

# Package ‘rcarbon’

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**Title** Calibration and Analysis of Radiocarbon Dates

**Version** 1.3.3

**Description** Enables the calibration and analysis of radiocarbon dates, often but not exclusively for the purposes of archaeological research. It includes functions not only for basic calibration, uncalibration, and plotting of one or more dates, but also a statistical framework for building demographic and related longitudinal inferences from aggregate radiocarbon date lists, including: Monte-Carlo simulation test (Timpson et al 2014 <doi:10.1016/j.jas.2014.08.011>), random mark permutation test (Crema et al 2016 <doi:10.1371/journal.pone.0154809>) and spatial permutation tests (Crema, Bevan, and Shennan 2017 <doi:10.1016/j.jas.2017.09.007>).

**Depends** R (>= 3.3.0)

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---

as.CalDates	<i>Convert to a CalDates object</i>
-------------	-------------------------------------

---

## Description

Convert other calibrated date formats to an rcarbon CalDates object.

## Usage

```
as.CalDates(x)
```

## Arguments

x	One or more calibrated dated to convert (currently only BchronCalibratedDates and oxCAARCalibratedDatesList objects are supported)
---	--

## Value

A CalDates object

## Examples

```
## Not run:
library(Bchron)
library(oxCAAR)
quickSetupOxcal()
dates <- data.frame(CRA=c(3200,2100,1900), Error=c(35,40,50))
bcaldates <- BchronCalibrate(ages=dates$CRA, ageSds=dates$Error,
calCurves=rep("intcal13", nrow(dates)))
rcaldates <- rcarbon::calibrate(dates$CRA, dates$Error, calCurves=rep("intcal13"))
ocaldates <- oxcalCalibrate(c(3200,2100,1900),c(35,40,50),c("a","b","c"))
## Convert to rcarbon format
caldates.b <- as.CalDates(bcaldates)
caldates.o <- as.CalDates(ocaldates)
## Comparison plot
plot(rcaldates$grids[[2]]$calBP,rcaldates$grids[[2]]$PrDens,
type="l", col="green", xlim=c(2300,1900))
lines(caldates.b$grids[[2]]$calBP,caldates.b$grids[[2]]$PrDens, col="red")
lines(caldates.o$grids[[2]]$calBP,caldates.o$grids[[2]]$PrDens, col="blue")
legend("topright", legend=c("rcarbon","Bchron","OxCal"), col=c("green","red","blue"), lwd=2)
```

```
## End(Not run)
```

---

```
as.CalGrid          Convert data to class CalGrid.
```

---

### Description

Tries to coerce any two-column matrix or data.frame to a calibrated probability distribution (an object of class "CalGrid") for use by the rcarbon package.

### Usage

```
as.CalGrid(x)
```

### Arguments

x                    A two-column matrix or data.frame class object.

### Value

A CalGrid class object of probabilities or summed probabilities per calendar year BP.

### Examples

```
df <- data.frame(calBP=5000:2000,PrDens=runif(length(5000:2000)))
mycalgrid <- as.CalGrid(df)
plot(mycalgrid)
```

---

```
as.UncalGrid        Convert data to class UncalGrid.
```

---

### Description

Tries to coerce any two-column matrix or data.frame to a uncalibrated probability distribution (an object of class "UncalGrid") for use by the rcarbon package.

### Usage

```
as.UncalGrid(x)
```

### Arguments

x                    A two-column matrix or data.frame class object.

### Value

A CalGrid class object of probabilities or summed probabilities per CRA.

**Examples**

```
df <- data.frame(CRA=5000:2000,PrDens=runif(length(5000:2000)))
mycalgrid <- as.UncalGrid(df)
```

---

barCodes

*Plot the median values of calibrated radiocarbon dates or bins*


---

**Description**

Plot the median values of multiple calibrated radiocarbon dates or bins in a barcode-like strip.

**Usage**

```
barCodes(
  x,
  yrng = c(0, 0.03),
  width = 20,
  col = rgb(0, 0, 0, 25, maxColorValue = 255),
  border = NA,
  ...
)
```

**Arguments**

x	A vector containing median values obtained from <a href="#">medCal</a> or <a href="#">binMed</a>
yrng	y-axis range of the bars.
width	width of the bars (optional)
col	color of the bars
border	the color to draw the border. Use border = NA to omit borders.
...	Additional arguments affecting the plot

**See Also**

[medCal](#); [binMed](#)

**Examples**

```
## Not run:
#Load EUROEVOL Data
data(euroevol)

#Subset Danish Dates
denmark <- subset(euroevol, Country=="Denmark")

#Calibrate and Bin
denmarkDates <- calibrate(x=denmark$C14Age, errors=denmark$C14SD)
```

```

denmarkBins <- binPrep(sites=denmark$SiteID,ages=denmark$C14Age,h=200) #200 years bin size

#Compute median date for each bin
bm <- binMed(x=denmarkDates,bins=denmarkBins)

#Compute median date for each date
dm <- medCal(denmarkDates)

#Compute SPD
denmarkSPD <- spd(x=denmarkDates,bins=denmarkBins,timeRange=c(10000,4000))

#Plot SPD and barCodes of median dates
plot(denmarkSPD,runm=200)
barCodes(dm,yrng=c(0,0.01))

#Plot SPD and barCodes of median bins in BC/AD
plot(denmarkSPD,runm=200,calendar="BCAD")
barCodes(BPtoBCAD(bm),yrng=c(0,0.01))

## End(Not run)

```

---

BCADtoBP

---

*Convert BC/AD dates to BP format*


---

## Description

Converts BC/AD dates to BP format while handling the absence of 'year 0'

## Usage

```
BCADtoBP(x)
```

## Arguments

x                    A numerical vector (currently only basic checks that these numbers are in a sensible range).

## Value

A vector with BP dates.

## Examples

```
BCADtoBP(-1268)
```

---

binMed	<i>Computes the median date of each bin</i>
--------	---

---

## Description

Function for generating a vector of median calibrated dates for each bin.

## Usage

```
binMed(x, bins, verbose = TRUE)
```

## Arguments

x	A CalDates class object.
bins	vector containing the bin names associated with each radiocarbon date. Can be generated using <a href="#">binPrep</a> .
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.

## Value

A vector of median dates in cal BP

## See Also

[binPrep](#), [barCodes](#)

## Examples

```
## Not run:  
#Load EUROEVOL Data  
data(euroevol)  
#Subset Danish Dates  
denmark <- subset(euroevol, Country=="Denmark")  
#Calibrate and Bin  
denmarkDates <- calibrate(x=denmark$C14Age, errors=denmark$C14SD)  
denmarkBins <- binPrep(sites=denmark$SiteID, ages=denmark$C14Age, h=200) #200 years bin size  
#Compute median date for each bin  
binMed(x=denmarkDates, bins=denmarkBins)  
  
## End(Not run)
```

---

binPrep

*Binning function of radiocarbon dates.*

---

## Description

Prepare a set of bins for controlling the aggregation of radiocarbon dates known to be from the same phase of same archaeological site (for use with [spd](#)). Used in cases where there is a concern that unusually high levels of sampling for radiocarbon at a given site or in a given site phase will impede comparison between sites or phases.

## Usage

```
binPrep(sites, ages, h, method = "complete")
```

## Arguments

sites	a vector of character strings (or number to coerce to character) of all sites or site phases. If character strings are used these should not contain underscores (see also below)
ages	a vector of uncalibrated conventional radiocarbon ages or a CalDates class object obtained using the <a href="#">calibrate</a> function.
h	a single numeric value passed to <a href="#">cutree</a> control degree of grouping of similar ages in a phase site.
method	the agglomeration method to be used, passed on to <a href="#">hclust</a> . Defaults to "complete" as in <a href="#">hclust</a> .

## Details

If ages is a CalDates class object, median dates are used for the clustering.

## Value

A vector of character strings with the same length of the object supplied for the argument ages identifying intra-site or intra-phase grouping, for use with [spd](#). The character strings effectively provide a "name" for each "phase" within a "site", using sequential integers after an underscore. For example if a site named "S001" had four dates grouped into two bins with two dates each, the resulting vector would be "S001\_1", "S001\_1", "S001\_2", and "S001\_2".

## See Also

[spd](#) for generating SPD; [binsense](#) for sensitivity analysis pertaining the choice of the parameter h.



---

binsense                      *Bin sensitivity Plot*

---

### Description

Visually explores how choosing different values for `h` in the `binPrep` function affects the shape of the SPD.

### Usage

```
binsense(
  x,
  y,
  h,
  timeRange,
  calendar = "BP",
  binning = "CRA",
  raw = F,
  verbose = T,
  legend = T,
  ...
)
```

### Arguments

<code>x</code>	A <code>CalDates</code> class object containing calibrated radiocarbon dates.
<code>y</code>	A vector containing the locations ID (e.g. site ID) of each calibrated date to be used for the binning process.
<code>h</code>	A vector of numbers containing values for the <code>h</code> parameter to be used in the <code>binPrep</code> function.
<code>timeRange</code>	A vector of length 2 indicating the start and end date of the analysis in cal BP.
<code>calendar</code>	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
<code>binning</code>	Either 'CRA' or 'calibrated'. Indicate whether the binning should be carried using the 14C age or using the median calibrated date. Default is 'CRA'.
<code>raw</code>	A logical variable indicating whether all SPDs should be returned or not. Default is FALSE.
<code>verbose</code>	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.
<code>legend</code>	A logical variable indicating whether the legend should be displayed. Default is TRUE
<code>...</code>	Additional arguments to be passed to the <code>spd</code> function.

**See Also**

[binPrep](#); [spd](#)

**Examples**

```
## Not run:
data(euroevol)
#subset Danish dates
denmark=subset(euroevol, Country=="Denmark")
denmarkDates=calibrate(x=denmark$C14Age, errors=denmark$C14SD)
binsense(x=denmarkDates, y=denmark$SiteID, h=seq(0, 200, 20), timeRange=c(10000, 4000), runm=200)

## End(Not run)
```

---

BPtoBCAD

*Convert BP dates to BC/AD format*

---

**Description**

Converts calibrated BP dates to BC/AD dates, omitting 'year 0'

**Usage**

```
BPtoBCAD(x)
```

**Arguments**

x                    A numerical vector (currently only basic checks that these numbers are in a sensible range).

**Value**

A vector with BC/BCE dates expressed as negative numbers and AD/CE dates as positive ones.

**Examples**

```
BPtoBCAD(4200)
```

---

calibrate	<i>Calibrate radiocarbon dates</i>
-----------	------------------------------------

---

## Description

Function for calibrating radiocarbon dates or uncalibrated SPDs.

Function for calibrating a SPD generated by summing uncalibrated dates.

## Usage

```
calibrate(x, ...)  
  
## Default S3 method:  
calibrate(  
  x,  
  errors,  
  ids = NA,  
  dateDetails = NA,  
  calCurves = "intcal13",  
  resOffsets = 0,  
  resErrors = 0,  
  timeRange = c(50000, 0),  
  normalised = TRUE,  
  F14C = FALSE,  
  calMatrix = FALSE,  
  eps = 1e-05,  
  ncores = 1,  
  verbose = TRUE,  
  ...  
)  
  
## S3 method for class 'UncalGrid'  
calibrate(  
  x,  
  errors = 0,  
  calCurves = "intcal13",  
  timeRange = c(50000, 0),  
  eps = 1e-05,  
  type = "fast",  
  datenormalised = FALSE,  
  spdnormalised = FALSE,  
  verbose = TRUE,  
  ...  
)
```

**Arguments**

x	A vector of uncalibrated radiocarbon ages or a UncalGrid class object.
...	ignored
errors	A vector of standard deviations corresponding to each estimated radiocarbon age.
ids	An optional vector of IDs for each date.
dateDetails	An optional vector of details for each date which will be returned in the output metadata.
calCurves	Either a character string naming a calibration curve already provided with the rcarbon package (currently 'intcal13', 'intcal13nhpine16', 'shcal13', 'shcal13shkauri16', 'marine13' and 'normal' (i.e. no calibration) are possible; default is 'intcal13') or a custom calibration curve as three-column matrix or data.frame (calibrated year BP, uncalibrated age bp, standard deviation). Different existing curves can be specified per dated sample, but only one custom curve can be provided for all dates.
resOffsets	A vector of offset values for any marine reservoir effect (default is no offset).
resErrors	A vector of offset value errors for any marine reservoir effect (default is no offset).
timeRange	Earliest and latest data to calibrate for, in calendar years. Posterior probabilities beyond this range will be excluded (the default is sensible in most cases).
normalised	A logical variable indicating whether the calibration should be normalised or not. Default is TRUE.
F14C	A logical variable indicating whether calibration should be carried out in F14C space or not. Default is FALSE.
calMatrix	a logical variable indicating whether the age grid should be limited to probabilities higher than eps
eps	Cut-off value for density calculation. Default is 1e-5.
ncores	Number of cores/workers used for parallel execution. Default is 1 (>1 requires doParallel package).
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.
type	Currently, two options are available. With "fast", a look-up is conducted for each calendar year that picks up the amplitude of the nearest CRA age. With "full", each CRA is calibrated individually and then summed (much slower).
datenormalised	Controls for calibrated dates with probability mass outside the timerange of analysis. If set to TRUE the total probability mass within the time-span of analysis is normalised to sum to unity. Should be set to FALSE when the parameter normalised in <a href="#">calibrate</a> is set to FALSE. Default is FALSE.
spdnormalised	A logical variable indicating whether the total probability mass of the SPD is normalised to sum to unity.

