

# Package ‘esd’

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**Title** Climate analysis and empirical-statistical downscaling (ESD) package for monthly and daily data.

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**Depends** ncdf4,zoo,R (>= 2.10.0)

**Description** The package contains R functions for retrieving data, making climate analysis and downscaling of monthly mean and daily mean global climate scenarios.

**License** GPL (>= 2)

**URL** <http://rcg.gvc.gu.se/edu/esd.pdf>

**ZipData** no

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 aggregate

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 aggregate

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## Description

The aggregation is based on the S3 method for zoo objects, but takes care of extra house keeping, such as attributes with meta data.

**Usage**

```
aggregate.station(x,by,FUN = mean, ..., regular = NULL, frequency = NULL)
aggregate.field(x,by,FUN = mean, ..., regular = NULL, frequency = NULL)
aggregate.comb(x,by,FUN = mean, ...,
               regular = NULL, frequency = NULL)
aggregate.area(x,is=NULL,it=NULL,FUN=sum,na.rm=TRUE,smallx=FALSE)
```

**Arguments**

x	A <a href="#">station</a> , <a href="#">spell</a> or a <a href="#">field</a> object
by	see <a href="#">aggregate.zoo</a>
FUN	see <a href="#">aggregate.zoo</a>
regular	see <a href="#">aggregate.zoo</a>
frequency	see <a href="#">aggregate.zoo</a>
is	spatial selection - see <a href="#">subset.field</a>
na.rm	TRUE: ignore NA - see <a href="#">mean</a>

**Value**

The call returns a station object

**Author(s)**

R.E. Benestad

**See Also**

[spatial.avg.field as.4seasons, annual](#)

**Examples**

```
# Example: use aggregate to compute annual mean temperature for Svalbard:
data(Svalbard)
y <- aggregate(Svalbard, year, FUN=mean, na.rm = FALSE)
plot(y)

# Example for getting seasonal aggregate of standard deviation of
data(Oslo)
ym <- as.4seasons(Oslo,FUN=mean)
ys <- as.4seasons(Oslo,FUN=sd)
y <- combine(ym,ys)
plot(y)

x <- t2m.NCEP()
z <- aggregate.area(x,FUN=mean)
plot(z)
```

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annual	<i>Conversion to esd objects.</i>
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### Description

annual aggregates time series into annual values (e.g. means). year, month, season return the years, months, and days associated with the data.

### Usage

```
annual(x,...)
annual.zoo(x,FUN="mean",na.rm=TRUE,nmin=NULL,...)
annual.default(x,FUN="mean",na.rm=TRUE,nmin=NULL,...)
annual.dsensemble(x,FUN="mean")
annual.station(x,FUN="mean",nmin=NULL,...)
annual.spell(x,FUN="mean",nmin=NULL,...)
annual.field(x,FUN="mean",na.rm=TRUE,nmin=NULL,...)
year(x,...)
month(x,...)
day(x,...)
season(x,format="character")
season.name()
pentad(x,l=5,it0=NULL,...)
```

### Arguments

x	a station, field object, or a date
FUN	see <a href="#">aggregate.zoo</a>
nmin	Minimum number of data points (e.g. days or months) with valid data accepted for annual estimate. NULL demands complete years.
format	'numeric' or 'character'
na.rm	TRUE: ignore NA - see <a href="#">mean</a>
l	length of window
it0	Value of specific year to include in the aggregated series (it0=1970 and l=1 will give year 1970, 1975, 1980, ...)

### Value

Same as x, or a numeric for year, month, day, or pentad.

### Author(s)

R.E. Benestad and A. Mezghanil

**See Also**

[as.annual,aggregate.station](#)

**Examples**

```
# Example: how to generate a new station object.
data <- round(matrix(rnorm(20*12),20,12),2); colnames(data) <- month.abb
x <- data.frame(year=1981:2000,data)
X <- as.station.data.frame(x,loc="",param="noise",unit="none")

# Example: how to generate a new field object.
year <- sort(rep(1991:2000,12))
month <- rep(1:12,length(1991:2000))
n <-length(year)
lon <- seq(-30,40,by=5); nx <- length(lon)
lat <- seq(40,70,by=5); ny <- length(lat)
# Time dimension should come first, space second.
y <- matrix(rnorm(nx*ny*n),n,nx*ny)
index <- as.Date(paste(year,month,1,sep="-"))
Y <- as.field(y,index,lon,lat,param="noise",unit="none")
map(Y)
plot EOF(Y))

data(Oslo)
plot(as.anomaly(Oslo))

data(ferder)
plot(annual(ferder,FUN="min"))
plot(annual(ferder,FUN="IQR",na.rm=TRUE))
plot(as.4seasons(ferder))

data(bjornholt)
plot(annual(bjornholt,FUN="exceedance",fun="count"))
plot(annual(bjornholt,FUN="exceedance",fun="freq"))
plot(annual(bjornholt,FUN="exceedance"))

# Test the as.4seasons function:
data(ferder)
#Daily data:
yd <- ferder
# Monthly data:
ym <- aggregate(ferder,as.yearmon)
ym <- zoo(coredata(ym),as.Date(index(ym)))
ym <- attrcp(ferder,ym)
plot(ym)

#Monthly reanalyses:
t2m <- t2m.ERAINT(lon=c(-30,40),lat=c(50,70))
T2m <- as.4seasons(t2m)
#Extract the grid point with location corresponding to that of the station:
x <- regrid(t2m,is=ferder)
x4s <- as.4seasons(x)
```

```

X4s <- regrid(T2m,is=ferder)
y4s1 <- as.4seasons(yd)
y4s2 <- as.4seasons(ym)
plot.zoo(y4s1,lwd=2,xlim=as.Date(c("1980-01-01","2000-01-01")),ylim=c(-10,20))
lines(y4s2,col="red",lty=2)
lines(x4s,col="darkblue",lwd=2)
lines(X4s,col="lightblue",lty=2)

