

Package ‘IsoplotR’

June 22, 2016

Title Statistical Toolbox for Radiometric Geochronology

Version 0.4

Description An R implementation of Ken Ludwig's popular Isoplot add-in to Microsoft Excel. Currently plots U-Pb data on Wetherill and Tera-Wasserburg concordia diagrams. Calculates concordia and discordia ages. Performs linear regression of measurements with correlated errors using the 'York' approach. Generates Kernel Density Estimates (KDEs) and Cumulative Age Distributions (CADs). Future versions will include functionality for the Ar-Ar, Rb-Sr, Sm-Nd, Re-Os, U-Th-He, fission track and cosmogenic nuclide methods, including isochrons, age spectra, ternary diagrams, radial plots, banana diagrams and multidimensional scaling plots. A graphical user interface is provided as an RStudio Shiny app at <http://isoplotr.london-geochron.com>. Offline access to this interface will be provided at a later point in time.

Author Pieter Vermeesch [aut, cre]

Maintainer Pieter Vermeesch <p.vermeesch@ucl.ac.uk>

Depends R (>= 3.0.0)

Imports methods

License GPL-2

LazyData true

RoxygenNote 5.0.1

R topics documented:

age	2
botev	3
cad	4
concordia	5
ellipse	6
examples	7
iratio	7
kde	8
lambda	11
read.data	11
settings	12
yorkfit	13

age	<i>Calculate isotopic ages</i>
-----	--------------------------------

Description

Calculates U-Pb ages and propagates their analytical uncertainties. Evaluates the equivalence of multiple ($^{206}\text{Pb}/^{238}\text{U}$ - $^{207}\text{Pb}/^{235}\text{U}$ or $^{207}\text{Pb}/^{206}\text{Pb}$ - $^{206}\text{Pb}/^{238}\text{U}$) compositions, computes the weighted mean isotopic composition and the corresponding concordia age using the method of maximum likelihood, computes the mswd of equivalence and concordance and their respective Chi-squared p-values. Performs linear regression of U-Pb data on Wetherill and Tera-Wasserburg concordia diagrams. Computes the upper and lower intercept ages (for Wetherill) or the lower intercept age and the $^{207}\text{Pb}/^{206}\text{Pb}$ intercept (for Tera-Wasserburg), taking into account error correlations and decay constant uncertainties.

Usage

```
age(x, ...)
```

Default S3 method:

```
age(x, method = "Pb206U238", dcu = TRUE, ...)
```

S3 method for class 'UPb'

```
age(x, concordia = 1, wetherill = TRUE, dcu = TRUE,
    i = NA, ...)
```

S3 method for class 'detritals'

```
age(x, ...)
```

Arguments

x	a scalar containing an isotopic ratio, a two element vector containing an isotopic ratio and its standard error, or an object of class UPb or detritals.
...	optional arguments
method	one of either 'Pb206U238', 'Pb207U235', or 'Pb207Pb206'
dcu	propagate the decay constant uncertainties?
concordia	scalar flag indicating whether each U-Pb analysis should be considered separately (concordia=1), a concordia age should be calculated from all U-Pb analyses together (concordia=2), or a discordia line should be fit through all the U-Pb analyses (concordia=2).
wetherill	boolean flag to indicate whether the data should be evaluated in Wetherill (TRUE) or Tera-Wasserburg (FALSE) space. This option is only used when concordia=2
i	(optional) index of a particular aliquot

Value

if **x** is a scalar or a vector, returns the age using the geochronometer given by **method** and its standard error.

if **x** has class **UPb** and **concordia=1**, returns a table with the following columns: 't.75', 'err[t.75]', 't.68', 'err[t.68]', 't.76', 'err[t.76]', 't.conc', 'err[t.conc]', containing the $^{207}\text{Pb}/^{235}\text{U}$ -age and standard error, the $^{206}\text{Pb}/^{238}\text{U}$ -age and standard error, the $^{207}\text{Pb}/^{206}\text{Pb}$ -age and standard error, and the concordia age and standard error, respectively.

if **x** has class **UPb** and **concordia=2**, returns a list with the following items:

x a named vector with the (weighted mean) U-Pb composition

cov the covariance matrix of the (mean) U-Pb composition

age the concordia age (in Ma)

age.err the standard error of the concordia age

mswd a list with two items (equivalence and concordance) containing the MSWD (Mean of the Squared Weighted Deviates, a.k.a the reduced Chi-squared statistic outside of geochronology) of isotopic equivalence and age concordance, respectively.

p.value a list with two items (equivalence and concordance) containing the p-value of the Chi-square test for isotopic equivalence and age concordance, respectively.