## Details

This function computes one or more calibrated radiocarbon ages using the method described in Bronk Ramsey 2008 (see also Parnell 2017). It is possible to specify different calibration curves or reservoir offsets individually for each date, and control whether the resulting calibrated distribution is normalised to 1 under-the-curve or not. Calculations can also be executed in parallel to reduce computing time. The function was modified from the BchronCalibrate function in the Bchron package developed by A.Parnell (see references below).

## Value

An object of class CalDates with the following elements:

- `metadata` A data.frame containing relevant information regarding each radiocarbon date and the parameter used in the calibration process.
- `grids` A list of calGrid class objects, containing the posterior probabilities for each calendar year. The most memory-efficient way to store calibrated dates, as only years with non-zero probability are stored, but aggregation methods such as `spd()` may then take longer to extract and combine multiple dates. NA when the parameter `calMatrix` is set to TRUE.
- `calMatrix` A matrix of probability values, one row per calendar year in `timeRange` and one column per date. By storing all possible years, not just those with non-zero probability, this approach takes more memory, but speeds up `spd()` and is suggested whenever the latter is to be used. NA when the parameter `calMatrix` is set to FALSE.

## References

Bronk Ramsey, C. 2008. Radiocarbon dating: revolutions in understanding, *Archaeometry* 50.2: 249-75. DOI: <https://doi.org/10.1111/j.1475-4754.2008.00394.x>

Parnell, A. 2017. Bchron: Radiocarbon Dating, Age-Depth Modelling, Relative Sea Level Rate Estimation, and Non-Parametric Phase Modelling, R package: <https://CRAN.R-project.org/package=Bchron>

## Examples

```
x1 <- calibrate(x=4000, errors=30)
plot(x1)
summary(x1)
# Example with a Marine Date, using a DeltaR of 300 and a DeltaR error of 30
x2 <- calibrate(x=4000, errors=30, calCurves='marine13', resOffsets=300, resErrors=30)
plot(x2)
```

## Description

Computes a Composite Kernel Density Estimate (CKDE) from multiple sets of randomly sampled calendar dates.

**Usage**

```
ckde(x, timeRange, bw, normalised = FALSE)
```

**Arguments**

x	A simdates class object, generated using <a href="#">sampleDates</a> .
timeRange	A vector of length 2 indicating the start and end date of the analysis in cal BP.
bw	Kernel bandwidth to be used.
normalised	A logical variable indicating whether the contribution of individual dates should be equal (TRUE), or weighted based on the area under the curve of non-normalised calibration (FALSE). Default is TRUE.

**Details**

The function computes Kernel Density Estimates using randomly sampled calendar dates contained in a simdates class object (generated using the `simulate.dates()` function). The output contains `nsim` KDEs, where `nsim` is the argument used in `simulate.dates()`. The resulting object can be plotted to visualise a CKDE (cf Brown 2017), and if `boot` was set to `TRUE` in `sampleDates` its bootstrapped variant (cf McLaughlin 2018 for a similar analysis). The shape of the CKDE is comparable to an SPD generated from non-normalised dates when the argument `normalised` is set to `FALSE`.

**Value**

An object of class `ckdeSPD` with the following elements

- `timeRange` The `timeRange` setting used.
- `res.matrix` A matrix containing the KDE values with rows representing calendar dates.

**References**

Brown, W. A. 2017. The past and future of growth rate estimation in demographic temporal frequency analysis: Biodemographic interpretability and the ascendance of dynamic growth models. *Journal of Archaeological Science*, 80: 96–108. DOI: <https://doi.org/10.1016/j.jas.2017.02.003>

McLaughlin, T. R. 2018. On Applications of Space–Time Modelling with Open-Source 14C Age Calibration. *Journal of Archaeological Method and Theory*. DOI <https://doi.org/10.1007/s10816-018-9381-3>

**See Also**

[sampleDates](#)

**Examples**

```
## Not run:
data(emedyd)
x = calibrate(x=emedyd$CRA, errors=emedyd$Error, normalised=FALSE)
bins = binPrep(sites=emedyd$SiteName, ages=emedyd$CRA, h=50)
s = sampleDates(x, bins=bins, nsim=100, boot=FALSE)
```

```
ckdeNorm = ckde(s,timeRange=c(16000,9000),bw=100,normalised=TRUE)
plot(ckdeNorm,type='multiline',calendar='BCAD')

## End(Not run)
```

---

combine

*Combine multiple CalDates Class Objects into one.*

---

## Description

Combine multiple CalDates Class Objects into one.

## Usage

```
combine(..., fixIDs = FALSE)
```

## Arguments

... CalDates class objects to be concatenated.

fixIDs logical. If set to TRUE, each date is given a new ID based on sequential integer. Default is FALSE

## Value

An object of class CalDates

## See Also

[calibrate](#)

## Examples

```
x1 = calibrate(c(2000,3400),c(20,20),ids=1:2)
x2 = calibrate(c(4000,3000),c(30,30),calCurves=c('intcal13','marine13'),
resOffsets=c(0,30),resErrors=c(0,20),ids=3:4)
mcurve <- mixCurves('intcal13','marine13',p=0.7,resOffsets=300,resErrors=20)
x3 = calibrate(5300,20,calCurves=mcurve,ids=5)
x = combine(x1,x2,x3)
## x$metadata
```

emedyd

*Radiocarbon dates for the Eastern Mediterranean around the Younger Dryas*

## Description

Radiocarbon dates (n=1915) and site coordinates (n=201) from a paper considering the relationship between human activity in the eastern Mediterranean/Middle East and early Holocene climate change, including the Younger Dryas.

## Usage

emedyd

## Format

A data.frame with the following variables:

LabID Laboratory ID assigned to each radiocarbon date (where known)

CRA Radiocarbon age in 14C years BP

Error Radiocarbon age error

Material Material of the dated sample

Species Species of the dated sample (where identified)

SiteName Name of the site from which the sample has been recovered

Country Country where the sampling site is located

Longitude Longitude of the sampling site in decimal degrees

Latitude Latitude of the sampling site in decimal degrees

Region One of three analytical regions (1=southern Levant, 2=Northern Levant, 3= South-central Anatolia)

## Source

Palmisano, A., Bevan, A. and S. Shennan 2017. Data and code for demographic trends in the paper "Human responses and non-responses to climatic variations during the Last Glacial-Interglacial transition in the eastern Mediterranean", UCL Discovery Archive 1570274. doi:10.14324/000.ds.1570274.

## References

Roberts, N., Woodbridge, J., Bevan, A., Palmisano, A., Shennan, S. and E. Asouti 2017. Human responses and non-responses to climatic variations during the Last Glacial-Interglacial transition in the eastern Mediterranean. *Quaternary Science Reviews*, 184, 47-67. doi:10.1016/j.quascirev.2017.09.011.



## Examples

```
## Not run:
data(emedyd)
northernlevant <- emedyd[emedyd$Region=="2",]
bins <- binPrep(northernlevant$SiteName, northernlevant$CRA, h=50)
x <- calibrate(northernlevant$CRA, northernlevant$error, normalised=FALSE)
spd.northernlevant <- spd(x, bins=bins, runm=50, timeRange=c(17000,8000))
plot(spd.northernlevant)

## End(Not run)
```

---

euroevol

*Radiocarbon dates from the EUROEVOL database*

---

## Description

Radiocarbon dates (n=14,053) and site coordinates (n=4,213) from the EUROEVOL project database. Sites without radiocarbon dates (n=544), phase-codes, and other data have been omitted.

## Usage

```
euroevol
```

## Format

A data.frame with the following variables:

C14ID ID of each radiocarbon date

C14Age Radiocarbon age in 14C years BP

C14SD Radiocarbon age error

LabCode Labcode of the radiocarbon date

Material Material of the dated sample

SiteID ID of the site from which the sample has been recovered

Latitude Latitude of the sampling site in decimal degrees

Longitude Longitude of the sampling site in decimal degrees

Country Country where the sampling site is located

## Source

Manning, K., Colledge, S., Crema, E., Shennan, S., Timpson, A., 2016. The Cultural Evolution of Neolithic Europe. EUROEVOL Dataset 1: Sites, Phases and Radiocarbon Data. Journal of Open Archaeology Data 5. doi:10.5334/joad.40

## References

Shennan, S., Downey, S.S., Timpson, A., Edinborough, K., Colledge, S., Kerig, T., Manning, K., Thomas, M.G., 2013. Regional population collapse followed initial agriculture booms in mid-Holocene Europe. *Nature Communications* 4, ncomms3486. doi:10.1038/ncomms3486

Timpson, A., Colledge, S., Crema, E., Edinborough, K., Kerig, T., Manning, K., Thomas, M.G., Shennan, S., 2014. Reconstructing regional population fluctuations in the European Neolithic using radiocarbon dates: a new case-study using an improved method. *Journal of Archaeological Science* 52, 549-557. doi:10.1016/j.jas.2014.08.011

## Examples

```
## Not run:
data(euroevol)
Ireland <- subset(euroevol, Country=="Ireland")
bins <- binPrep(Ireland$SiteID, Ireland$C14Age, h=200)
x <- calibrate(Ireland$C14Age, Ireland$C14SD)
spd.ireland <- spd(x, bins=bins, runm=200, timeRange=c(8000, 4000))
plot(spd.ireland)

## End(Not run)
```

---

ewdates

---

*Subset of EUROEVOL radiocarbon dates from Great Britain*


---

## Description

Radiocarbon dates (n=2,324) and site coordinates (n=652) from England and Wales collected from the EUROEVOL project database. See [euroevol](#) for more details regarding the source data.

## Usage

```
ewdates
```

## Format

A data.frame with the following variables:

C14ID ID of each radiocarbon date

C14Age Radiocarbon age in <sup>14</sup>C years BP

C14SD Radiocarbon age error

LabCode Labcode of the radiocarbon date

Material Material of the dated sample

SiteID ID of the site from which the sample has been recovered

Eastings Easting coordinates of the sampling site in meters (OSGB 1936 epsg:27700)

Northings Northing coordinates of the sampling site in meters (OSGB 1936 epsg:27700)

**Source**

Manning, K., Colledge, S., Crema, E., Shennan, S., Timpson, A., 2016. The Cultural Evolution of Neolithic Europe. EUROEVOL Dataset 1: Sites, Phases and Radiocarbon Data. Journal of Open Archaeology Data 5. doi:10.5334/joad.40

---

ewowin

*Polygonal window of England and Wales*

---

**Description**

An [owin](#) class polygonal window of England and Wales.

**Usage**

```
ewowin
```

**Format**

An owin class object.

**Source**

South,A.2011.rworldmap: A New R package for Mapping Global Data. The R Journal Vol. 3/1 : 35-43.

**Examples**

```
## Not run:
data(ewowin)
# Obtained from rworldmap:
library(maptools)
library(rgeos)
library(rworldmap)
bng <- CRS("+init=epsg:27700")
sbrit <- getMap(resolution="high")
sbrit <- spTransform(sbrit[sbrit$GEOUNIT == "United Kingdom",], bng)
tmp <- SpatialPolygons(list(Polygons(list(Polygon(cbind(c(130000,130000,310000,
425000,700000,700000,130000), c(0,300000,560000,598000,300000,0,0))),
"1")), 1:1, proj4string=bng)
sbrit <- gIntersection(sbrit,tmp)
ewowin <- as.owin(sbrit)

## End(Not run)
```

---

gaussW	<i>Gaussian weighting of dates relative to</i>
--------	--

---

**Description**

Rescale a numeric vector to a specified minimum and maximum.

**Usage**

```
gaussW(x, mean, sd, type = "weights")
```

**Arguments**

x	A numeric vector or an object of class CalDates.
mean	A single numeric value indicating the value to centre the Gaussian kernel on.
sd	A single numeric value indicating the standard deviation of the Gaussian kernel to be used.
type	The type of output to produce: currently either "weighted" (for a simple total weight value for each date) or "raw" (a list of reweighted calibrated radiocarbon probabilities for each calibrated date).

**Value**

A numeric vector of weights (or optionally a list of reweighted calibrated radiocarbon probabilities).

**Examples**

```
## Example weighting fo a set of dates versus a focal date of 5950 calBP
years <- seq(6500, 5500, -10)
plot(cbind(years, gaussW(years, 5950, 50)))
## Example weighting of three calibrated dates versus a focal date of 5950 calBP
dates <- calibrate(c(5280, 5180, 5080), c(30,30,30), normalised=FALSE)
gaussW(dates, 5950, 50)
## Or the same with raw output
dateswt <- gaussW(dates, 5950, 50, type="raw")
head(dateswt[[1]])
```

---

hpd	<i>Computes the highest posterior density interval (HPDI) for calibrated dates</i>
-----	--

---

**Description**

Computes the highest posterior density interval (HPDI) for calibrated dates

**Usage**

```
hpdi(x, credMass = 0.95)
```

**Arguments**

x	A CalDates class object.
credMass	Interval probability mass

**Value**

A list of matrices with HPDI ranges in cal BP

**See Also**

[calibrate](#)

**Examples**

```
x <- calibrate(c(3050,2950),c(20,20))
hpdi(x)
```

---

medCal

*Computes the median date of calibrated dates*

---

**Description**

Function for generating a vector median calibrated dates from a CalDates class object.