```

---

anomaly

*Anomaly and Climatology*


---

### Description

S3-method that computes anomalies and/or climatology for time series and fields. `clim2pca` is unfinished

### Usage

```

anomaly(x,...)
anomaly.default(x,...)
anomaly.comb(x,...)
anomaly.field(x,...)
anomaly.station(x,...)
anomaly.dsensemble(x,ref=NULL,...)
anomaly.annual(x,ref=1961:1990)
anomaly.month(x,ref=NULL)
anomaly.season(x,ref=NULL)
anomaly.day(x,ref=NULL)
climatology(x,...)
climatology.default(x)
climatology.field(x)
climatology.station(x)
clim2pca(x,...)
clim2pca.default(x)
clim2pca.month(x)
clim2pca.day(x)

```

### Arguments

<code>x</code>	A station or field object
<code>ref</code>	vector defining the reference interval

### Value

The call returns a similar object as `x` containing anomalies and data climatology.

In ‘`anomaly.dsensemble`’, the default value of the reference period is taken as the available time period from observations, i.e. same time period as in attribute ‘`station`’ is used as baseperiod to compute anomalies of GCM downscaled results.

**Author(s)**

R.E. Benestad

**See Also**[as.anomaly](#), [as.climatology](#)**Examples**

```
data(ferder)
plot(anomaly(ferder))
```

as

*Conversion to esd objects.***Description**

Various methods for converting objects from one shape to another. These methods do the house keeping, keeping track of attributes and metadata.

`as.field.station` uses `regrid` to generate a field based on bi-linear interpolation of station values and their coordinates. Unfinished...

**Usage**

```
as.4seasons(x, FUN=mean, ...)
as.4seasons.default(x, FUN=mean, ...)
as.4seasons.station(x, FUN=mean, ...)
as.4seasons.day(x, FUN=mean, na.rm=TRUE, dateindex=TRUE, nmin=85, ...)
as.4seasons.field(x, FUN=mean, ...)
as.4seasons.spell(x, FUN="mean", ...)
as.seasons(x, start=01-01, end=12-31, FUN=mean, ...)
as.annual(x, ...)
as.annual.default(x, ...)
as.annual.numeric(x, ...)
as.annual.integer(x, ...)
as.annual.yearqtr(x, frac = 0, ...)
as.annual.spell(x, ...)
as.annual.station(x, ...)
as.anomaly(x, ...)
as.anomaly.default(x, ref=NULL, na.rm=TRUE)
as.anomaly.station(x, ref=NULL, na.rm=TRUE)
as.anomaly.field(x, ref=NULL, na.rm=TRUE)
as.anomaly.zoo(x, ref=NULL, na.rm=TRUE)
as.appended(x, ...)
as.appended.ds.comb(x, iapp=1)
as.appended.eof.comb(x, iapp=1)
```

```
as.appended.field.comb(x,iapp=1)
as.eof(x,...)
as.eof.zoo(x,...)
as.eof.ds(x,iapp=NULL)
as.eof.eof(x,iapp=NULL)
as.eof.comb(x,iapp=NULL)
as.eof.field(x,iapp=NULL,...)
as.eof.appendix(x,iapp=1)
as.calibrationdata(x)
as.calibrationdata.ds(x)
as.calibrationdata.station(x)
as.climatology(x,...)
as.comb(x,...)
as.comb.eof(x,...)
as.ds(x)
as.field(x,...)
as.field.zoo(x,lon,lat,param,unit,
             longname=NA,quality=NA,src=NA,url=NA,
             reference=NA,info=NA,calendar=gregorian,
             greenwich=TRUE, method= NA,type=NA,aspect=NA)
as.field.default(x,index,lon,lat,param,unit,
                 longname=NA,quality=NA,src=NA,url=NA,
                 reference=NA,info=NA,calendar=gregorian,
                 greenwich=TRUE, method= NA,type=NA,aspect=NA)
as.field.eof(x,...)
as.field.comb(x,iapp=NULL,...)
as.field.ds(x,iapp=NULL,...)
as.field.station(x,...)
as.fitted.values(x)
as.fitted.values.ds(x)
as.fitted.values.station(x)
as.monthly(x,fun=mean)
as.observed.station(x)
as.original.data(x)
as.original.data.ds(x)
as.original.data.station(x)
as.pattern(x)
as.pattern.ds(x)
as.pattern.eof(x)
as.pattern.cca(x)
as.pattern.mvr(x)
as.pattern.field(x)
as.pattern.trend(x)
as.pattern.corfield(x)
as.pca(x)
as.pca.ds(x)
as.pca.station(x)
as.residual(x)
```



```

as.residual.ds(x)
as.residual.station(x)
as.station(x,...)
as.station.zoo(x,loc=NA,param=NA,unit=NA,lon=NA,lat=NA,alt=NA,
               cntr=NA,longname=NA,stid=NA,quality=NA,src=NA,url=NA,
               reference=NA,info=NA, method= NA)
as.station.data.frame(x,location=NA,param=NA,unit=NA,lon=NA,lat=NA,alt=NA,
                     cntr=NA,longname=NA,stid=NA,quality=NA,src=NA,url=NA,
                     reference=NA,info=NA, method= NA)
as.station.zoo(x,location=NA,param=NA,unit=NA,lon=NA,lat=NA,alt=NA,
              cntr=NA,longname=NA,stid=NA,quality=NA,src=NA,url=NA,
              reference=NA,info=NA, method= NA,type=NA,aspect=NA)
as.trajectory(x,n=10,loc=NA,param=NA,unit=NA,lon=NA,
              lat=NA,alt=NA,cntr=NA,longname=NA,stid=NA,
              quality=NA,src=NA,url=NA,reference=NA,info=NA,
              method= NA,verbose=FALSE)

as.station.list(x)
as.station.ds(x)
as.station.pca(x)
as.station.field(x)
as.station.spell(x)
as.station.eof(x,pattern=1:10)
as.pca.ds(x)
as.pca.station(x)