if **x** has class **UPb** and **concordia=3**, returns a list with the following items:

x a two element vector with the upper and lower intercept ages (if **wetherill==TRUE**) or the lower intercept age and $^{207}\text{Pb}/^{206}\text{Pb}$ intercept (for Tera-Wasserburg)

cov the covariance matrix of the elements in **x**

Examples

```
data(examples)
print(age(examples$UPb))
print(age(examples$UPb, concordia=1))
print(age(examples$UPb, concordia=2))
```

botev

Compute the optimal kernel bandwidth

Description

Uses the diffusion algorithm of Zdravko Botev (2011) to calculate the bandwidth for kernel density estimation

Usage

```
botev(x)
```

Arguments

x a vector of ordinal data

Value

a scalar value with the optimal bandwidth

Author(s)

Dzdravko Botev

References

Botev, Z. I., J. F. Grotowski, and D. P. Kroese. "Kernel density estimation via diffusion." The Annals of Statistics 38.5 (2010): 2916-2957.

Examples

```
data(examples)
samp <- examples$DZ[['N1']]
bw <- botev(samp)
print(bw)
```

cad	<i>Plot continuous data as cumulative age distributions</i>
-----	---

Description

Plot a dataset as a Cumulative Age Distribution (CAD), also known as a ‘empirical cumulative distribution function’.

Usage

```
cad(x, method = NA, pch = NA, verticals = TRUE, xlab = "age [Ma]",
    colmap = "heat.colors", col = "black", ...)
```

Arguments

x	an object of class UPb or detritals
method	a string indicating what kind of age should be plotted. If x has class UPb, type could be one of either t.75, t.68 (default), t.76 or t.conc
pch	(optional) plot character
verticals	boolean flag indicating if the horizontal lines of the CAD should be connected by vertical lines
xlab	x-axis label

colmap	an optional string with the name of one of R's built-in colour palettes (e.g., heat.colors, terrain.colors, topo.colors, cm.colors), which are to be used for plotting data of class detritals.
col	colour to give to single sample datasets (i.e. not of class detritals)
...	optional arguments to the generic plot function

Examples

```
data(examples)
cad(examples$DZ)
```

concordia	<i>Concordia diagram</i>
-----------	--------------------------

Description

Plot U-Pb data on Wetherill and Tera-Wasserburg concordia diagrams, calculate concordia ages and compositions, evaluates the equivalence of multiple ($^{206}\text{Pb}/^{238}\text{U}$ - $^{207}\text{Pb}/^{235}\text{U}$ or $^{207}\text{Pb}/^{206}\text{Pb}$ - $^{206}\text{Pb}/^{238}\text{U}$) compositions, computes the weighted mean isotopic composition and the corresponding concordia age using the method of maximum likelihood, computes the msd of equivalence and concordance and their respective Chi-squared p-values. Performs linear regression and computes the upper and lower intercept ages (for Wetherill) or the lower intercept age and the $^{207}\text{Pb}/^{206}\text{Pb}$ intercept (for Tera-Wasserburg), taking into account error correlations and decay constant uncertainties.

Usage

```
concordia(x, limits = NULL, alpha = 0.05, wetherill = TRUE,
  show.numbers = FALSE, ellipse.col = rgb(0, 1, 0, 0.5),
  concordia.col = "darksalmon", dcu = TRUE, show.age = 1)
```

Arguments

x	an object of class UPb
limits	age limits of the concordia line
alpha	confidence cutoff for the error ellipses
wetherill	boolean flag (FALSE for Tera-Wasserburg)
show.numbers	boolean flag (TRUE to show grain numbers)
ellipse.col	background colour of the error ellipses
concordia.col	colour of the concordia line
dcu	show decay constant uncertainty?
show.age	one of either 1: don't show the age 2: calculate the concordia age 3: fit a discordia line

Examples

```
data(examples)
concordia(examples$UPb)
```

 ellipse

Get coordinates of error ellipse for plotting

Description

Construct an error ellipse age a given confidence level from its centre and covariance matrix

Usage

```
ellipse(x, y, covmat, alpha = 0.05)
```

Arguments

x	x-coordinate (scalar) for the centre of the ellipse
y	y-coordinate (scalar) for the centre of the ellipse
covmat	covariance matrix of the x-y coordinates
alpha	the probability cutoff for the error ellipses

Value

a [50x2] matrix of plot coordinates

Examples

```
x = 99; y = 101;
covmat <- matrix(c(1,0.9,0.9,1),nrow=2)
ell <- ellipse(x,y,covmat)
plot(c(90,110),c(90,110),type='l')
polygon(ell,col=rgb(0,1,0,0.5))
points(x,y,pch=21,bg='black')
```

examples

*Example datasets for testing IsoplotR***Description**

U-Pb and detrital zircon datasets

Details

examples is a list with two items

UPb: an object of class 'UPb' containing a high precision U-Pb dataset packaged with Ken Ludwig's Isoplot program.