**Usage**

```
medCal(x)
```

**Arguments**

x	A CalDates class object.
---	--------------------------

**Value**

A vector of median dates in cal BP

**See Also**

[calibrate](#), [barCodes](#)

**Examples**

```
x <- calibrate(c(3050,2950),c(20,20))
medCal(x)
```

---

mixCurves	<i>Creates mixed calibration curves.</i>
-----------	--

---

### Description

Function for generating mixed calibration curves (e.g. intcal13/marine13, intcal13/shcal13) with user-defined proportions.

### Usage

```
mixCurves(
  calCurve1 = "intcal13",
  calCurve2 = "intcal13",
  p = 0.5,
  resOffsets = 0,
  resErrors = 0
)
```

### Arguments

calCurve1	Name of the first curve, chosen from 'intcal13','intcal13nhpine16','shcal13','shcal13shkauri16','marine13'. Default is 'intcal13'.
calCurve2	Name of the second curve, from the same list. Default is 'marine13'.
p	The proportion of contribution of the first curve. Default is 1.
resOffsets	Offset value for the marine reservoir effect to be applied to calCurve2. Default is 0.
resErrors	Error of the marine reservoir effect offset to be applied to calCurve2. Default is 0.

### Details

The function is based on the `mix.calibrationcurves` function of the `clam` package.

### Value

A three-column matrix containing calibrated year BP, uncalibrated age bp, and standard deviation. To be used as custom calibration curve for the `calibrate` function.

### References

Blaauw, M. and Christen, J.A.. 2011. Flexible paleoclimate age-depth models using an autoregressive gamma process. *Bayesian Analysis*, 6, 457-474. Blaauw, M. 2018. clam: Classical Age-Depth Modelling of Cores from Deposits. R package version 2.3.1. <https://CRAN.R-project.org/package=clam>  
 Marsh, E.J., Bruno, M.C., Fritz, S.C., Baker, P., Capriles, J.M. and Hastorf, C.A., 2018. IntCal, SHCal, or a Mixed Curve? Choosing a 14 C Calibration Curve for Archaeological and Paleoenviromental Records from Tropical South America. *Radiocarbon*, 60(3), pp.925-940.

**See Also**[calibrate](#)**Examples**

```
myCurve <- mixCurves('intcal13', 'marine13', p=0.7, resOffsets=300, resErrors=20)
x <- calibrate(4000, 30, calCurves=myCurve)
```

---

`modelTest`*Monte-Carlo simulation test for SPDs*

---

**Description**

Comparison of an observed summed radiocarbon date distribution (aka SPD) with simulated outcomes from a theoretical model.

**Usage**

```
modelTest(  
  x,  
  errors,  
  nsim,  
  bins = NA,  
  runm = NA,  
  timeRange = NA,  
  backsight = 50,  
  changexpr = expression((t1/t0)^(1/d) - 1),  
  gridclip = TRUE,  
  raw = FALSE,  
  model = c("exponential"),  
  method = c("uncalsample"),  
  predgrid = NA,  
  normalised = NA,  
  datenormalised = NA,  
  spdnormalised = FALSE,  
  ncores = 1,  
  fitonly = FALSE,  
  a = 0,  
  b = 0,  
  edgeSize = 500,  
  verbose = TRUE  
)
```

**Arguments**

x	A CalDates object containing calibrated radiocarbon ages
errors	A vector of errors corresponding to each radiocarbon age
nsim	Number of simulations
bins	A vector indicating which bin each radiocarbon date is assigned to.
runm	A number indicating the window size of the moving average to smooth both observed and simulated SPDs. If set to NA no moving average is applied. Default is NA.
timeRange	A vector of length 2 indicating the start and end date of the analysis in cal BP. The fitting process is applied considering the SPD within the interval defined by this parameter. If no values are supplied the earliest and latest median calibrated dates of the observed data will be used.
backsight	A single numeric value defining the distance in time between the focal year and the backsight year for computing the rate of change. Default is 10.
changexpr	An expression for calculating the rate of change in SPD between the focal year and a backsight year. Available input options are t1 (the SPD for the focal year), t0 (the SPD for the backsight year), d (the distance between t0 and t1), and any other standard constants and mathematical operators. A sensible default is provided.
gridclip	Whether the sampling of random dates is constrained to the observed range (TRUE) or not (FALSE). Default is TRUE.
raw	A logical variable indicating whether all permuted SPDs should be returned or not. Default is FALSE.
model	A vector indicating the model to be fitted. Currently the acceptable options are 'uniform', 'linear', 'exponential' and 'custom'. Default is 'exponential'.
method	Method for the creation of random dates from the fitted model. Either 'uncalsample' or 'calsample'. Default is 'uncalsample'. See below for details.
predgrid	A data.frame containing calendar years (column calBP) and associated summed probabilities (column PrDens). Required when model is set to 'custom'.
normalised	Whether the simulated dates should be normalised or not. Default based on whether x is normalised or not.
datenormalised	Argument kept for backward compatibility with previous versions.
spdnormalised	A logical variable indicating whether the total probability mass of the SPD is normalised to sum to unity for both observed and simulated data.
ncores	Number of cores used for for parallel execution. Default is 1.
fitonly	A logical variable. If set to TRUE, only the the model fitting is executed and returned. Default is FALSE.
a	Starter value for the exponential fit with the <code>nls</code> function using the formula $y \sim \exp(a + b * x)$ where y is the summed probability and x is the date. Default is 0.
b	Starter value for the exponential fit with the <code>nls</code> function using the formula $y \sim \exp(a + b * x)$ where y is the summed probability and x is the date. Default is 0.



edgeSize	Controls edge effect by expanding the fitted model beyond the range defined by timeRange.
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.

## Details

The function implements a Monte-Carlo test for comparing a theoretical or fitted statistical model to an observed summed radiocarbon date distribution (aka SPD) and associated rates of changes. A variety of theoretical expectations can be compared to the observed distribution by setting the `model` argument, for example to fit basic 'uniform' (the mean of the SPD), 'linear' (fitted using the `lm` function) or `model='exponential'` models (fitted using the `nls` function). Models are fitted to the period spanned by `timeRange` although `x` can contain dates outside this range to mitigate possible edge effects (see also `bracket`). Alternatively, it is possible for the user to provide a model of their own by setting `model='custom'` and then supplying a two-column `data.frame` to `predgrid`. The function generates `nsim` theoretical SPDs from the fitted model via Monte-Carlo simulation, this is then used to define a 95% critical envelope for each calendar year. The observed SPD is then compared against the simulation envelope; local departures from the model are defined as instances where the observed SPD is outside such an envelope, while an estimate of the global significance of the observed SPD is also computed by comparing the total areas of observed and simulated SPDs that fall outside the simulation envelope. The theoretical SPDs can be generated using two different sampling approaches defined by the parameter `method`. If `method` is set to 'uncalsample' each date is drawn after the fitted model is backcalibrated as a whole and adjusted for a baseline expectation; if it is set to 'calsample' samples are drawn from the fitted model in calendar year then individually back calibrated and recalibrated (the approach of Timpson et al. 2014). For each simulation, both approaches produces  $n$  samples, with  $n$  equal to the number of bins or number of dates (when bins are not defined). Differences between these two approaches are particularly evident at dates coincident with steeper portions of the calibration curve. If more than one type of calibration curve is associated with the observed dates, at each Monte-Carlo iteration, the function randomly assigns each bin to one of the calibration curves with probability based on the proportion of dates within the bin associated to the specific curves. For example, if a bin is composed of four dates and three are calibrated with 'intcal13' the probability of that particular bin being assigned to 'intcal13' is 0.75.

## Value

An object of class `SpdModelTest` with the following elements

- `result` A four column `data.frame` containing the observed probability density (column `PrDens`) and the lower and the upper values of the simulation envelope (columns `lo` and `hi`) for each calendar year column `calBP`
- `result.roc` A four column `data.frame` containing the observed rates of change (column `roc`) and the lower and the upper values of the simulation envelope (columns `lo.roc` and `hi.roc`) for the mid point between two chronological blocks `calBP`
- `sim` A matrix containing the simulation results of the summed probabilities. Available only when `raw` is set to TRUE
- `sim.roc` A matrix containing the simulation results of the rate of change of summed probabilities. Available only when `raw` is set to TRUE

- `pval` A numeric vector containing the p-value of the global significance test for the summed probabilities
- `pval.roc` A numeric vector containing the p-value of the global significance test for the rates of change
- `fit` A data.frame containing the probability densities of the fitted model for each calendar year within the time range of analysis
- `fitobject` Fitted model. Not available when `model` is 'custom'
- `n` Number of radiocarbon dates.
- `nbins` Number of bins.
- `nsim` Number of Monte-Carlo simulations.
- `backsight` Backsight size.

### Note

- Windows users might receive a memory allocation error with larger time span of analysis (defined by the parameter `timeRange`). This can be avoided by increasing the memory limit with the `memory.limit` function.
- Users experiencing a `Error: cannot allocate vector of size ...` error message can increase the memory size using the `Sys.setenv()`, for example: `Sys.setenv("R_MAX_VSIZE" = 16e9)`.
- The function currently supports only dates calibrated with 'intcal13', 'intcal13nhpine16', 'shcal13', 'shcal13shkauri16', and 'marine13'.

### References

Timpson, A., Colledge, S., Crema, E., Edinborough, K., Kerig, T., Manning, K., Thomas, M.G., Shennan, S., (2014). Reconstructing regional population fluctuations in the European Neolithic using radiocarbon dates: a new case-study using an improved method. *Journal of Archaeological Science*, 52, 549-557. doi:10.1016/j.jas.2014.08.011

### Examples

```
## Example with Younger Dryas period Near East, including site bins
## Not run:
data(emedyd)
caldates <- calibrate(x=emedyd$CRA, errors=emedyd$Error, normalised=FALSE)
bins <- binPrep(sites=emedyd$SiteName, ages=emedyd$CRA, h=50)
nsim=5 #toy example
expnull <- modelTest(caldates, errors=emedyd$Error, bins=bins, nsim=nsim, runm=50,
timeRange=c(16000,9000), model="exponential", datenormalised=FALSE)
plot(expnnull, xlim=c(16000,9000))
round(expnnull$pval,4) #p-value
summary(expnnull)

## End(Not run)
```

multiplot

*Plot multiple dates***Description**

Plot multiple radiocarbon dates.

**Usage**

```

multiplot(
  x,
  type = "d",
  calendar = "BP",
  HPD = FALSE,
  credMass = 0.95,
  decreasing = NULL,
  label = TRUE,
  xlim = NULL,
  xlab = NA,
  ylab = NA,
  col.fill = "grey50",
  col.fill2 = "grey82",
  col.line = "black",
  lwd = 1,
  cex.id = 1,
  cex.lab = 1,
  cex.axis = 1,
  ydisp = FALSE,
  gapFactor = 0.2
)

```

**Arguments**

x	A CalDates class object with length > 1.
type	Whether the calibrated dates are displayed as distributions ('d') or as horizontal bars ('b') spanning the HPD interval. Default is 'd'.
calendar	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
HPD	Logical value indicating whether intervals of higher posterior density should be displayed. Default is FALSE.
credMass	A numerical value indicating the size of the higher posterior density interval. Default is 0.95.
decreasing	Whether dates should be plotted with decreasing order of median calibrated date (i.e. old to new; TRUE) or increasing order (i.e. new to old; FALSE). If set to NULL the dates plotted in the supplied order. Default is NULL

label	Whether the ID of each date should be displayed. Default is TRUE.
xlim	the x limits of the plot. In BP or in BC/AD depending on the choice of the parameter calender. Notice that if BC/AD is selected BC ages should have a minus sign (e.g. c(-5000, 200) for 5000 BC to 200 AD).
xlab	(optional) Label for the x axis. If unspecified the default setting will be applied ("Year BP" or "Year BC/AD").
ylab	(optional) Label for the y axis.
col.fill	A vector of primary fill color for the calibrated date distribution. Default is 'grey50'.
col.fill2	A vector of secondary secondary colour fill color for the calibrated date distribution, used for regions outside the higher posterior interval. Ignored when HPD=FALSE. Default is 'grey82'.
col.line	A vector of line color (ignored when type is set to 'd'). Default is 1.
lwd	Line width (ignored when type is set to 'd'). Default is 1.
cex.id	The magnification to be used the date labels relative to the current setting of cex. Default is adjusted to 1.
cex.lab	The magnification to be used for x and y labels relative to the current setting of cex. Default is adjusted to 1.
cex.axis	The magnification to be used for axis annotation relative to the current setting of cex. Default is adjusted to 1.
ydisp	Whether the y axis should be displayed. Ignored when type is set to 'b'. Default is FALSE
gapFactor	Defines spacing between calibrated dates (when type is set to 'd') or the distance between the lines and the labels (when type is set to 'b') as proportion of individual y-axis ranges. Default is 0.2.