```

### Arguments

x	Data object
location	define location attribute attr(x,location)
param	define variable attribute attr(x,variable)
unit	define unit attribute attr(x,unit)
lon	define longitude attribute attr(x,longitude)
lat	define latitude attribute attr(x,latitude)
alt	define altitude attribute attr(x,altitude)
cntr	define country attribute attr(x,country)
longname	define long-name attribute attr(x,loongname)
stid	define station ID attribute attr(x,station_id)
quality	define quality attribute attr(x,quality)
src	define source attribute attr(x,source)
url	define URL attribute attr(x,URL)
reference	define reference attribute attr(x,reference)
info	define info attribute attr(x,info)
method	define method attribute attr(x,method)
FUN	function

<code>na.rm</code>	TRUE: ignore NA values
<code>nmin</code>	Minimum number of valid data points. Default value is set to <code>nmin=85</code> for daily values, <code>nmin=3</code> for monthly values, and <code>nmin=300</code> for annual values.
<code>dateindex</code>	
<code>monthly</code>	
<code>aspect</code>	
<code>iapp</code>	For values greater than 1, select the corresponding appended field in 'comb' objects (e.g. 1 gives <code>attr(x,appendix.1)</code> )
<code>pattern</code>	Which EOF pattern (mode) to extract as a series for PC

### Details

'field' object must be a two dimensional object (time,nlon\*nlat). Time index must be the first dimension. The second dimension is a product of the number of longitudes by the number of latitudes. For a different format, users can use 'aperm' function (Transpose an array by permuting its dimensions) to rearrange the object to meet the required format and 'dim' function can be used to transform a 3D (d1,d2,d3) object into 2D (d1,d2\*d3) object as `dim(3D) <- c(d1,d2*d3)`

### Value

A field object

### Author(s)

R.E. Benestad and A. Mezghani

### Examples

```
# Example: how to generate a new station object.
data <- round(matrix(rnorm(20*12),20,12),2); colnames(data) <- month.abb
x <- data.frame(year=1981:2000,data)
X <- as.station.data.frame(x,loc="",param="noise",unit="none")

# Example: how to generate a new field object.
year <- sort(rep(1991:2000,12))
month <- rep(1:12,length(1991:2000))
n <-length(year)
lon <- seq(-30,40,by=5); nx <- length(lon)
lat <- seq(40,70,by=5); ny <- length(lat)
# Time dimension should come first, space second.
y <- matrix(rnorm(nx*ny*n),n,nx*ny)
index <- as.Date(paste(year,month,1,sep="-"))
Y <- as.field(y,index,lon,lat,param="noise",unit="none")
map(Y)
plot EOF(Y))

data(Oslo)
plot(as.anomaly(Oslo))
```

```

data(ferder)
plot(annual(ferder,FUN="min"))
plot(annual(ferder,FUN="IQR",na.rm=TRUE))
plot(as.4seasons(ferder))

data(bjornholt)
plot(annual(bjornholt,FUN="exceedance",fun="count"))
plot(annual(bjornholt,FUN="exceedance",fun="freq"))
plot(annual(bjornholt,FUN="exceedance"))

# Test the as.4seasons function:
data(ferder)
#Daily data:
yd <- ferder
# Monthly data:
ym <- aggregate(ferder,as.yearmon)
ym <- zoo(coredata(ym),as.Date(index(ym)))
ym <- attrcp(ferder,ym)
plot(ym)

#Monthly reanalyses:
t2m <- t2m.ERAINT(lon=c(-30,40),lat=c(50,70))
T2m <- as.4seasons(t2m)
#Extract the grid point with location corresponding to that of the station:
x <- regrid(t2m,is=ferder)
x4s <- as.4seasons(x)
X4s <- regrid(T2m,is=ferder)
y4s1 <- as.4seasons(yd)
y4s2 <- as.4seasons(ym)
plot.zoo(y4s1,lwd=2,xlim=as.Date(c("1980-01-01","2000-01-01")),ylim=c(-10,20))
lines(y4s2,col="red",lty=2)
lines(x4s,col="darkblue",lwd=2)
lines(X4s,col="lightblue",lty=2)

# Select a random season
data(bjornholt)
data(ferder)
plot(as.seasons(ferder,FUN=CDD))
plot(as.seasons(ferder,start=05-17,end=11-11,FUN=HDD))

```

## Description

Applies a canonical correlation analysis (CCA) to two data sets. The CCA here can be carried out based on an [svd](#) based approach (after Bretherton et al. (1992), J. Clim. Vol 5, p. 541, also documented in Benestad (1998): "Evaluation of Seasonal Forecast Potential for Norwegian Land Temperatures and Precipitation using CCA", DNMI KLIMA Report 23/98 at <http://met.no/>

[english/r\\_and\\_d\\_activities/publications/1998.html](http://english/r_and_d_activities/publications/1998.html)) or ii) a covariance-eigenvalue approach (after Wilks, 1995, "Statistical methods in the Atmospheric Sciences", Academic Press, p. 401).

The analysis can also be applied to either EOFs or fields.

Note: the analysis has sometimes been somewhat unstable, returning inconsistent results. The recommendation is to use EOFs and SVD option.