DZ: an object of class 'detrital' containing a detrital zircon U-Pb dataset from Namibia.

Author(s)

Ken Ludwig and Pieter Vermeesch

References

Ludwig, K. R. User's manual for Isoplot 3.00: a geochronological toolkit for Microsoft Excel. No. 4. Kenneth R. Ludwig, 2003.

Vermeesch, Pieter, and Eduardo Garzanti. "Making geological sense of 'Big Data' in sedimentary provenance analysis." Chemical Geology 409 (2015): 20-27.

Examples

```
data(examples)
concordia(examples$UPb)
dev.new()
kde(examples$DZ)
```

iratio

*Isotopic ratios***Description**

Gets or sets natural isotopic ratios.

Usage

```
iratio(ratio, x = NULL, e = NULL)
```

Arguments

ratio	one of either 'U238U235', 'Ar40Ar36', 'Ar38Ar36', 'Rb85Rb87', 'Sr88Sr86', 'Sr87Sr86', 'Sr84Sr86', 'Re185Re187', 'Os184Os192', 'Os186Os192', 'Os187Os192', 'Os188Os192', 'Os189Os192'
x	new value for ratio
e	new value for its standard error

Value

if x=e=NULL, returns a two-item vector containing the mean value of the requested ratio and its standard error, respectively.

References

- Ar: Lee, Jee-Yon, et al. "A redetermination of the isotopic abundances of atmospheric Ar." *Geochimica et Cosmochimica Acta* 70.17 (2006): 4507-4512.
- Rb: Catanzaro, E. J., et al. "Absolute isotopic abundance ratio and atomic weight of terrestrial rubidium." *J. Res. Natl. Bur. Stand. A* 73 (1969): 511-516.
- Sr: Moore, L. J., et al. "Absolute isotopic abundance ratios and atomic weight of a reference sample of strontium." *J. Res. Natl. Bur. Stand.* 87.1 (1982): 1-8.
- Re: Gramlich, John W., et al. "Absolute isotopic abundance ratio and atomic weight of a reference sample of rhenium." *J. Res. Natl. Bur. Stand. A* 77 (1973): 691-698.
- Os: Voelkening, Joachim, Thomas Walczyk, and Klaus G. Heumann. "Osmium isotope ratio determinations by negative thermal ionization mass spectrometry." *Int. J. Mass Spect. Ion Proc.* 105.2 (1991): 147-159.
- U: Hiess, Joe, et al. "²³⁸U/²³⁵U systematics in terrestrial uranium-bearing minerals." *Science* 335.6076 (2012): 1610-1614.

Examples

```
# returns the 238U/235U ratio of Hiess et al. (2012):
print(iratio('U238U235'))
# use the 238U/235U ratio of Steiger and Jaeger (1977):
iratio('U238U235',138.88,0)
print(iratio('U238U235'))
```

kde

Create (a) kernel density estimate(s)

Description

Creates one or more kernel density estimates using a combination of the Botev (2010) bandwidth selector and the Abramson (1982) adaptive kernel bandwidth modifier.

Usage

```

kde(x, ...)

## Default S3 method:
kde(x, from = NA, to = NA, bw = NA, adaptive = TRUE,
    log = FALSE, n = 512, plot = TRUE, pch = NA, xlab = "age [Ma]",
    ylab = "", kde.col = rgb(1, 0, 1, 0.6), hist.col = rgb(0, 1, 0, 0.2),
    show.hist = TRUE, bty = "n", binwidth = NA, ncol = NA, ...)

## S3 method for class 'UPb'
kde(x, from = NA, to = NA, bw = NA, adaptive = TRUE,
    log = FALSE, n = 512, plot = TRUE, pch = NA, xlab = "age [Ma]",
    ylab = "", kde.col = rgb(1, 0, 1, 0.6), hist.col = rgb(0, 1, 0, 0.2),
    show.hist = TRUE, bty = "n", binwidth = NA, ncol = NA, type = 4,
    cutoff.76 = 1100, cutoff.disc = c(-15, 5), ...)

## S3 method for class 'detritals'
kde(x, from = NA, to = NA, bw = NA, adaptive = TRUE,
    log = FALSE, n = 512, plot = TRUE, pch = NA, xlab = "age [Ma]",
    ylab = "", kde.col = rgb(1, 0, 1, 0.6), hist.col = rgb(0, 1, 0, 0.2),
    show.hist = TRUE, bty = "n", binwidth = NA, ncol = NA,
    samebandwidth = TRUE, normalise = TRUE, ...)