### See Also

[calibrate](#)

### Examples

```
data("emedyd")
tellaswad = subset(emedyd, SiteName=='Tell Aswad')
x = calibrate(tellaswad$CRA, tellaswad$Error, ids=tellaswad$LabID)
multiplot(x, HPD=TRUE, decreasing=TRUE, label=FALSE, gapFactor = 0.1)
multiplot(x, type='b', calendar='BCAD', cex.id = 0.5, lwd=2, gapFactor = 0.5)
```

---

p2pTest *Point to point test of SPD*

---

### Description

Test for evaluating the difference in the summed probability values associated with two points in time.

### Usage

```
p2pTest(x, p1 = NA, p2 = NA, interactive = TRUE, plot = FALSE)
```

### Arguments

x	result of <a href="#">modelTest</a> with raw=TRUE.
p1	calendar year (in BP) of start point.
p2	calendar year (in BP) of end point.
interactive	if set to TRUE enables an interactive selection of p1 and p2 from a graphical display of the SPD. Disabled when p1 and p2 are defined.
plot	if set to TRUE the function plots the location of p1 and p2 on the SPD. Default is FALSE.

### Details

The function compares observed differences in the summed probability values associated with two points in time against a distribution of expected values under the null hypothesis defined with the [modelTest](#) function. The two points can be specified manually (assigning BP dates to the arguments p1 and p2) or interactively (clicking on a SPD plot). Note that [modelTest](#) should be executed setting the argument raw to TRUE (default is FALSE).

### Value

A list with: the BP dates for the two points and the p-value obtained from a two-sided test.

### References

Edinburgh, K., Porcic, M., Martindale, A., Brown, T.J., Supernant, K., Ames, K.M., (2017). Radiocarbon test for demographic events in written and oral history. PNAS 201713012. doi:10.1073/pnas.1713012114

### See Also

[modelTest](#).

**Examples**

```
## Example with Younger Dryas period Near East, including site bins
## Not run:
data(emedyd)
caldates <- calibrate(x=emedyd$CRA, errors=emedyd$Error, normalised=FALSE)
bins <- binPrep(sites=emedyd$SiteName, ages=emedyd$CRA, h=50)
nsim=10 #toy example
expnull <- modelTest(caldates, errors=emedyd$Error, bins=bins, nsim=nsim, runm=50,
timeRange=c(16000,9000), model="exponential", datenormalised=FALSE, raw=TRUE)
p2pTest(x=expnull,p1=13000,p2=12500) #non-interactive mode
p2pTest(x=expnull) #interactive mode

## End(Not run)
```

---

permTest

*Random mark permutation test for SPDs*


---

**Description**

Global and local significance test for comparing shapes of multiple SPDs using random permutations.

**Usage**

```
permTest(
  x,
  marks,
  timeRange,
  backsight = 10,
  changexpr = expression((t1/t0)^(1/d) - 1),
  nsim,
  bins = NA,
  runm = NA,
  datenormalised = FALSE,
  spdnormalised = FALSE,
  raw = FALSE,
  verbose = TRUE
)
```

**Arguments**

x	A CalDates class object containing the calibrated radiocarbon dates.
marks	A numerical or character vector containing the marks associated to each radiocarbon date.
timeRange	A vector of length 2 indicating the start and end date of the analysis in cal BP.
backsight	A single numeric value defining the distance in time between the focal year and the backsight year for computing the rate of change. Default is 10.

changexpr	An expression for calculating the rate of change in SPD between the focal year and a backsight year. Available input options are t1 (the SPD for the focal year), t0 (the SPD for the backsight year), d (the distance between t0 and t1), and any other standard constants and mathematical operators. A sensible default is provided.
nsim	Number of random permutations
bins	A vector indicating which bin each radiocarbon date is assigned to.
runm	A number indicating the window size of the moving average to smooth the SPD. If set to NA no moving average is applied. Default is NA.
datenormalised	If set to TRUE the total probability mass of each calibrated date will be made to sum to unity (the default in most radiocarbon calibration software). This argument will only have an effect if the dates in x were calibrated without normalisation (via normalised=FALSE in the <code>calibrate</code> function), in which case setting datenormalised=TRUE here will rescale each dates probability mass to sum to unity before aggregating the dates, while setting datenormalised=FALSE will ensure unnormalised dates are used for both observed and simulated SPDs. Default is FALSE.
spdnormalised	A logical variable indicating whether the total probability mass of the SPD is normalised to sum to unity.
raw	A logical variable indicating whether all permuted SPDs should be returned or not. Default is FALSE.
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.

## Details

The function generates a distribution of expected SPDs by randomly shuffling the marks assigned to each *bin* (see `spd` for details on binning). The resulting distribution of probabilities for each *mark* (i.e. group of dates) for each calendar year is z-transformed, and a 95% simulation envelope is computed. Local significant departures are defined as instances where the observed SPD (which is also z-transformed) is outside such envelope. A global significance is also computed by comparing the total "area" outside the simulation envelope in the observed and simulated data.

## Value

An object of class `SpdPermTest` with the following elements

- `observed` A list containing data.frames with the summed probability (column *PrDens* for each calendar year (column *calBP* for each mark/group
- `envelope` A list containing matrices with the lower and upper bound values of the simulation envelope for each mark/group
- `pvalueList` A list of p-value associated with each mark/group

## References

Crema, E.R., Habu, J., Kobayashi, K., Madella, M., (2016). Summed Probability Distribution of 14 C Dates Suggests Regional Divergences in the Population Dynamics of the Jomon Period in Eastern Japan. PLOS ONE 11, e0154809. doi:10.1371/journal.pone.0154809

**Examples**

```
## compare demographic trajectories in Netherlands and Denmark
## Not run:
data(euroevol)
nld.dnk = subset(euroevol, Country=="Netherlands"|Country=="Denmark")
bins = binPrep(nld.dnk$SiteID, nld.dnk$C14Age, h=200)
dates = calibrate(nld.dnk$C14Age, nld.dnk$C14SD, normalised=FALSE)
res = permTest(dates, marks=as.character(nld.dnk$Country), nsim=1000,
bins=bins, runm=200, timeRange=c(10000, 4000))
round(res$pValueList, 4) #extract p-values
summary(res)
par(mfrow=c(2, 1))
plot(res, focalm="Netherlands", main="Netherlands")
plot(res, focalm="Denmark", main="Denmark")

## End(Not run)
```

---

plot.CalDates

*Plot calibrated dates*


---

**Description**

Plot calibrated radiocarbon dates.

**Usage**

```
## S3 method for class 'CalDates'
plot(
  x,
  ind = 1,
  label = NA,
  calendar = "BP",
  type = "standard",
  xlab = NA,
  ylab = NA,
  axis4 = TRUE,
  HPD = FALSE,
  credMass = 0.95,
  customCalCurve = NA,
  add = FALSE,
  col = "grey50",
  col2 = "grey82",
  cex.axis = 0.75,
  cex.xylab = 0.75,
  cex.label = 0.75,
  ...
)
```



**Arguments**

x	CalDates class object containing calibrated radiocarbon dates.
ind	Number indicating the index value of the calibrated radiocarbon date to be displayed. Default is 1.
label	(optional) Character vector to be shown on the top-right corner of the display window.
calendar	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
type	Either 'standard' or 'auc'. If set to 'auc', displays both the normalised (dashed line) and unnormalised curves. Default is 'standard'.
xlab	(optional) Label for the x axis. If unspecified the default setting will be applied ("Year BP" or "Year BC/AD").
ylab	(optional) Label for the y axis. If unspecified the default setting will be applied ("Radiocarbon Age").
axis4	Logical value indicating whether an axis of probabilities values should be displayed. Default is TRUE.
HPD	Logical value indicating whether intervals of higher posterior density should be displayed. Default is FALSE.
credMass	A numerical value indicating the size of the higher posterior density interval. Default is 0.95.
customCalCurve	A three column data.frame or matrix that allows you to pass and plot a custom calibration curve if you used one during calibration. You can currently only provide one such custom curve which is used for all dates.
add	if set to TRUE the calibrated date is displayed over the existing plot. Default is FALSE.
col	The primary fill color for the calibrated date distribution.
col2	The secondary colour fill color for the calibrated date distribution, used for regions outside the higher posterior interval. Ignored when HPD=FALSE.
cex.axis	The magnification to be used for axis annotation relative to the current setting of cex. Default is adjusted to 0.75.
cex.xylob	The magnification to be used for x and y labels relative to the current setting of cex. Default is adjusted to 0.75.
cex.label	The magnification to be used for the label on the top right corner defined by the argument label. Default is adjusted to 0.75.
...	Additional arguments affecting the plot.

**See Also**

[calibrate](#)

**Examples**

```
x <- calibrate(x=c(3402,3490,4042),errors=c(20,20,30))
plot(x) #display the first date
plot(x,2) #displays the second date
plot(x,3, calendar="BCAD", HPD=TRUE) #display in BC/AD with higher posterior density interval
```

---

plot.CalGrid

*Plot a summed probability distribution (from a CalGrid object)*


---

**Description**

Plot a summed radiocarbon probability distribution. This is a basic function for plotting SPDs that have been constructed manually or by calibrating a summed or otherwise irregular CRA grid. In most instances, it is sensible to use plot.CalSPD instead.

**Usage**

```
## S3 method for class 'CalGrid'
plot(
  x,
  runm = NA,
  calendar = "BP",
  fill.p = "grey50",
  border.p = NA,
  xlim = NA,
  ylim = NA,
  cex.lab = 0.75,
  cex.axis = cex.lab,
  mar = c(4, 4, 1, 3),
  add = FALSE,
  ...
)
```

**Arguments**

x	A "CalGrid" class object of summed probabilities per calendar year BP.
runm	A number indicating the window size of the moving average to smooth the SPD. If set to NA no moving average is applied. Default is NA
calendar	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
fill.p	Fill colour of the polygon depicting the summed probability distribution.
border.p	Border colour of the polygon depicting the summed probability distribution.
xlim	the x limits of the plot. In BP or in BC/AD depending on the choice of the parameter calendar. Notice that if BC/AD is selected BC ages should have a minus sign (e.g. c(-5000,200) for 5000 BC to 200 AD).

ylim	Adjust y axis limits (otherwise sensible default).
cex.lab	Size of label text.
cex.axis	Size of axis text.
mar	Adjust margins around plot.
add	Whether or not the new graphic should be added to an existing plot.
...	Additional arguments affecting the plot

**See Also**

[spd](#); [plot.CalSPD](#)

**Examples**

```
data(euroevol)
mycaldates <- calibrate(euroevol[1:10,"C14Age"], euroevol[1:10,"C14SD"], normalised=FALSE)
myspd <- spd(mycaldates, timeRange=c(8000,2000))
plot(myspd) #ordinary plot using \code{plot.CalSPD}
plot(myspd$grid) #working plot using the internal CalGrid object
```

---

plot.CalSPD

*Plot a summed probability distribution*

---

**Description**

Plot a summed probability distribution (SPD) of radiocarbon dates

**Usage**

```
## S3 method for class 'CalSPD'
plot(
  x,
  runm = NA,
  calendar = "BP",
  type = "standard",
  xlim = NA,
  ylim = NA,
  ylab = "Summed Probability",
  spdnormalised = FALSE,
  rescale = FALSE,
  fill.p = "grey75",
  border.p = NA,
  xaxt = "s",
  yaxt = "s",
  add = FALSE,
  cex.axis = 1,
  ...
)
```

**Arguments**

x	A CalSPD class object.
runm	A number indicating the window size of the moving average to smooth the SPD. If set to NA no moving average is applied. Default is NA
calendar	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
type	Either 'standard' or 'simple'. The former visualise the SPD as an area graph, while the latter as line chart.
xlim	the x limits of the plot. In BP or in BC/AD depending on the choice of the parameter calendar. Notice that if BC/AD is selected BC ages should have a minus sign (e.g. c(-5000,200) for 5000 BC to 200 AD).
ylim	the y limits of the plot.
ylab	(optional) Label for the y axis. If unspecified the default setting will be applied ("Summed Probability")
spdnormalised	A logical variable indicating whether the total probability mass of the SPD is normalised to sum to unity.
rescale	A logical variable indicating whether the SPD should be rescaled to range 0 to 1.
fill.p	Fill colour for the SPD
border.p	Border colour for the SPD
xaxt	Whether the x-axis tick marks should be displayed (xaxt='s', default) or not (xaxt='n').
yaxt	Whether the y-axis tick marks should be displayed (xaxt='s', default) or not (xaxt='n').
add	Whether or not the new graphic should be added to an existing plot.
cex.axis	The magnification to be used for axis annotation relative to the current setting of cex. Default is 1.
...	Additional arguments affecting the plot

**See Also**

[spd](#); [plot.CalGrid](#)

**Examples**

```
## Not run:
data(emedyd)
levant <- emedyd[emedyd$Region=="1"|emedyd$Region=="2",]
bins <- binPrep(levant$SiteName, levant$CRA, h=50)
x <- calibrate(levant$CRA, levant$error, normalised=FALSE)
spd.levant <- spd(x, bins=bins, timeRange=c(17000,8000))
spd.northernlevant <- spd(x[levant$Region=="2"], bins=bins[levant$Region=="2"],
timeRange=c(17000,8000))
plot(spd.levant, runm=50, xlim=c(16000,9000))
plot(spd.northernlevant, runm=50, add=TRUE, fill.p="black")
```

```

legend("topleft", legend=c("All Levant dates", "Northern Levant only"),
fill=c("grey75", "black"), border=NA)
plot(spd.levant, runm=50, xlim=c(16000, 9000), type="simple")
plot(spd.northernlevant, runm=50, col="red", type="simple", add=TRUE)

## End(Not run)

```

---

plot.compositeKDE	<i>Plots a Composite Kernel Density Estimate of sampled radiocarbon dates.</i>
-------------------	--

---

## Description

Plots a Composite Kernel Density Estimate of sampled radiocarbon dates.