The CCA analysis can be used to develop statistical models according to:

$$Y = \Psi X$$

Where Y is the predictand and X the predictor. `plotCCA` plots the CCA results, `testCCA` is for code verification, and `Psi` returns the matrix

$$\Psi$$

`stations2field` turns a group of station objects into a field by the means of a simple and crude interpolation/gridding. `check.repeat` is a quality-control function that eliminates repeated years in the station objects.

Try the same type of argument as in `lm('y ~ x, data=')`

## Usage

```
CCA(Y,X,...)
CCA.default(Y,X,...)
CCA.eof(Y,X,i.eof=1:8)
CCA.pca(Y,X,i.eof=1:8)
CCA.field(Y,X,i.eof=1:8)
test.cca(method="CCA",reconstr=FALSE,mode=1,test=TRUE,LINPACK=TRUE,
         SVD=TRUE,n.pc=4,synthetic=TRUE)

predict.CCA <- function(object, newdata=NULL, ...)
```

## Arguments

Y	An object with climate data: field, eof, pca.
X	Same as Y.
SVD	Use a singular value decomposition as a basis for the PCA.
i.eof	Which EOFs to include in the CCA.
LINPACK	an option for <a href="#">svd</a> .
object	The result from CCA.
newdata	The same as X.

## Value

A CCA object: a list containing a.m, b.m, u.k, v.k, and r, describing the Canonical Correlation variates, patterns and correlations. a.m and b.m are the patterns and u.k and v.k the vectors (time evolution).

**Author(s)**

R.E. Benestad

**Examples**

```
# CCA with two eofs
slp <- slp.NCEP(lat=c(-40,40),anomaly=TRUE)
sst <- sst.NCEP(lat=c(-40,40),anomaly=TRUE)
eof.1 <- EOF(slp,it=1)
eof.2 <- EOF(sst,it=1)
cca <- CCA(eof.1,eof.2)
plot(cca)

# CCA with PCA and EOF:
NACD <- station.nacd()
plot(annual(NACD))
map(NACD,FUN="sd")
pca <- PCA(NACD)
plot(pca)
naslp <- slp.NCEP(lon=c(-30,40),lat=c(30,70),anomaly=TRUE)
map(naslp)
eof <- EOF(naslp,it=1)
nacca <- CCA(pca,eof)
plot(nacca)
```

CCI

*Calculus Cyclone identification.***Description**

Identifies cyclones (low pressure systems) in a gridded data set using a Calculus Cyclone Identification (CCI) method (EMS04-A-00146, Benestad, R.E.; Sorteberg, A.; Chen, D. 'Storm statistics derived using a calculus-based cyclone identification method', [http://www.cosis.net/members/meetings/sessions/oral\\_programme.php?p\\_id=110&s\\_id=1845](http://www.cosis.net/members/meetings/sessions/oral_programme.php?p_id=110&s_id=1845), European Meteorological Society AC2, Nice, Sept 28, 2004). Storms are identified with longitude, latitude, and date. Also returned are estimates of local minimum pressure, max pressure gradient near storm, max geostrophic and gradient windspeed near storm, and radius of the storm. The storm location is by means of finding where first derivatives of north-south and east-west gradients both are zero. The storm radius is estimated from the points of inflexion along the latitude and longitude lines running through the centre of the storm.

If a north-south or east-west sea level pressure (p) profile can be approximated as

$$p(\theta) = p_0 + \sum_{i=1}^{N_\theta} [a_\theta(i) \cos(\omega_\theta(i)\theta) + b_\theta(i) \sin(\omega_\theta(i)\theta)]$$

Then the pressure gradient can be estimated as:

$$\frac{\partial \hat{p}(\theta)}{\partial \theta} = \sum_{i=1}^{N_\theta} \omega_\theta(i) [-\hat{a}_\theta(i) \sin(\omega_\theta(i)\theta) + \hat{b}_\theta(i) \cos(\omega_\theta(i)\theta)]$$

The gradient in x and y directions are found after the transform

$$\frac{d\hat{p}(x)}{dx} = \frac{1}{a \cos(\phi)} \frac{d\hat{p}(\theta)}{d\theta}$$

and

$$\frac{d\hat{p}(y)}{dy} = \frac{1}{a} \frac{d\hat{p}(\phi)}{d\phi}$$

(Gill, 1982).

This code is based on the CCI method of the R-package 'cyclones' and has been adapted for 'esd'.

The maximum gradient wind (max.speed) is estimated as described in Fleagle and Businger (1980) p. 163. (eq 4.27).

Reference: Benestad & Chen (2006) 'The use of a Calculus-based Cyclone Identification method for generating storm statistics', Tellus A, in press. Benestad (2005)

### Usage

```
CCI(Z,m=14,it=NULL,is=NULL,label=NULL,cyclones=TRUE,
    mindistance=1E6,dpmin=5E-4,pmax=NULL,rmin=1E4,rmax=2E6,
    accuracy=NULL,nsim=NULL,fname="cyclones.rda",
    lplot=FALSE,verbose=FALSE)
```

### Arguments

Z	A field object.
m	Number of harmonics used for fit to profile (Fourier truncation).
it	A list providing time index, e.g. month.
is	A list providing space index, lon and/or lat.
label	Label for ID-purposes.
cyclones	TRUE: identifies cyclones, FALSE: anticyclones.
mindistance	Min distance between cyclones. Unit: m.
dpmin	Min pressure gradient at points of inflection around cyclone. Unit: Pa/m (10hPa/km).
pmax	Max pressure in center of cyclone. Unit: hPa.
rmin	Min average radius of cyclone. Unit: m.
rmax	Max average radius of cyclone. Unit: m.
nsim	Number of simultaneous cyclones identified and saved ordered according to depth/strength.
fname	Name of output file.
lplot	TRUE: produce plots.
verbose	Print out diagnostics.
accuracy	Not yet finished.