```

Arguments

<code>x</code>	a vector of numbers or an object of class UPb or detrital
<code>...</code>	optional arguments to be passed on to density
<code>from</code>	minimum age of the time axis. If NULL, this is set automatically
<code>to</code>	maximum age of the time axis. If NULL, this is set automatically
<code>bw</code>	the bandwidth of the KDE. If NULL, bw will be calculated automatically using <code>botev()</code>
<code>adaptive</code>	boolean flag controlling if the adaptive KDE modifier of Abramson (1982) is used
<code>log</code>	transform the ages to a log scale if TRUE
<code>n</code>	horizontal resolution of the density estimate
<code>plot</code>	show the KDE as a plot
<code>pch</code>	the symbol used to show the samples. May be a vector. Set <code>pch = NA</code> to turn them off.
<code>xlab</code>	the label of the x-axis
<code>ylab</code>	the label of the y-axis
<code>kde.col</code>	the fill colour of the KDE specified as a four element vector of r, g, b, alpha values
<code>hist.col</code>	the fill colour of the histogram specified as a four element vector of r, g, b, alpha values

<code>show.hist</code>	boolean flag indicating whether a histogram should be added to the KDE
<code>bty</code>	change to "o", "l", "7", "c", "u", or "]" if you want to draw a box around the plot
<code>binwidth</code>	scalar width of the histogram bins, in Myr if <code>x\$log==FALSE</code> , or as a fractional value if <code>x\$log==TRUE</code> . Sturges' Rule is used if <code>binwidth==NA</code>
<code>ncol</code>	scalar value indicating the number of columns over which the KDEs should be divided. This option is only used if <code>x</code> is of class <code>detritals</code> .
<code>type</code>	scalar indicating whether to plot the $^{207}\text{Pb}/^{235}\text{U}$ age (<code>type=1</code>), the $^{206}\text{Pb}/^{238}\text{U}$ age (<code>type=2</code>), the $^{207}\text{Pb}/^{206}\text{Pb}$ age (<code>type=3</code>), the $^{207}\text{Pb}/^{206}\text{Pb}$ - $^{206}\text{Pb}/^{238}\text{U}$ age (<code>type=4</code>), or the (Wetherill) concordia age (<code>type=5</code>)
<code>cutoff.76</code>	the age (in Ma) below which the $^{206}\text{Pb}/^{238}\text{U}$ and above which the $^{207}\text{Pb}/^{206}\text{Pb}$ age is used. This parameter is only used if <code>type=4</code> .
<code>cutoff.disc</code>	two element vector with the maximum and minimum percentage discordance allowed between the $^{207}\text{Pb}/^{235}\text{U}$ and $^{206}\text{Pb}/^{238}\text{U}$ age (if $^{206}\text{Pb}/^{238}\text{U} < \text{cutoff.76}$) or between the $^{206}\text{Pb}/^{238}\text{U}$ and $^{207}\text{Pb}/^{206}\text{Pb}$ age (if $^{206}\text{Pb}/^{238}\text{U} > \text{cutoff.76}$). Set <code>cutoff.disc=NA</code> if you do not want to use this filter.
<code>samebandwidth</code>	boolean flag indicating whether the same bandwidth should be used for all samples. If <code>samebandwidth = TRUE</code> and <code>bw = NULL</code> , then the function will use the median bandwidth of all the samples.
<code>normalise</code>	boolean flag indicating whether or not the KDEs should all integrate to the same value.

Value

if `plot==TRUE`, returns an object of class `KDE`, i.e. a list containing the following items:

x horizontal plot coordinates
y vertical plot coordinates
bw the base bandwidth of the density estimate
ages the data values from the input to the KDE function

or, if `class(x)=='detritals'`, an object of class `KDEs`, i.e. a list containing the following items:

kdes a named list with objects of class `KDE`
from the beginning of the common time scale
to the end of the common time scale
themax the maximum probability density of all the KDEs
xlabel the x-axis label to be used by `plot.KDEs`

Examples

```
data(examples)
kde(examples$DZ[['N1']],kernel="epanechnikov")
kde(examples$DZ,from=0,to=3000)
```

lambda	<i>Decay constants</i>
--------	------------------------

Description

Gets or sets the decay constants of radioactive istopes

Usage

```
lambda(nuclide, x = NULL, e = NULL)
```

Arguments

nuclide	the nuclide name
x	new value for the decay constant
e	new value for the decay constant uncertainty

Value

if x==e==NULL, returns a two-item vector containing the decay constant [in Ma-1] and its standard error, respectively.