## Usage

```

## S3 method for class 'compositeKDE'
plot(
  x,
  calendar = "BP",
  type = "envelope",
  ylim = NA,
  xlim = NA,
  fill.col = "lightgrey",
  interval = 0.95,
  line.col = "black",
  line.type = 2,
  multiline.alpha = NA,
  multiline.col = "black",
  ...
)

```

## Arguments

x	A compositeKDE class object generated using the <a href="#">ckde</a> function.
calendar	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
type	Either envelope or multiline. Default is envelope.
ylim	the y limits of the plot.
xlim	the x limits of the plot. In BP or in BC/AD depending on the choice of the parameter calendar. Notice that if BC/AD is selected BC ages should have a minus sign (e.g. c(-5000, 200) for 5000 BC to 200 AD).
fill.col	Envelope color when type='envelope'. Default is 'lightgrey'.
interval	Quantile interval for the envelope. Default is 0.95.
line.col	Line color when type='envelope'. Default is 'black'.

`line.type` Line type when `type='envelope'`. Default is 2.  
`multiline.alpha` Alpha level for line transparency when `type='multiline'`. Default is  $10/\text{nsim}$ , where `nsim` is the number of simulations. If `nsim` is smaller than 10, `multiline.alpha` will be set to 1.  
`multiline.col` Line color when `type='multiline'`. Default is 'black'.  
`...` Additional arguments affecting the plot

### Details

Visualise a `compositeKDE` class object. If `type` is set 'envelope' an envelope of the percentile interval defined by the parameter `interval` is shown along with the mean KDE. If `type` is set 'multiline' all KDEs are shown.

### See Also

[ckde](#);

---

<code>plot.spatialTest</code>	<i>Plot results of the local spatial permutation test of summed probability distributions.</i>
-------------------------------	--

---

### Description

Displays local growth rates, p-values, and q-values retrieved from a `spatialTest` class object.

### Usage

```

## S3 method for class 'spatialTest'
plot(
  x,
  index = 1,
  option,
  breakRange = NA,
  breakLength = 7,
  rd = 5,
  baseSize = 0.5,
  plim = 0.05,
  qlim = 0.05,
  legend = FALSE,
  legSize = 1,
  location = "bottomright",
  ...
)

```

**Arguments**

x	A spatialTest class object
index	A numerical value indicating which transition to display. Ignored when option="rawLegend" or option="testLegend". Default is 1.
option	Indicates what to display. Either "raw" (the local growth rate) or "test" (the test results, i.e. q and p values).
breakRange	A vector of length 2 defining the minimum and maximum values of growth rate to be displayed in the legend. If set to NA its computed from data range (default).
breakLength	A numerical vector defining the number of breaks for growth rates to be displayed in the legend.
rd	Number of decimal places of the growth rate to be displayed in the Legend
baseSize	Numerical value giving the amount by which points should be magnified relative to the default settings in R. Default is 0.5
plim	Threshold value for the p-values. Default is 0.05.
qlim	Threshold value for the q-values. Default is 0.05.
legend	Logical values specifying whether the legend should be displayed or not. Default is FALSE.
legSize	Numerical value giving the amount by which points should be magnified relative to the default settings in R for the Legend. Default is 1.
location	A single keyword from the list "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright", "right" and "center" to specify the location of the Legend. Default is "bottomright".
...	Graphical parameters to be passed to methods.

**Details**

The function displays a distribution map of local growth rates (when option="raw"), q- and p-values (when option="test"), and the associated legends (when option="rawLegend" or option="testLegend").

**See Also**

[sptest](#)

---

plot.SpdModelTest	<i>Plot result of Monte-Carlo simulation of observed versus modelled SPDs</i>
-------------------	---

---

**Description**

The function visualises the observed summed probability distribution of radiocarbon dates along with a simulation envelope for the null model and regions of positive and negative deviation.

**Usage**

```
## S3 method for class 'SpdModelTest'
plot(
  x,
  calendar = "BP",
  type = "spd",
  ylim = NA,
  xlim = NA,
  col.obs = "black",
  lwd.obs = 0.5,
  xaxs = "i",
  yaxs = "i",
  bbtty = "f",
  bbttyRet = TRUE,
  drawaxes = TRUE,
  ...
)
```

**Arguments**

<code>x</code>	A <code>SpdModelTest</code> class object generated using the <code>modelTest</code> function.
<code>calendar</code>	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
<code>type</code>	Whether to display SPDs ('spd') or rates of change ('roc'). Default is 'spd'.
<code>ylim</code>	the y limits of the plot.
<code>xlim</code>	the x limits of the plot. In BP or in BC/AD depending on the choice of the parameter <code>calendar</code> . Notice that if BC/AD is selected BC ages should have a minus sign (e.g. <code>c(-5000, 200)</code> for 5000 BC to 200 AD).
<code>col.obs</code>	Line colour for the observed SPD.
<code>lwd.obs</code>	Line width for the observed SPD.
<code>xaxs</code>	The style of x-axis interval calculation (see <a href="#">par</a> )
<code>yaxs</code>	The style of y-axis interval calculation (see <a href="#">par</a> )
<code>bbtty</code>	Display options; one between 'b', 'n', and 'f'. See details below.
<code>bbttyRet</code>	Whether details on the intervals of positive and negative deviations are returned. Default is TRUE.
<code>drawaxes</code>	A logical value determining whether the axes should be displayed or not. Default is TRUE.
<code>...</code>	Additional arguments affecting the plot

**Details**

The argument `bbtty` controls the display options of the Monte-Carlo Test. Default settings (`bbtty='f'`) displays the observed SPD (solid black line), the simulation envelope of the fitted model (shaded grey polygon) and regions of significance positive (red semi-transparent rectangle) and negative



(blue semi-transparent rectangle) deviation. The option `bbty='b'` removes the regions of positive/negative deviations, whilst the option `bbty='n'` displays the simulation envelope on existing plot.

### See Also

[modelTest](#)

---

plot.SpdPermTest      *Plot result of mark permutation test of SPDs*

---

### Description

Visualises the observed SPD along with the simulation envelope generated from [permTest](#), with regions of positive and negative deviations highlighted in red and blue.

### Usage

```
## S3 method for class 'SpdPermTest'
plot(
  x,
  focalm = 1,
  type = "spd",
  calendar = "BP",
  xlim = NA,
  ylim = NA,
  col.obs = "black",
  col.env = rgb(0, 0, 0, 0.2),
  lwd.obs = 0.5,
  xaxs = "i",
  yaxs = "i",
  bbty = "f",
  drawaxes = TRUE,
  ...
)
```

### Arguments

<code>x</code>	A <code>SpdPermTest</code> class object. Result of random mark permutation test (see <a href="#">permTest</a> )
<code>focalm</code>	Value specifying the name of the focal mark (group) to be plotted.
<code>type</code>	Whether to display SPDs ('spd') or rates of change ('roc'). Default is 'spd'.
<code>calendar</code>	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
<code>xlim</code>	the x limits of the plot. In BP or in BC/AD depending on the choice of the parameter calendar. Notice that if BC/AD is selected BC ages should have a minus sign (e.g. <code>c(-5000, 200)</code> for 5000 BC to 200 AD).

ylim	the y limits of the plot.
col.obs	Line colour for the observed SPD
col.env	Colour for the simulation envelope
lwd.obs	Line width for the observed SPD
xaxs	The style of x-axis interval calculation (see <a href="#">par</a> )
yaxs	The style of y-axis interval calculation (see <a href="#">par</a> )
bbty	Display options; one between 'b', 'n', and 'f'. See details in <a href="#">plot.SpdModelTest</a> .
drawaxes	A logical value determining whether the axes should be displayed or not. Default is TRUE.
...	Additional arguments affecting the plot

**See Also**

[permTest](#); [plot.SpdModelTest](#);

---

plot.spdRC	<i>Plot spdRC class objects</i>
------------	---------------------------------

---

**Description**

Plot rates of change between time-blocks

**Usage**

```
## S3 method for class 'spdRC'
plot(
  x,
  calendar = "BP",
  col.obs = "black",
  lwd.obs = 0.5,
  xaxs = "i",
  yaxs = "i",
  xlim = NA,
  ...
)
```

**Arguments**

x	spdRC class object containing geometric growth rates.
calendar	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
col.obs	Line colour for the observed SPD
lwd.obs	Line width for the observed SPD
xaxs	The style of x-axis interval calculation (see <a href="#">par</a> )

yaxs            The style of y-axis interval calculation (see [par](#))  
 xlim            Limits for the x axis.  
 ...             Additional arguments affecting the plot.

### See Also

[spd2rc](#)

---

plot.stackCalSPD            *Plot a stack of SPDs*

---

### Description

Visualises multiple SPDs grouped as a stackCalSPD object.

### Usage

```

## S3 method for class 'stackCalSPD'
plot(
  x,
  type = "stacked",
  calendar = "BP",
  spdnormalised = FALSE,
  rescale = FALSE,
  runm = NA,
  xlim = NA,
  ylim = NA,
  xaxt = "s",
  yaxt = "s",
  gapFactor = 0.2,
  col.fill = NA,
  col.line = NA,
  lwd.obs = 1,
  lty.obs = 1,
  cex.lab = 1,
  cex.axis = cex.lab,
  legend = TRUE,
  legend.arg = NULL,
  ylab = NA,
  ymargin = 1.1,
  ...
)

```

**Arguments**

<code>x</code>	A <code>stackCalSPD</code> class object. Result of <code>stackspd</code> function.
<code>type</code>	How to display the SPDs. Current options are 'stacked', 'lines', 'proportion', and 'multipanel'. Default is 'stacked'.
<code>calendar</code>	Either 'BP' or 'BCAD'. Indicate whether the calibrated date should be displayed in BP or BC/AD. Default is 'BP'.
<code>spdnormalised</code>	A logical variable indicating whether the total probability mass of the SPDs are normalised to sum to unity. Default is FALSE.
<code>rescale</code>	A logical variable indicating whether the SPDs should be rescaled to range 0 to 1. Default is FALSE.
<code>runm</code>	A number indicating the window size of the moving average to smooth the SPD. If set to NA no moving average is applied. Default is NA
<code>xlim</code>	the x limits of the plot. In BP or in BC/AD depending on the choice of the parameter <code>calendar</code> . Notice that if BC/AD is selected BC ages should have a minus sign (e.g. <code>c(-5000, 200)</code> for 5000 BC to 200 AD).
<code>ylim</code>	the y limits of the plot.
<code>xaxt</code>	Whether the x-axis tick marks should be displayed ( <code>xaxt='s'</code> , default) or not ( <code>xaxt='n'</code> ).
<code>yaxt</code>	Whether the y-axis tick marks should be displayed ( <code>yaxt='s'</code> , default) or not ( <code>yaxt='n'</code> ).
<code>gapFactor</code>	Defines spacing between SPDs as proportion of the y-axis range for multipanel plots. Default is 0.2.
<code>col.fill</code>	Vector of fill color for the observed SPDs. The default color scheme is based on the Dark2 palette of RColorBrewer package.
<code>col.line</code>	Line colour for the observed SPDs. The default color scheme is based on the Dark2 palette of RColorBrewer package.
<code>lwd.obs</code>	Line width for the observed SPDs. Default is 1.
<code>lty.obs</code>	Line type for the observed SPDs. Default is 1.
<code>cex.lab</code>	The magnification to be used for x and y labels relative to the current setting of <code>cex</code> . Default is adjusted to 1.
<code>cex.axis</code>	The magnification to be used for axis annotation relative to the current setting of <code>cex</code> . Default is adjusted to 1.
<code>legend</code>	Whether legend needs to be displayed. Item names will be retrieved from the values supplied in the argument group in <code>stackspd</code> . Default is TRUE.
<code>legend.arg</code>	list of additional arguments to pass to <code>legend</code> ; names of the list are used as argument names. Only used if <code>legend</code> is set to TRUE.
<code>ylab</code>	a title for the y axis
<code>ymargin</code>	multiplier for the maximum value on <code>ylim</code> range. Default is 1.1.
<code>...</code>	Additional arguments affecting the plot.

## References

Erich Neuwirth (2014). RColorBrewer: ColorBrewer Palettes. R package version 1.1-2. <https://CRAN.R-project.org/package=RColorBrewer>.

## Examples

```
## Not run:
data(emedyd)
emedyd = subset(emedyd,Region==1)
x = calibrate(x=emedyd$CRA, errors=emedyd$Error,normalised=FALSE)
bins = binPrep(sites=emedyd$SiteName, ages=emedyd$CRA,h=50)
res = stackspd(x=x,timeRange=c(16000,8000),bins=bins,group=emedyd$Region)
plot(res,type='stacked')
plot(res,type='lines')
plot(res,type='proportion')
plot(res,type='multipanel')

## End(Not run)
```

---

plot.stKde

*Plot map(s) of the spatio-temporal intensity of radiocarbon dates.*

---

## Description

Plotting function for single or multiple maps of the spatio-temporal intensity of radiocarbon dates at given focal years.