**Value**

An 'events' object: `as.events(X=data.frame(date,time,lon,lat,pcent,max.dspl, max.speed,radius,qf))`  
 The subobjects `lon` (longitude: units degrees), `lat` (latitude: units: degrees), `pcent` (local minimum pressure: units hPa), `max.dspl` (pressure gradient: units hPa/m), `max.speed` (windspeed: units m/s), and `radius` (radius of the storm: units km), `qf` (quality flag, 1 = OK, 2 = less spatially precise, identified after widening the pressure gradient zero-crossings) are `[1:nt]`-vectors.

**Author(s)**

K.M. Parding & R.E. Benestad

---

coherence

*Coherence spectrum - cross-spectrum analysis*

---

**Description**

Based on: [http://en.wikipedia.org/wiki/Wiener-Khinchin\\_theorem](http://en.wikipedia.org/wiki/Wiener-Khinchin_theorem); Press et al. (1989) 'Numerical Recipes in Pascal', Cambridge, section 12.8 'Maximum Entropy (All Poles) Method'; von Storch & Zwiers (1999) 'Statistical Analysis in climate Research', Cambridge, section 11.4, eq 11.67, p. 235;

A test with two identical series the original equation (eq 11.67) from von Storch & Zwiers (1999) gave uniform values: 1. The denominator was changed from  $(\Gamma_{xx} * \Gamma_{yy})$  to  $(\sqrt{\Gamma_{xx} * \Gamma_{yy}})$ .

**Usage**

```
coherence(x,y,dt=1,M=NULL,plot=TRUE)
testcoherence(x=NULL,y=NULL)
```

**Arguments**

<code>x</code>	A vector (time series).
<code>y</code>	A vector (time series).
<code>dt</code>	time incremet - for plotting.
<code>M</code>	Window length - default= half series length
<code>plot</code>	Flag: plot the diagnostics.

**Value**

A complex vector .

**Author(s)**

R.E. Benestad

## Examples

```
## Not run:
data(DNMI.t2m)
data(DNMI.slp)
eof.1 <- EOF(DNMI.t2m,mon=1)
eof.2 <- EOF(DNMI.slp,mon=1)
cca <- CCA(eof.1,eof.2)
# Testing routine:
testCCA()

data(oslo.dm)
testcoherence(oslo.dm$t2m,oslo.dm$t2m)

## End(Not run)
```

---

col.bar

*Color bar*


---

## Description

Display a color bar object on an existing plot.

## Usage

```
col.bar(breaks, horiz = TRUE, pch = 21, v = 1, h = 1, col = col, cex =
2, cex.lab = 0.6, type = "r", verbose = FALSE, vl = 0.5, border = FALSE,
...)
colbar(breaks,col,fig=c(0.15,0.2,0.15,0.3),horiz=FALSE)
```

## Arguments

breaks	A numeric vector of breakpoints for the colours
horiz	
pch	<a href="#">par</a>
v	Vertical space between color bar points
h	horizontal space between color bar points
col	<a href="#">par</a>
cex	<a href="#">par</a>
cex.lab	<a href="#">par</a>
type	r : rectangular shape , p : for points
verbose	
vl	Vertical lines
border	Color bar borders
...	More graphical parameters to be passed



## Details

Insert a color bar or color points into an existing plot or map

## Author(s)

A. Mezghani

## Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.

## The function is currently defined as
function (breaks, horiz = TRUE, pch = 21, v = 1, h = 1, col = col,
        cex = 2, cex.lab = 0.6, type = "r", verbose = FALSE, vl = 0.5,
        border = FALSE, ...)
{
  par0 <- par()
  xleft <- par()$usr[1]
  xright <- par()$usr[2]
  ybottom <- par()$usr[4] - 1 - h
  ytop <- par()$usr[4] - 1
  by <- (xright - xleft - v * (length(col)))/(length(breaks))
  steps <- seq(0, (xright - xleft - v * (length(col))), by = by)
  nsteps <- length(steps)
  if (verbose)
    print(steps)
  if (verbose)
    print(breaks)
  if (verbose)
    print(nsteps)
  k <- 1/2
  for (i in 1:(nsteps - 2)) {
    if (!is.null(v))
      if (i == 1)
        k <- k + v/2
      else k <- k + v
    if (type == "r") {
      rect(xleft = k + xleft + steps[i], xright = k + xleft +
        steps[i + 1], ybottom = ybottom, ytop = ytop,
        col = col[i], border = border)
    }
    else if (type == "p") {
      points(x = k + xleft + (steps[i] + steps[i + 1])/2,
        y = (ybottom + ytop)/2, pch = pch, bg = col[i],
        cex = cex, ...)
    }
    text(x = k + xleft + (steps[i] + steps[i + 1])/2, y = ybottom -
      vl, labels = levels(cut(breaks, breaks))[i], col = "grey50",
      cex = cex.lab)
  }
}
```