Examples

```
print(lambda('U238'))
# use the decay constant of Kovarik and Adams (1932)
lambda('U238',0.0001537,0.0000068)
print(lambda('U238'))
```

read.data	<i>Read geochronology data</i>
-----------	--------------------------------

Description

Cast a .csv file or a matrix into one of IsoplotR's data classes

Usage

```
read.data(x, ...)
```

Default S3 method:

```
read.data(x, method = "Pb206U238", format = 1, ...)
```

S3 method for class 'matrix'

```
read.data(x, method = "U-Pb", format = 1, ...)
```

S3 method for class 'data.frame'

```
read.data(x, method = "U-Pb", format = 1, ...)
```

Arguments

x	a file name (.csv format) or matrix
...	optional arguments to the read.csv function
method	one of 'U-Pb', 'Ar-Ar', 'Rb-Sr', 'Sm-Nd', 'Re-Os', 'U-Th-He', 'fission tracks', 'cosmogenic nuclides' or 'other'
format	formatting option, depends on the value of method. If method = 'U-Pb', then format is one of either: 1: 7/6, s[7/6], 6/8, s[6/8], 7/5, s[7/5] (other formats will be added later)

Value

an object of class 'UPb', 'ArAr', 'RbSr', 'SmNd', 'ReOs', 'UThHe', 'fission', 'cosmogenics', or 'other'

Examples

```
# load one of the built-in .csv files:
fname <- system.file("UPb.csv",package="IsoplotR")
UPb <- read.data(fname, 'U-Pb')
concordia(UPb)
```

settings	<i>Load settings to and from json</i>
----------	---------------------------------------

Description

Get and set preferred values for decay constants and isotopic abundances from and to a .json file format

Usage

```
settings(fname = NULL)
```

Arguments

fname	the path of a .json file
-------	--------------------------

Value

if fname==NULL, returns a .json string

Examples

```
json <- system.file("defaults.json",package="IsoplotR")
settings(json)
print(settings())
```

yorkfit

*Linear regression of X,Y-variables with correlated errors***Description**

Implements the unified regression algorithm of York et al. (2004) which, although based on least squares, yields results that are consistent with maximum likelihood estimates of Ludwig and Titterton (1994)

Usage

```
yorkfit(X, Y, sX, sY, rXY)
```

Arguments

X	vector of measurements
Y	vector of measurements
sX	standard errors of X
sY	standard errors of Y
rXY	correlation coefficients between X and Y

Value

a five element list containing

- a** the intercept of the straight line fit
- b** the slope of the fit
- sa** the standard error of the intercept
- sb** the standard error of the slope

References

Ludwig, K. R., and D. M. Titterton. "Calculation of $^{230}\text{Th}/^{238}\text{U}$ isochrons, ages, and errors." *Geochimica et Cosmochimica Acta* 58.22 (1994): 5031-5042.

York, Derek, et al. "Unified equations for the slope, intercept, and standard errors of the best straight line." *American Journal of Physics* 72.3 (2004): 367-375.

Examples

```
X <- c(1.550, 12.395, 20.445, 20.435, 20.610, 24.900,
      28.530, 50.540, 51.595, 86.51, 106.40, 157.35)
Y <- c(.7268, .7849, .8200, .8156, .8160, .8322,
      .8642, .9584, .9617, 1.135, 1.230, 1.490)
n <- length(X)
sX <- X*0.01
sY <- Y*0.005
```

```
rXY <- rep(0.8,n)
fit <- yorkfit(X,Y,sX,sY,rXY)
covmat <- matrix(0,2,2)
plot(range(X),fit$a+fit$b*range(X),type='l',ylim=range(Y))
for (i in 1:n){
  covmat[1,1] <- sX[i]^2
  covmat[2,2] <- sY[i]^2
  covmat[1,2] <- rXY[i]*sX[i]*sY[i]
  covmat[2,1] <- covmat[1,2]
  ell <- ellipse(X[i],Y[i],covmat,alpha=0.05)
  polygon(ell)
}
```

Index

age, [2](#)

botev, [3](#)

cad, [4](#)
concordia, [5](#)

ellipse, [6](#)
examples, [7](#)

iratio, [7](#)

kde, [8](#)

lambda, [11](#)

read.data, [11](#)

settings, [12](#)

yorkfit, [13](#)