## Usage

```
## S3 method for class 'stKde'
plot(
  x,
  focalyears = NULL,
  type = "focal",
  plotdir = NULL,
  imnames = "byyear",
  zlim = NULL,
  box = FALSE,
  main = "auto",
  col.mask = "grey75",
  ncolours = 256,
  ramptype = "raw",
  imdim = c(10, 10),
  mars = c(0.5, 0.5, 2.5, 2),
  ribbon = TRUE,
  ribargs = list(),
  cex.ribbon = 0.5,
  withpts = "n",
```

```

    cex.main = 0.8,
    pch.pts = 19,
    col.pts = "grey50",
    cex.pts = 0.1,
    tidydevs = TRUE,
    verbose = TRUE,
    ...
)

```

### Arguments

x	An object of class stKde.
focalyears	A vector of numeric values for focal years, in calBP, that will be timesteps at which date intensity maps will be plotted.
type	A single character string stipulating which type of plot to create. Current options are "nonfocal", "mask", "focal", "proportion", "change" and "all".
plotdir	Optional output directory for plots. If NULL, then only a single plot is made to the current device. If a valid output directory is provided then one or more timeslices maps are saved as png files (e.g. as source images for an animation).
imnames	The format of the output files if output is as png files to a directory. The current two options are "basic" (labelling the images in basic sequence as preferred by animation software such as ffmpeg) or "byyear" (labelling the images by calBP year).
zlim	Numeric vector of length=2 which controls the maximum or minimum of the colour ramp.
box	Logical. Plot a border around the map or not.
main	Single character string specifying a main title. "auto" implies internal default titles are used.
col.mask	The colour used to depict any areas that are being masked out.
ncolours	The maximum number of colours to use in the colour ramp.
ramptype	What kind of treatment for the colour ramp. Current options are "raw" (do not try to standardise the ramps across timeslices), "std" (standardise each plot, typically by capturing the first 3sd in the main colour ramp and then outliers beyond that with the extreme colours of the ramp), "unl" (do not standardise but plot generalised high/low ramp labels) and "mmx" (scale to the minimum and maximum values across all timeslices).
indim	Height and width of the plot to png in cm.
mars	Margins of the plot to png.
ribbon	Whether to plot the colour ramp legend or not.
ribargs	Whether to plot the colour ramp legend or not.
cex.ribbon	The size of the ribbon font.
withpts	Plot with the original sample locations shown (current options are "y" and "n").
cex.main	The size of the title font.

pch.pts	The symbols used for the plotted points.
col.pts	The colours used for the plotted points.
cex.pts	The size used for the plotted points.
tidydevs	Logical for whether to clean up any open graphics devices or not (default is TRUE).
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.
...	Additional arguments affecting the plot.

### Details

This function plots to a screen device if a single focal year is stipulated and no output directory. Or if an output directory is stipulated, it plots one or more focal years to png files, with some basic formatting options and optional cross-year standardisation of the colour ramps (with a view to them being stills-for-video). For even more control of plotting, call this function one year at a time in a loop.

### Examples

```
## Not run:
## Example with a subset of English and Welsh dates from the Euroevol dataset
data(ewdates)
data(ewowin)
dir.create(file.path("im"), showWarnings=FALSE)
x <- calibrate(x=ewdates$C14Age, errors=ewdates$C14SD, normalised=FALSE)
bins1 <- binPrep(sites=ewdates$SiteID, ages=ewdates$C14Age, h=50)
stkde1 <- stkde(x=x, coords=ewdates[,c("Eastings", "Northings")],
win=ewowin, sbw=40000, cellres=2000, focalyears=seq(6500, 5000, -100),
tbw=50, bins=bins1, backsight=200, outdir="im")

## Plot just the proportion surface for just 5900 calBP
plot(stkde1, 5900, type="proportion")

## Plot an example of all four basic outputs for just 5900 calBP
dev.new(height=2.5, width=8)
par(mar=c(0.5, 0.5, 2.5, 2))
plot(stkde1, 5900, type="all")

## Plot standardised change surfaces to a sub-directory called
## /png for all timeslices and save to file.
dir.create(file.path("png"), showWarnings=FALSE)
plot(stkde1, seq(6500,5000,-100), type="change", ramptype="std", withpts=TRUE, plotdir="png")

## Plot all four summary surfaces in one image, saving them to a sub-directory call 'pngall',
## and with the output of the change map standardised to a common ramp
## (but leaving the focal and proportion maps unstandardised with simple ramp labelling)
dir.create(file.path("pngall"), showWarnings=FALSE)
plot(stkde1, seq(6500,5000,-100), type="all", ramptype=c("unl","unl","std"), imdim=cm(c(2.5,8)),
withpts=TRUE, plotdir="pngall")
```

```
## End(Not run)
```

---

rcarbon

*rcarbon: Calibration and analysis of radiocarbon dates*

---

## Description

The **rcarbon** package handles the calibration and analysis of radiocarbon, often but not exclusively for the purposes of archaeological research. It includes functions not only for basic calibration, uncalibration and plotting of one or more dates, but also a statistical framework for building demographic and related longitudinal inferences from aggregate radiocarbon date lists.

## Details

Core functions in the **rcarbon** package can be grouped as follows:

**Calibration Functions** `calibrate` and `uncalibrate` enable the calibration and back-calibration for a variety of curves.

**Aggregation Functions** `spd` generates a summed probability distribution (SPD) of radiocarbon dates; `binPrep` can be used to define clusters of radiocarbon dates associated with the same context/phase

**Statistical Test Functions** `modelTest` compares the observed SPD against a variety of theoretical models (most typically an exponential curve) using the Monte-Carlo approach; `p2pTest` compares observed differences in SPD between two user-specified points in time against differences expected from a theoretical model; `permTest` compares two or more SPDs and test for the null hypothesis that all sets are derived from the same population; `SPpermTest` identifies, for defined intervals, locations with significantly higher or lower growth rate in the SPD compared to the pan-regional trend in the data

## Note

Up-to-date development version, bug-reports, and further information concerning the **rcarbon** package can be found on GitHub (<https://github.com/ahb108/rcarbon>). To see the preferred citation for the package, type `citation("rcarbon")`.

## Author(s)

The **rcarbon** is developed and maintained by Andrew Bevan and Enrico Crema

## References

See individual functions for references.



---

reScale	<i>Rescale a numeric vector to a specified minimum and maximum</i>
---------	--

---

**Description**

Rescale a numeric vector to a specified minimum and maximum.

**Usage**

```
reScale(x, type = "simple", to = c(0, 1), na.rm = TRUE)
```

**Arguments**

x	numeric vector to smooth.
type	what kind of rescaling to perform. Current options are 'simple' (default) and 'normal' which produces a z-score and 'custom' for which the 'to' argument must be specified.
to	numeric vector of length 2 specifying the minimum and maximum value to perform a linear rescale between (default is 0 and 1)
na.rm	Set to TRUE, this removes NAs before rescaling.

**Value**

A numeric vector of rescaled values.

**Examples**

```
reScale(15:200)
```

---

runMean	<i>Calculate a running mean from a numeric vector.</i>
---------	--

---

**Description**

Calculate a running mean from a numeric vector.

**Usage**

```
runMean(x, n, edge = "NA")
```

**Arguments**

x	numeric vector to smooth.
n	the size of the window in which to smooth.
edge	How to treat edge cases where a full window is unavailable. Current options are 'NA' to fill with NAs or 'fill' to fill with original values

**Value**

A numeric vector of smoothed values.

**Examples**

```
x <- rnorm(1000)
y <- c(1:1000)
plot(y,x, type="l")
lines(runMean(x,50), col="red")
```

---

sampleDates	<i>Sample random calendar dates</i>
-------------	-------------------------------------

---

**Description**

Randomly samples calendar dates from each calibrated date or bin.

**Usage**

```
sampleDates(x, bins = NA, nsim, boot = FALSE, verbose = TRUE)
```

**Arguments**

x	A 'CalDates' class object.
bins	A vector containing the bin names associated with each radiocarbon date. If set to NA, binning is not carried out.
nsim	Number of sampling repetitions.
boot	A logical value indicating whether bootstrapping is carried out (see details below). Default is FALSE.
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.

**Details**

The function randomly samples calendar dates based from calibrated probability distributions. When the bins argument is supplied a single calendar date is sampled from each bin. When boot=TRUE, dates (or bins) are randomly sampled with replacement before calendar dates are sampled.

**Value**

An object of class simdates with the following elements

- sdates A matrix containing the randomly sampled calendar dates, with rows containing each of the nsim repetitions.
- weight A vector (or matrix in when boot=TRUE) containing the total area under the curve of each date, normalised to sum to unity. Notice this will be identical for all dates if the calibration is carried out with the argument normalised set to TRUE.

---

`smoothGauss`*Smooth a numeric vector using a Gaussian window*

---

**Description**

Smooth a numeric vector using a Gaussian window

**Usage**

```
smoothGauss(x, alpha, window = 0.1)
```

**Arguments**

<code>x</code>	numeric vector of values to smooth.
<code>alpha</code>	numeric value controlling the size of the gaussian smoothing window. Proportional to the standard deviation of the Gaussian smoothing kernel where $sd=(N-1)/(2*\alpha)$ with N being the length of the input vector.
<code>window</code>	a fraction between 0 and 1 representing the proportion of the input vector to include in the moving window.

**Details**

Adapted from `smth.gaussian` in the `smoother` package.

**References**

Hamilton, N. (2015). `smoother`: Functions Relating to the Smoothing of Numerical Data, R package version 1.1, <https://CRAN.R-project.org/package=smoother>

**Examples**

```
smoothGauss(runif(200), alpha=5)
```

---

`spd`*Summed probability distributions (SPD) of radiocarbon dates.*

---

**Description**

The function generates Summed probability distributions (SPD) of radiocarbon dates, with optional binning routine for controlling inter-site or inter-phase variation in sampling intensity.

**Usage**

```
spd(
  x,
  timeRange,
  bins = NA,
  datenormalised = FALSE,
  spdnormalised = FALSE,
  runm = NA,
  verbose = TRUE,
  edgeSize = 500
)
```

**Arguments**

x	A CalDates class object containing the calibrated radiocarbon dates.
timeRange	A vector of length 2 indicating the start and end date of the analysis in cal BP.
bins	A vector containing the bin names associated with each radiocarbon date. If set to NA, binning is not carried out.
datenormalised	Controls for calibrated dates with probability mass outside the timerange of analysis. If set to TRUE the total probability mass within the time-span of analysis is normalised to sum to unity. Should be set to FALSE when the parameter normalised in <a href="#">calibrate</a> is set to FALSE. Default is FALSE.
spdnormalised	A logical variable indicating whether the total probability mass of the SPD is normalised to sum to unity.
runm	A number indicating the window size of the moving average to smooth the SPD. If set to NA no moving average is applied. Default is NA
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.
edgeSize	Extra margin in C14 Age time to handle edge effect when datenormalise is set to TRUE. Default is 500.

**Details**

The binning routine consists of computing summed probability distribution of all dates associated to a given bin, divided by the number of contributing dates. This controls for any striking differences in sampling intensity, and ensures that each site phase is equally contributing to the final SPD (see Timpson et al 2014 for details). Bins can be generated using the [binPrep](#), whilst the sensitivity to parameter choice can be explored with the [binsense](#) function.

**Value**

An object of class CalSPD with the following elements

- metadata A data.frame containing relevant information regarding the parameters used to create the SPD as well as sample size and number of bins
- grid A CalGrid class object containing the summed probability associated to each calendar year between timeRange[1] and timeRange[2]

**References**

Timpson, A., et al, (2014). Reconstructing regional population fluctuations in the European Neolithic using radiocarbon dates: a new case-study using an improved method. *Journal of Archaeological Science* 52: 549-557. DOI:10.1016/j.jas.2014.08.011

**See Also**

[calibrate](#) for calibrating radiocarbon dates; [binPrep](#) for preparing bins.

---

 spd2rc

---

*Computes rates of change from SPDs*


---

**Description**

Function for computing rates of change between abutting user-defined time-blocks.

**Usage**

```
spd2rc(
  spd,
  breaks = NULL,
  backsight = NULL,
  changexpr = expression((t1/t0)^(1/d) - 1)
)
```

**Arguments**

spd	Summed Probability Distribution obtained using the <a href="#">spd</a> function.
breaks	A vector giving the breakpoints between the time-blocks.
backsight	A single numeric value defining the distance in time between the focal year and the backsight year for computing the rate of change.
changexpr	An expression defining how the rate of change is calculated, where $t_1$ is the summed probability for a focal block or year, $t_0$ is the summed probability for previous block or backsight year, and $d$ is the duration of the block or the length of the backsight. Default is a geometric growth rate (i.e $\text{expression}((t_2/t_1)^{(1/d)}-1)$ ).