```

    par(fig = par0$fig)
  }

```

---

combine

*Combine*

---

## Description

combine is a S3 method for combining esd objects, e.g. into groups of stations, stations and eof object, or fields. The function is based on [merge](#), and is also used to synchronise the esd objects.

For fields, combine.field is used to append different data sets, e.g. for the purpose of computing common EOFs (see [EOF](#) or for mixing fields (coupled EOFs).

For stations, combine.station can work two ways: (1) to combine a set of stations and group them into one data object; (2) combine series with different monthly values for one specific site into one record for the monthly data. E.g. January, February, ..., December months can be combined into one complete series of monthly data.

For DS-results, combine.ds is based on combine.station, but also takes care of the additional meta data (the original series and predictor patterns). For instance, this method can combine separate downscaled results for each calendar months at a single location into one complete time series.

g2d1 transform objects between grid starting at the greenwich (greenwich=TRUE) and the data line (greenwich=FALSE).

sp2np re-arranges field objects according to a grid going from 90S (South Pole) to 90N (North Pole) for SP2NP=TRUE. Otherwise, the object is arranged from 90N to 90S.

softattr copies the names of a subset of the attributes excluding "index", "dim" and others specified by ignore. attrcp passes on the attributes from one object (x) to another (y).

zeros counts the occurrence of zero values in a vector.

Other operations, such as `c(...)`, `rbind(...)` (combine along the time dimension), and `cbind(...)` (combine along the space dimension) also work.

## Usage

```

attrcp(x,y,ignore=NULL)
combine(x,y,...)
combine.default(x,y,all=FALSE,orig.format=TRUE)
combine.ds(...,all=TRUE)
combine.ds.comb(...,all=TRUE)
combine.ds.station(...,all=TRUE)
combine.ds.pca(...,all=TRUE)
combine.field(x,y,all=FALSE,dimension="time",approach="field",orig.format=TRUE,verbose=FALSE)
combine.field.station(x,y,all=FALSE,orig.format=TRUE,verbose=FALSE)
combine.list(...,all=TRUE)
combine.station(...,all=TRUE)
combine.station.month(...)
combine.station.eof(x,y,all=FALSE,orig.format=TRUE)
combine.station.field(x,y,all=FALSE,orig.format=TRUE)

```

```

combine.stations(...,all=TRUE)
combine.zoo(...)
g2dl(x,greenwich=TRUE,...)
g2dl.field(x,greenwich=TRUE)
g2dl.corfield(x,greenwich=TRUE)
g2dl.default(x,greenwich=TRUE,lon=NULL,lat=NULL,d=NULL)
g2dl.eof(x,greenwich=TRUE)
sp2np(x,SP2NP=TRUE)
softattr(x,ignore=NULL)
zeros(x)

```

### Arguments

x	station, eof, or field object
all	See link{merge.zoo}
orig.format	TRUE: the result will be formatted the same way as the input.
dimension	Which dimension to combine - in time or in space
approach	How to combine
greenwich	TRUE: center map on the Greenwich line (0E)
SP2NP	TRUE: order from south pole (bottom of plot) to north pole (top of plot)
ignore	List of attributes to ignore.

### Value

A field object

### Author(s)

R.E. Benestad

### Examples

```

library(esd)
t2m <- t2m.NCEP(lon=c(-40,40),lat=c(30,70))
T2m <- t2m.NorESM.M(lon=c(-40,40),lat=c(30,70))

# Combine in time to compute common EOFs:
X <- combine(t2m,T2m)
ceof <- EOF(X,it=1)
plot(ceof)

# Use combine to synchronise field and station data:
data(Oslo)
y <- combine.field.station(Oslo,t2m)
plot(y$y)

```

corfield

*Correlation***Description**

Compute the correlation between field objects and station/field.

**Usage**

```
corfield(x,y,...)
corfield.zoo(x,y,plot=TRUE,use=pairwise.complete.obs,verbose=FALSE,
             colbar=list(breaks=seq(-1,1,by=0.05),rev=TRUE))
corfield.field(x,y,plot=TRUE,use=pairwise.complete.obs,verbose=FALSE,
               colbar=list(breaks=seq(-1,1,by=0.05),rev=TRUE),...)
corfield.field.station(x,y,plot=TRUE,verbose=FALSE,
                       colbar=list(breaks=seq(-1,1,by=0.05),rev=TRUE),
                       use=pairwise.complete.obs,...)
corfield.station(x,y,plot=TRUE,verbose=FALSE,
                 use=pairwise.complete.obs,na.action=na.omit,
                 colbar=list(breaks=seq(-1,1,by=0.05),rev=TRUE),...)
corfield.eof(x,y,pattern=1,plot=TRUE,
             colbar=list(breaks=seq(-1,1,by=0.05),rev=TRUE),
             use=pairwise.complete.obs,na.action=na.omit,...)
corfield.trajectory(x,y,it=NULL,is=NULL,param=NULL,FUN="count",
                   unit=NULL,longname=NULL,loc=NULL,
                   use="pairwise.complete.obs",
                   colbar=list(breaks=seq(-1,1,by=0.05),rev=TRUE),
                   method="pearson")
```

**Arguments**

x	data object
y	data object
plot	TRUE: plot the results
use	see <a href="#">cor</a> .

**Value**

Map of correlation

**Author(s)**

R.E. Benestad and A. Mezghani

**Examples**

```

x <- t2m.ERAINT(lon=c(-40,30),lat=c(0,50))
y <- t2m.NCEP(lon=c(-40,30),lat=c(0,50))
r <- corfield(annual(x),annual(y))

data(Oslo)
t2m <- t2m.ERAINT()
x <- subset(Oslo,it=1)
y <- subset(t2m,it=1)
r <- corfield(x,y)

```

crossval

*Cross-validation***Description**

Applies a cross-validation of DS results, using the same strategy as in the DS exercise. Any step-wise screening is applied for each iteration independently of that used to identify the subset of skillful predictors in the original analysis. The model coefficients (beta) is saved for each iteration, and both correlation and root-mean-squared-error are returned as scores.

**Usage**

```

crossval(x, m=5, ...)
crossval.ds(x, m=5, ...)
crossval.list(x, m=5, ...)

```

**Arguments**

x	The results from <a href="#">DS</a> .
m	window with - leave m-out for each iteration. There are also some pre-set options: 'cordex-esd-exp1', 'value-exp1', and 'loo' for experiments defined at CORDEX-ESD, COST-VALUE, and leave-one-out ('loo') cross-validation.

**Value**

Cross-validation object.

**Author(s)**

R.E. Benestad

## Examples

