from a JMC 475 standard solution.

Details

Variables:

- Time (POSIXct format)
- 172 Mass
- 174 Mass
- 175 Mass
- 176 Mass
- 177 Mass
- 179 Mass
- 180 Mass
- 181 Mass
- 182 Mass
- Sample ID

See Also

BHVO.Lu,BHVO.Hf

```
## Not run:
#Preloaded data
data (Standard.Hf)
head (Standard.Hf)
## End(Not run)
```

Standards.Lu_Re 17

Standards.Lu_Re

Standards.Lu_Re

Description

The 'Standards.Lu.Re' data frame has 320 rows and 11 columns, which are the measurements from a standard solution containing both Re and Lu, which is measured interspersed with unknown samples in an analytical session.

Details

Variables:

- **Time** (POSIXct format)
- 170 Mass
- 172 Mass
- 173 Mass
- 174 Mass
- 175 Mass
- 176 Mass
- 177 Mass
- 185 Mass
- 187 Mass
- Sample ID

See Also

BHVO.Lu, SlopeReLu, LuIDcorr

```
## Not run:
#Preloaded data
data (Standards.Lu.Re)
head (Standards.Lu.Re)
## End(Not run)
```

18 York

Description

Computes the linear weighted regression (with correlated errors in bivariate data).

Usage

York (data)

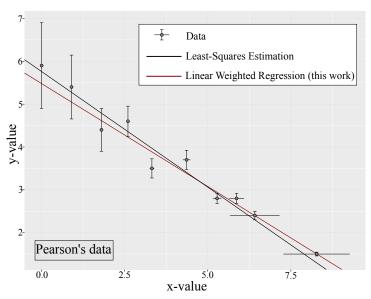
Arguments

data

data frame with x,y,sx, and sy values (sx and sy must be absolute errors).

Details

This function generates a best-fit line using the weighted regression method following the approach of York et al. (2004). In some analytical applications, it may be required. In this case of Pearson's data (Pearson, 1901), the results are compared with other softwares (see Table below). A graphical output displaying the fitted trend line with the data points is also shown for comparison.



Tool	Intercept	Std. error	Slope	Std. error	r^2
1 LSM	5.47991	0.295713	-0.485033	0.058256	NA
² Excel®	5.47991	0.359247	-0.4805333	0.070620	NA
3 Matlab ${\mathbb R}$	5.47991	0.294971	-0.4805334	0.057985	0.96
This function	5.47991	0.294971	-0.4805336	0.057985	0.96
R (LSE)	5.76119	0.18949	-0.53958	0.04213	0.95

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Value

The output displays the slope, the intercept, and their uncertainties as well as the linear correlation coefficient of the fitted data.

References

Pearson, K., 1901, On Lines and planes of closest fit to systems of points in space: Philosophical Magazzine, 2(6), 559-572.

¹York, D., Evensen, N.M., López-Martínez, M., De Basabe-Delgado, J., 2004, Unified equations for the slope, intercept, and standard errors of the best straight line: American Journal of Physics, 72, 367-375.

²Cantrell, C.A., 2008, Technical Note: Review of methods for linear least-squares fitting of data and application to atmospheric chemistry problems. Atmospheric Chemistry and Physics Discussions, 8, 6409-6436.

³ Thirumalai, K., Singh, A., Ramesh, R., 2011, A MATLAB code to perform weighted linear regression with (correlated or uncorrelated) errors in bivariate data: Journal of The Geological Society of India, 77(4), 377-380.

See Also

Pearson, SlopeReLu, LuIDcorr

```
## Not run:
#from preloaded data
data (Pearson)
York(Pearson)
##############

##-- Pearson's data --##
x <- c(0.0, 0.9, 1.8, 2.6, 3.3, 4.4, 5.2, 6.1, 6.5, 7.4)
y <- c(5.9, 5.4, 4.4, 4.6, 3.5, 3.7, 2.8, 2.8, 2.4, 1.5)
sx <- c(0.031623, 0.031623, 0.044721, 0.035355, 0.070711, 0.111803, 0.129099, 0.223607, 0.745356, 1.000000)
sy <- c(1.000000, 0.745356, 0.500000, 0.353553, 0.223607, 0.223607, 0.119523, 0.119523, 0.100000, 0.044721)
data <- data.frame(x,y,sx,sy)
York(data)
## End(Not run)</pre>
```

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