**Details**

When the argument `breaks` is supplied the function aggregates the summed probability within each time-block and compared them across abutting blocks using the expression defined by `changexpr`. When the argument `backsight` is provided he expression is based on the comparison between the summed probability of each year and the associated backsight year.

**Value**

An object of class `spdRC`.

**Examples**

```
## Not run:
data(emedyd)
caldates <- calibrate(x=emedyd$CRA, errors=emedyd$Error, normalised=FALSE, calMatrix=TRUE)
bins <- binPrep(sites=emedyd$SiteName, ages=emedyd$CRA, h=50)
emedyd.spd <- spd(caldates,bins,timeRange=c(16000,9000),runm=100)
emedyd.gg <- spd2rc(emedyd.spd,breaks=seq(16000,9000,-1000))
emedyd.gg2 <- spd2rc(emedyd.spd,backsight=10)
plot(emedyd.gg)
plot(emedyd.gg2)

## End(Not run)
```

---

spkde	<i>Map the spatial intensity of a set of radiocarbon dates for a given focal year.</i>
-------	--

---

**Description**

Function for mapping the spatial intensity of radiocarbon dates for a given geographical region and focal year.

**Usage**

```
spkde(
  x,
  coords,
  sbw,
  focyear,
  tbw,
  win,
  cellres,
  bins = NA,
  backsight = NA,
  nsim = NULL,
  maskthresh = 0,
  changexpr = expression((t1 - t0)/tk),
  raw = FALSE,
  spjitter = TRUE,
  amount = NULL,
  verbose = TRUE,
  ...
)
```

**Arguments**

x An object of class CalDates with calibrated radiocarbon ages.

coords	A two column matrix of geographical coordinates from a projected coordinate system (no checks are made for this) and with the same number of rows as length(x).
sbw	A single numeric value for the spatial bandwidth to be applied around each raster cell, expressed as the standard deviation of a continuous Gaussian kernel (passed as the sigma argument to density.ppp()).
focalyear	A single numeric value for the focal year for the intensity map.
tbw	A single numeric value for the temporal bandwidth to be applied around each focal year, expressed as the standard deviation of a continuous Gaussian kernel.
win	The bounding polygon for the mapping (must be an object of class 'owin', see the spatstat package)
cellres	The cell or pixel resolution of the output raster maps.
bins	A vector of labels corresponding to site names, ids, bins or phases (same length as x)
backsight	A single numeric value (which will be coerced to be positive) that specifies a comparison timestep in the past for a mapping of temporal change.
nsim	How many bootstrap simulations to run (default is none).
maskthresh	A single numeric value for a lower-bound cut-off for all maps, based on a minimum required spatial intensity of all dates in x.
changexpr	An expression for calculating the change in spatial intensity between the focal year and a backsight year (as defined via the backsight argument). Available input options are t1 (the spatial intensity for the focal year), t0 (the spatial intensity for the backsight year) and tk (the overall spatial intensity for all dates irrespective of year), plus any other standard constants and mathematical operators. A sensible default is provided.
raw	Whether to output the raw simulations (if nsim is set) or just the summaries (the latter is default).
spjitter	Whether noise is applied to the spatial coordinates or not. Default is TRUE.
amount	Amount of jitter applied to the spatial coordinates when spjitter=TRUE. Default is d/5, where d is difference between the closest coordinates.
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.
...	ignored or passed to internal functions.

### Details

This function is not really intended for general use, but rather as an internal function for `stkde()`. Most applications should use the latter, but `spkde` has been exported and made externally available, both because this function retains the result in memory (in contrast to `stkde`) and with a view to possible addition of bootstrap methods in the future. Some function arguments therefore remain experimental. The function computes timeslice maps of the spatio-temporal kernel intensity of radiocarbon dates across a geographic region for a specific focal year. The user specifies the arbitrary size of both the spatial and the temporal Gaussian kernels that will be used to summarise radiocarbon date intensity per grid cell.

**Value**

A list object of class spKde with the following elements:

- A series of list items storing some of the input parameters such as the focal year, sbw, tbw, backsight, maskthresh.
- nonfocal An im object mapping the basic spatial intensity of all dates, without reference to a focal year.
- focal An im object mapping the spatial intensity of dates for the focal year (i.e. weighted by how much each dates probability distribution overlaps with a Gaussian kernel centred on the focal year with a standard deviation of tbw).
- proportion An im object mapping the proportional intensity of dates for the focal year (i.e. the focal surface divided by the nonfocal surface).
- change An im object mapping the amount of change between the intensity of dates for the focal year and a chosen backsight year (i.e. as defined by changexpr).

**Examples**

```
## Not run:
## Example for the focal year 5600 calBP (also with site binning),
## using a subset of English and Welsh dates from the Euroevol dataset
data(ewdates)
data(ewowin)
x <- calibrate(x=ewdates$C14Age, errors=ewdates$C14SD, normalised=FALSE)
bins1 <- binPrep(sites=ewdates$SiteID, ages=ewdates$C14Age, h=50)
spkde1 <- spkde(x=x, coords=ewdates[,c("Eastings", "Northings")], win=ewowin,
sbw=40000, cellres=2000, focyear=5600, tbw=50, bins=bins1, backsight=200, amount=1)
plot(spkde1$focal)
plot(spkde1$proportion)

## End(Not run)
```

---

sptest

*Spatial Permutation Test of summed probability distributions.*


---

**Description**

This function carries out local spatial permutation tests of summed radiocarbon probability distributions in order to detect local deviations in growth rates (Crema et al 2017).

**Usage**

```
sptest(
  calDates,
  timeRange,
  bins,
  locations,
```



```

breaks,
spatialweights,
rate = expression((t2/t1)^(1/d) - 1),
nsim = 1000,
runm = NA,
permute = "locations",
ncores = 1,
datenormalised = FALSE,
verbose = TRUE,
raw = FALSE
)

```

### Arguments

calDates	A CalDates class object.
timeRange	A vector of length 2 indicating the start and end date of the analysis in cal BP
bins	A vector indicating which bin each radiocarbon date is assigned to. Must have the same length as the number of radiocarbon dates. Can be created using the <a href="#">binPrep</a> function. Bin names should follow the format "x_y", where x refers to a unique location (e.g. a site) and y is a integer value (e.g. "S023_1", "S023_2", "S034_1", etc.).
locations	A SpatialPoints or a SpatialPointsDataFrame class object. Rownames of each point should much the first part of the bin names supplied (e.g. "S023", "S034")
breaks	A vector of break points for defining the temporal slices.
spatialweights	A spatialweights class object defining the spatial weights between the locations (cf. <a href="#">spweights</a> )
rate	An expression defining how the rate of change is calculated, where t1 is the summed probability for a focal block, t2 is the summed probability for next block, and d is the duration of the blocks. Default is a geometric growth rate (i.e. $\text{expression}((t2/t1)^{(1/d)-1})$ ).
nsim	The total number of simulations. Default is 1000.
runm	The window size of the moving window average. Must be set to NA if the rates of change are calculated from the raw SPDs.
permute	Indicates whether the permutations should be based on the "bins" or the "locations". Default is "locations".
ncores	Number of cores used for for parallel execution. Default is 1.
datenormalised	A logical variable indicating whether the probability mass of each date within timeRange is equal to 1. Default is FALSE.
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.
raw	A logical variable indicating whether permuted sets of geometric growth rates for each location should be returned. Default is FALSE.

## Details

The function consists of the following seven steps: 1) for each location (e.g. a site) generate a local SPD of radiocarbon dates, weighting the contribution of dates from neighbouring sites using a weight scheme provided by the `spatialweights` class object; 2) define temporal slices (using breaks as break values), then compute the total probability mass within each slice; 3) compute the rate of change between abutting temporal slices by using the formula:  $(SPD_t/SPD_{t+1}^{1/\Delta t} - 1)$ ; 4) randomise the location of individual bins or the entire sequence of bins associated with a given location and carry out steps 1 to 3; 5) repeat step 4 `nsim` times and generate, for each location, a distribution of growth rates under the null hypothesis (i.e. spatial independence); 6) compare, for each location, the observed growth rate with the distribution under the null hypothesis and compute the p-values; and 7) compute the false-discovery rate for each location.

## Value

A `spatialTest` class object

## References

Crema, E.R., Bevan, A., Shennan, S. (2017). Spatio-temporal approaches to archaeological radiocarbon dates. *Journal of Archaeological Science*, 87, 1-9.

## See Also

`permTest` for a non-spatial permutation test; `plot.spatialTest` for plotting; `spweights` for computing spatial weights; `spd2rc` for computing geometric growth rates.

## Examples

```
## Reproduce Crema et al 2017 ##
## Not run:
data(euroevol) #load data

## Subset only for 8000 to 5000 Cal BP (c7200-4200 C14BP)
edge=800
timeRange=c(8000,5000)
euroevol2=subset(euroevol,C14Age<=c(timeRange[1]-edge)&C14Age>=c(timeRange[2]-edge))

## define chronological breaks
breaks=seq(8000,5000,-500)

## Create a SpatialPoints class object
library(sp)
sites = unique(data.frame(SiteID=euroevol2$SiteID,
Longitude=euroevol2$Longitude,Latitude=euroevol2$Latitude))
locations=data.frame(Longitude=sites$Longitude,Latitude=sites$Latitude)
rownames(locations)=sites$SiteID
coordinates(locations)<-c("Longitude","Latitude")
proj4string(locations)<- CRS("+proj=longlat +datum=WGS84")

## Compute Distance and Spatial Weights
distSamples=spDists(locations,locations,longlat = TRUE)
```

```

spatialweights=spweights(distSamples,h=100) #using a kernal bandwidth of 100km

## Calibration and binning
bins=binPrep(sites=euroevol2$SiteID,ages=euroevol2$C14Age,h=200)
calDates=calibrate(x=euroevol2$C14Age,errors=euroevol2$C14SD,
timeRange=timeRange,normalised=FALSE)

## Main Analysis (over 2 cores; requires doParallel package)
## NOTE: the number of simulations should be ideally larger
## to ensure a better resolution of the p/q-values.
res.locations=sptest(calDates,timeRange=timeRange,bins=bins,locations=locations,
spatialweights=spatialweights,breaks=breaks,ncores=2,nsim=100,
permute="locations",datenormalised=FALSE)

## Plot results
library(rworldmap)
base=getMap(resolution="low") #optionally add base map
#retrieve coordinate limits#
xrange=bbbox(res.locations$locations)[1,]
yrange=bbbox(res.locations$locations)[2,]

par(mfrow=c(2,2))
par(mar=c(0.1,0.1,0,0.5))
plot(base,col="antiquewhite3",border="antiquewhite3",xlim=xrange,ylim=yrange)
plot(res.locations,index=4,add=TRUE,legend=TRUE,option="raw",breakRange=c(-0.005,0.005))
par(mar=c(0.1,0.1,0,0.5))
plot(base,col="antiquewhite3",border="antiquewhite3",xlim=xrange,ylim=yrange)
plot(res.locations,index=4,add=TRUE,legend=TRUE,option="test")

## End(Not run)

```

---

spweights

*Compute weights from distance matrix*


---

### Description

Function for computing a matrix of gaussian or fixed weights from distance matrix

### Usage

```
spweights(distmat, h = NULL, kernel = "gaussian")
```

### Arguments

distmat	a symmetric matrix of inter-site distances (in km).
h	parameter of the Gaussian distance decay function.
kernel	indicates the type of weighting function, either 'fixed' or 'gaussian'. Default is 'gaussian'.

**Details**

This function generates a weight matrix (required for the `SPpermTest`) function. When `kernel=="fixed"`, the weight  $w_{ij}$  between site  $i$  and  $j$  is equal to 1 when their interdistance  $d_{ij}$  is below  $h$ , and equal to 0 when  $d_{ij} > h$ . When `kernel=="gaussian"`, the weight is calculated with formula  $\exp(-d_{ij}^2/h^2)$ .

**Value**

An object of class `spatialweights`

**Examples**

```
lon <- c(11.3426,0.1278,0.1218)
lat <- c(44.4949,51.5074,52.2053)
library(sp)
d <- spDists(x=cbind(lon,lat),y=cbind(lon,lat))
spweights(d,h=100)
spweights(d,h=100,kernel="fixed")
```

---

stackspd

*Stacked Summed Probability Distribution*


---

**Description**

Generates and combines multiple SPDs based on a user defined grouping.

**Usage**

```
stackspd(
  x,
  timeRange,
  bins = NA,
  group = NULL,
  datenormalised = FALSE,
  runm = NA,
  verbose = TRUE,
  edgeSize = 500
)
```

**Arguments**

<code>x</code>	A <code>CalDates</code> class object containing the calibrated radiocarbon dates.
<code>timeRange</code>	A vector of length 2 indicating the start and end date of the analysis in cal BP.
<code>bins</code>	A vector containing the bin names associated with each radiocarbon date. If set to <code>NA</code> , binning is not carried out.
<code>group</code>	A vector containing the grouping variable.

datenormalised	Controls for calibrated dates with probability mass outside the timerange of analysis. If set to TRUE the total probability mass within the time-span of analysis is normalised to sum to unity. Should be set to FALSE when the parameter normalised in <code>calibrate</code> is set to FALSE. Default is FALSE.
runm	A number indicating the window size of the moving average to smooth the SPD. If set to NA no moving average is applied. Default is NA
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.
edgeSize	Controls edge effect by expanding the fitted model beyond the range defined by <code>timeRange</code> .