```
data(Oslo)
t2m <- t2m.NCEP(lon=c(-20,40),lat=c(45,65))
eof <- EOF(t2m,1)

ds <- DS(Oslo,eof)
xv <- crossval(ds)
plot(xv)
```

---

Data

*Sample data.*

---

## Description

Different data sets: station data from northern Europe (NACD, NARP) and historic reconstructions (Oslo, Svalbard) from Dr. Nordli, Met Norway.

The object `station.meta` contains station information, used in the methods `station`.

Also reduced representation of re-analyses, where the data have been sampled by skipping grid points to reduce the spatial dimensions and stored as 20 EOFs (30 for precipitation). The data compression facilitated by the EOFs can provide 80-90% of the variance in the data. ESD uses the large-scale features from these reanalyses, and hence this information loss may be acceptable for downscaling work.

A reduced copy of the NorESM (M RCP 4.5) is also provided for the examples and demonstrations on how the downscaling can be implemented. Note: downscaling for end-users should never be based on one GCM simulation alone.

The object `geoborders` contains data on coastlines and borders, used in the methods `map`.

`glossstations` contains META-data for GLOSS stations taken from the table in [http://www.gloss-sealevel.org/station\\_handbook/stations/#.Vqtc6kL4phg](http://www.gloss-sealevel.org/station_handbook/stations/#.Vqtc6kL4phg)

Some data sets, such as NINO3.4 and NAOI come with a 'frozen' version in the package, but there are also functions that read the most recent version of these indices from the Internet. GSL reads the global mean sea level.

## Usage

```
data(bjornholt)
data(ferder)
data(dse.ferder)
data(vardo)
data(eca.meta)
data(station.meta)
data(NACD)
data(NARP)
data(Oslo)
data(Svalbard)
data(eof.t2m.ERAINT)
```

```

data(eof.t2m.ERA40)
data(eof.t2m.NCEP)
data(eof.precip.ERAINT)
data(eof.slp.ERAINT)
data(eof.slp.MERRA)
data(eof.slp.NCEP)
data(eof.t2m.NorESM.M)
data(eof.t2m.DNMI)
data(eof.sst.DNMI)
data(eof.slp.DNMI)
data(geoborders)
slp.MERRA(lon=NULL,lat=NULL,anomaly=FALSE)
t2m.MERRA(lon=NULL,lat=NULL,anomaly=FALSE)
t2m.NCEP(lon=NULL,lat=NULL,anomaly=FALSE)
sst.NCEP(lon=NULL,lat=NULL,anomaly=FALSE)
slp.NCEP(lon=NULL,lat=NULL,anomaly=FALSE)
t2m.ERAINT(lon=NULL,lat=NULL,anomaly=FALSE)
precip.ERAINT(lon=NULL,lat=NULL,anomaly=FALSE)
slp.ERAINT(lon=NULL,lat=NULL,anomaly=FALSE)
t2m.ERA40(lon=NULL,lat=NULL,anomaly=FALSE)
t2m.DNMI(lon=NULL,lat=NULL,anomaly=FALSE)
slp.DNMI(lon=NULL,lat=NULL,anomaly=FALSE)
sst.DNMI(lon=NULL,lat=NULL,anomaly=FALSE)
t2m.NorESM.M(lon=NULL,lat=NULL,anomaly=FALSE)
data(sunspots)
data(NINO3.4)
data(NAOI)
NINO3.4(url=ftp://ftp.cpc.ncep.noaa.gov/wd52dg/data/indices/ersst3b.nino.mth.ascii,header=TRUE)
NAO(url=http://www.cpc.ncep.noaa.gov/products/precip/CWlink/pna/norm.nao.monthly.b5001.current.ascii)
SOI(url=ftp://ftp.bom.gov.au/anon/home/ncc/www/sco/soi/soiplaintext.html,header=FALSE)
data(dse.Oslo)
HadCRUT4(url="http://www.metoffice.gov.uk/hadobs/hadcrut4/data/current/time_series/HadCRUT.4.2.0.0.")
NASAgiss(url=http://data.giss.nasa.gov/gistemp/taledata_v3/GLB.Ts+dSST.txt)
GSL <- function(url=http://www3.epa.gov/climatechange/images/indicator_downloads/sea-level_fig-1.csv)

```

### Arguments

lon	longitude range c(lon.min,lon.max)
lat	latitude range
anomaly	TRUE: return anomaly
url	source of data
plot	TRUE:plot

### Value

Numeric vectors/matrices with a set of attributes describing the data.

**Author(s)**

R.E. Benestad

**See Also**[aggregate.area as.4seasons, annual](#)**Examples**

```
data(Oslo)
year <- as.numeric( format(index(Oslo), %Y) )
plot(aggregate(Oslo, by=year,FUN=mean, na.rm = FALSE))

data(etopo5)
z <- subset(etopo5,is=list(lon=c(-10,30),lat=c(40,60)))
map(z)
```

diagnose

*Diagnose***Description**

Diagnose and examine combined fields, MVR, and CCA results. applies some tests to check for consistency.

The method `diagnose.comb.eof` which estimates the difference in the mean for the PCs of the calibration data and GCMs over a common period in addition to the ratio of standard deviations and lag-one autocorrelation. This 'bias correction' is described in Imbert and Benestad (2005), *Theor. Appl. Clim.* <http://dx.doi.org/10.1007/s00704-005-0133-4>.

`climvar` estimates the climatological variance, e.g. how the inter-annual variance varies with seasons.

**Usage**