### Value

An object of class `stackCalSPD`

### Examples

```
## Not run:
data(emedyd)
x = calibrate(x=emedyd$CRA, errors=emedyd$Error,normalised=FALSE)
bins = binPrep(sites=emedyd$SiteName, ages=emedyd$CRA,h=50)
res = stackspd(x=x,timeRange=c(16000,8000),bins=bins,group=emedyd$Region)

## End(Not run)
```

---

stkde

*Map the spatio-temporal intensity of a set of radiocarbon dates*

---

### Description

Function for mapping the spatio-temporal intensity of radiocarbon dates for a given geographical region in one or more tim.

### Usage

```
stkde(
  x,
  coords,
  sbw,
  focalyears,
  tbw,
  win,
  cellres,
  outdir = ".",
  bins = NA,
  backsight = NA,
  maskthresh = 0,
```

```

    changexpr = expression((t1 - t0)/tk),
    spjitter = TRUE,
    amount = NULL,
    verbose = TRUE,
    ...
)

```

### Arguments

x	An object of class CalDates with calibrated radiocarbon ages.
coords	A two column matrix of geographical coordinates from a projected coordinate system (no checks are made for this) and with the same number of rows as length(x).
sbw	A single numeric value for the spatial bandwidth to be applied around each raster cell, expressed as the standard deviation of a continuous Gaussian kernel (passed as the sigma argument to density.ppp()).
focalyears	A vector of numeric values for focal years, in calBP, that will be timesteps at which date intensity maps will be produced.
tbw	A single numeric value for the temporal bandwidth to be applied around each focal year, expressed as the standard deviation of a continuous Gaussian kernel.
win	The bounding polygon for the mapping (must be an object of class 'owin', see the spatstat package)
cellres	The cell or pixel resolution of the output raster maps.
outdir	The output directory for timeslice maps and data that are saved to file.
bins	A vector of labels corresponding to site names, ids, bins or phases (same length as x)
backsight	A single numeric value (which will be coerced to be positive) that specifies a comparison timestep in the past for a mapping of temporal change.
maskthresh	A single numeric value for a lower-bound cut-off for all maps, based on a minimum required spatial intensity of all dates in x.
changexpr	An expression for calculating the change in spatial intensity between the focal year and a backsight year (as defined via the backsight argument). Available input options are t1 (the spatial intensity for the focal year), t0 (the spatial intensity for the backsight year) and tk (the overall spatial intensity for all dates irrespective of year), plus any other standard constants and mathematical operators. A sensible default is provided.
spjitter	Whether noise is applied to the spatial coordinates or not. Default is TRUE.
amount	Amount of jitter applied to the spatial coordinates when spjitter=TRUE. Default is d/5, where d is difference between the closest coordinates.
verbose	A logical variable indicating whether extra information on progress should be reported. Default is TRUE.
...	ignored or passed to internal functions.

## Details

This function computes one or more timeslice maps of the spatio-temporal kernel intensity of radiocarbon dates across a geographic region and for a specific focal year. The user specifies the arbitrary sizes of both the spatial and the temporal Gaussian kernels that will be used to summarise radiocarbon date intensity per grid cell per timestep. The results allow standardisation of colour ramps, etc. across timesteps and are amenable to plotting individually via `plot.stKde` and/or for output to png for animation.

## Value

A list object of class `stKde` with the following elements:

- A series of list items storing some of the input parameters such as the `focalyears`, `sbw`, `tbw`, `backsight`, `maskthresh`
- `nonfocal` An `im` object mapping the basic spatial intensity of all dates, without reference to a focal year.
- `impaths` A character vector of the paths to the individual timeslices stored on file. Maps are not stored in memory (see `spkde()` for further details of what is stored).
- `stats` A list of `data.frames` offering summary statistics on each of the different types of output surface across all timeslices. Used primarily to allow consistent colour ramps across timeslices.
- `ppp` The `ppp` object for all dates and the observation window.

## Examples

```
## Not run:
## Example with a subset of English and Welsh dates from the Euroevol dataset
data(ewdates)
data(ewowin)
x <- calibrate(x=ewdates$C14Age, errors=ewdates$C14SD, normalised=FALSE)
## Create centennial timeslices (also with site binning)
bins1 <- binPrep(sites=ewdates$SiteID, ages=ewdates$C14Age, h=50)
stkde1 <- stkde(x=x, coords=ewdates[,c("Eastings", "Northings")], win=ewowin,
sbw=40000, cellres=2000, focalyears=seq(6500, 5000, -100), tbw=50, bins=bins1,
backsight=200, outdir="im", amount=1)
## Plot an example of all four basic outputs for 5900 calBP
dev.new(height=2.5, width=8)
par(mar=c(0.5, 0.5, 2.5, 2))
plot(stkde1, 5900, type="all")

## End(Not run)
```

---

subset.CalDates      *Subsetting calibrated dates*

---

### Description

Subsets calibrated dates (CalDates class object) based on Logical expressions of time intervals.

### Usage

```
## S3 method for class 'CalDates'  
subset(x, s, p, ...)
```

### Arguments

x	A CalDates class object
s	Logical expression indicating dates to keep. The expression should include the term BP which refers to specific dates.
p	Probability mass meeting the condition defined by ss.
...	Further arguments to be passed to or from other methods (ignored).

### Details

The function subsets CalDates class objects by identifying all dates that have a probability mass larger than p for a user defined logical expression of temporal interval containing the term BP, where BP refers to radiocarbon date. See examples for further details

### Value

A CalDates class object.

### Examples

```
## Generate some calibrated dates  
x = calibrate(c(12100,5410,5320,3320),errors=c(20,20,30,30))  
## Subsets all dates that have a probability mass above 0.8 before 10000 BP  
x2 = subset(x,BP>10000,p=0.8)  
## Subsets all dates that have a probability mass above 0.5 between 6000 and 6300 BP  
x3 = subset(x,BP>6000&&BP<6300,p=0.5)
```



---

summary.CalDates      *Summarise a CalDates class object*

---

### Description

Returns summary statistics of calibrated dates.

### Usage

```
## S3 method for class 'CalDates'
summary(object, prob = NA, calendar = "BP", ...)
```

### Arguments

object	A CalDates class object.
prob	A vector containing probabilities for the higher posterior density interval. Default is <code>c(0.683, 0.954)</code> , i.e. 1 and 2-Sigma range.
calendar	Whether the summary statistics should be computed in cal BP ("BP") or in BCAD ("BCAD").
...	further arguments passed to or from other methods.

### Value

A data.frame class object containing the ID of each date, along with the median date and one and two sigma (or a user specified probability) higher posterior density ranges.

---

summary.SpdModelTest      *Summarise a SpdModelTest class object*

---

### Description

summary method for class "SpdModelTest"

### Usage

```
## S3 method for class 'SpdModelTest'
summary(object, type = "spd", ...)
```

### Arguments

object	A SpdModelTest class object produced using the <code>link{modelTest}</code> function.
type	Specifies whether the summary should be based on SPD ('spd') or associated rates of change ('roc'). Default is 'spd'.
...	Ignored

**Details**

The summary function returns metadata (number of radiocarbon dates, bins, and simulations), the p-value of the global significance test, and the chronological interval of local positive and negative deviations from the simulation envelope.

**See Also**

[modelTest](#).

---

summary.SpdPermTest    *Summarise a SpdPermTest class object*

---

**Description**

summary method for class "SpdPermTest"

**Usage**

```
## S3 method for class 'SpdPermTest'
summary(object, type = "spd", ...)
```

**Arguments**

object	A SpdPermTest class object produced using the link{permTest} function.
type	Specifies whether the summary should be based on SPD ('spd') or associated rates of change ('roc'). Default is 'spd'.
...	Ignored

**Details**

The summary function returns metadata (number of radiocarbon dates, bins, and simulations), the p-value of the global significance test, and the chronological interval of local positive and negative deviations from the simulation envelope for each set.

**See Also**

[permTest](#).

---

thinDates	<i>Sampling function to select a maximum number of dates per site, bin or phase.</i>
-----------	--

---

### Description

Function to select a subset of uncalibrated radiocarbon dates up to a maximum sample size per site, bin or phase.

### Usage

```
thinDates(ages, errors, bins, size, thresh = 0.5, method = "random", seed = NA)
```

### Arguments

ages	A vector of uncalibrated radiocarbon ages
errors	A vector of uncalibrated radiocarbon errors (same length as ages)
bins	A vector of labels corresponding to site names, ids, bins or phases (same length as ages)
size	A single integer specifying the maximum number of desired dates for each label stated bin.
thresh	A single numeric value between 0 and 1 specifying the approximate proportion (after rounding) of the resulting sample that will be chosen according to lowest date errors. At the extremes, 0 produces a simple random sample whereas 1 selects the sample dates with the lowest errors. Ignored if method="random".
method	The method to be applied where "random" simple selects a random sample, whereas "splitsample", picks some proportion (see thresh) of the sample to minimise errors, and randomly samples the rest. At present, these are the only two options.
seed	Allows setting of a random seed to ensure reproducibility.

### Value

A numeric vector of the row indices corresponding to those of the input data.

### See Also

[binPrep](#)

### Examples

```
data(euroevol)
foursites <- euroevol[euroevol$SiteID %in% c("S2072", "S4380", "S6139", "S9222"),]
table(as.character(foursites$SiteID))
## Thin so each site has 10 dates each max, with random selection
thinInds<- thinDates(ages=foursites$C14Age, errors=foursites$C14SD,
```

```

bins=foursites$SiteID, size=10, method="random", seed=123)
tdates <- foursites[thinInds,]
tdates
## Same but choose the first 60% (i.e. 6 dates) from the lowest errors
## and then fill in the rest randomly.
thinInds<- thinDates(ages=foursites$C14Age, errors=foursites$C14SD,
bins=foursites$SiteID, size=10, method="splitsample", thresh=0.6, seed=123)
tdates1 <- foursites[thinInds,]
tdates1

```

---

uncalibrate	<i>Uncalibrate (back-calibrate) a calibrated radiocarbon date (or summed probability distribution).</i>
-------------	---

---

## Description

Function for uncalibrating one or more radiocarbon dates.

## Usage

```

uncalibrate(x, ...)

## Default S3 method:
uncalibrate(x, CRAerrors = 0, roundyear = TRUE, calCurves = "intcal13", ...)

## S3 method for class 'CalGrid'
uncalibrate(
  x,
  calCurves = "intcal13",
  eps = 1e-05,
  compact = TRUE,
  verbose = TRUE,
  ...
)

```

## Arguments

x	Either a vector of calibrated radiocarbon ages or an object of class CalGrid.
...	ignored
CRAerrors	A vector of standard deviations corresponding to each estimated radiocarbon age (ignored if x is a CalGrid object).
roundyear	An optional vector of IDs for each date (ignored if x is a CalGrid object).
calCurves	A string naming a calibration curve already provided with the rcarbon package (currently 'intcal13', 'intcal13nhpine16', 'shcal13', 'shcal13shkauri16', and 'marine13' are possible) or a custom curve provided as matrix/data.frame in three columns ("CALBP", "C14BP", "Error"). The default is the 'intcal13' curve and only one curve can currently be specified for all dates.

eps	Cut-off value for density calculation (for CalGrid objects only).
compact	A logical variable indicating whether only uncalibrated ages with non-zero probabilities should be returned (for CalGrid objects only).
verbose	A logical variable indicating whether extra information on progress should be reported (for CalGrid objects only).

### Details

This function takes one or more calibrated calendars and looks-up the corresponding uncalibrated age, error of the stated calibration curve at that point. It also provides a randomised estimate of the uncalibrated age based on the curve error (and optionally also a hypothetical measurement error).

### Value

A data.frame with specifying the original data, the uncalibrated age without the calibration curve error (ccCRA), the calibration curve error at this point in the curve (ccError), a randomised uncalibrated age (rCRA) given both the stated ccError and any further hypothesised instrumental error provided by the CRAerrors argument (rError).

### Examples

```
# Uncalibrate two calendar dates
uncalibrate(c(3050,2950))
```

---

which.CalDates	<i>Which indices for calibrated dates</i>
----------------	---

---

### Description

Gives the TRUE indices of calibrated dates (CalDates class object) based on Logical expressions of time intervals.

### Usage

```
which.CalDates(x, s, p)
```

### Arguments

x	A CalDates class object
s	Logical expression indicating dates to keep. The expression should include the term BP which refers to specific dates.
p	Probability mass meeting the condition defined by ss.

### Details

The function subsets CalDates class objects by identifying all dates that have a probability mass larger than p for a user defined logical expression of temporal interval containing the term BP, where BP refers to radiocarbon date. See examples for further details

**Value**

A CalDates class object.

**Examples**

```
## Generate some calibrated dates
x = calibrate(c(12100,5410,5320,3320),errors=c(20,20,30,30))
## Subsets all dates that have a probability mass above 0.8 before 10000 BP
x2 = which.CalDates(x,BP>10000,p=0.8)
## Subsets all dates that have a probability mass above 0.5 between 6000 and 6300 BP
x3 = which.CalDates(x,BP>6000&&BP<6300,p=0.5)
```

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