```
diagnose(x,...)
diagnose.default(x,...)
diagnose.comb(x)
diagnose.eof(x)
diagnose.comb.eof(x)
diagnose.mvr(x)
diagnose.cca(x)
diagnose.ds(x,plot=FALSE)
diagnose.station(x,it=NULL,...)
diagnose.matrix(X,xlim=NULL,ylim=NULL,verbose=FALSE,...)
diagnose.dsensemble(x,plot=TRUE,map=TRUE,plot.type=target,...)
diagnose.distr(x,main=Mean - quantile,
               xlab=mean,ylab=expression(q[p]),
               sub=src(x),probs=0.95)
```



**Arguments**

**x** data object  
**it** teporal selection - see [subset](#)  
**plot**  
**plot.type**

**Value**

A 'diag' object containing test results

**Author(s)**

R.E. Benestad

**Examples**

```

t2m <- t2m.NCEP(lon=c(-40,40),lat=c(30,70))
T2m <- t2m.NorESM.M(lon=c(-40,40),lat=c(30,70))
# Combine in time to compute common EOFs:
X <- combine(t2m,T2m)
diagnose(X)

ceof <- EOF(X,it=1)
plot(diagnose(ceof))

slp <- slp.NCEP(lat=c(-40,40),anomaly=TRUE)
sst <- sst.NCEP(lat=c(-40,40),anomaly=TRUE)
eof.1 <- EOF(slp,it=1)
eof.2 <- EOF(sst,it=1)
cca <- CCA(eof.1,eof.2)
diagnose(cca)

```

---

DS

*Downscale*


---

**Description**

Identifies statistical relationships between large-scale spatial climate patterns and local climate variations for monthly and daily data series.

The function calibrates a linear regression model using step-wise screening and common EOFs ([EOF](#)) as basis functions. It then valuates the statistical relationship and predicts the local climate parameter from predictor fields.

The function is a S3 method that Works with ordinary EOFs, common EOFs ([combine](#)) and mixed-common EOFs. DS can downscale results for a single station record as well as a set of stations. There are two ways to apply the downscaling to several stations; either by looping through each

station and carrying out the DS individually or by using PCA to describe the characteristics of the whole set. Using PCA will preserve the spatial covariance seen in the past. It is also possible to compute the PCA prior to carrying out the DS, and use the method DS.pca. DS.pca differs from the more generic DS by (default) invoking different regression modules (link{MVR} or CCA).

The rationale for using mixed-common EOFs is that the coupled structures described by the mixed-field EOFs may have a more physical meaning than EOFs of single fields [Benestad et al. (2002), "Empirically downscaled temperature scenarios for Svalbard", *Atm. Sci. Lett.*, doi.10.1006/asle.2002.0051].

The function DS() is a generic routine which in principle works for when there is any real statistical relationship between the predictor and predictand. The predictand is therefore not limited to a climate variable, but may also be any quantity affected by the regional climate. *It is important to stress that the downscaling model must reflect a well-understood (physical) relationship.*

The routine uses a step-wise regression (step) using the leading EOFs. The calibration is by default carried out on de-trended data [ref: Benestad (2001), "The cause of warming over Norway in the ECHAM4/OPYC3 GHG integration", *Int. J. Clim.*, 15 March, vol 21, p.371-387.].

DS.seasonalcycle is an experimental set-up where the calibration is carried out based on the similarity of the seasonal variation to make most use of available information on a 'worst-case' basis, taking the upper limit view that at most, all the seasonal cycle is connected to the corresponding seasonal cycle in the predictor. See Benestad (2009) 'On Tropical Cyclone Frequency and the Warm Pool Area' *Nat. Hazards Earth Syst. Sci.*, 9, 635-645, 2009 <http://www.nat-hazards-earth-syst-sci.net/9/635/2009/nhess-9-635-2009.html>.

The function biasfix provides a type of 'bias correction' based on the method diagnose which estimates the difference in the mean for the PCs of the calibration data and GCMs over a common period in addition to the ratio of standard deviations and lag-one autocorrelation. This 'bias correction' is described in Imbert and Benestad (2005), *Theor. Appl. Clim.* <http://dx.doi.org/10.1007/s00704-005-0133-4>.

## Usage

```
DS(y,X,verbose=TRUE,plot=FALSE,...)
DS.default(y, X, mon = NULL, method = "lm", swsm = "step", m = 5,
  rmtrend = TRUE, eofs = 1:7,
  area.mean.expl = FALSE, verbose = FALSE, weighted = TRUE, ...)
DS.station(y,X,biascorrect=FALSE,mon=NULL,
  method="lm",swsm="step",m = 5,
  rmtrend=TRUE,eofs=1:7,area.mean.expl=FALSE,
  verbose=FALSE,pca=TRUE,neofs=20,...)
DS.eof(X,y,mon=NULL,
  method="lm",swsm="step",m=5,
  rmtrend=TRUE,eofs=1:7,area.mean.expl=FALSE,
  verbose=FALSE,pca=TRUE,npca=20,...)
DS.comb(X,y,biascorrect=FALSE,mon=NULL,
  method="lm",swsm="step",m=5,
  rmtrend=TRUE,eofs=1:7,area.mean.expl=FALSE,
  verbose=FALSE,...)
DS.field(X,y,biascorrect=FALSE,mon=NULL,
  method="lm",swsm="step",m=5,
  rmtrend=TRUE,eofs=1:7,area.mean.expl=FALSE,
```

```

        verbose=FALSE,...)
DS.t2m.month.field(y,X,biascorrect=FALSE,mon=NULL,m=5,
                  method="lm",swsm="step",
                  rmtrend=TRUE,eofs=1:7,area.mean.expl=FALSE,
                  verbose=FALSE,station=TRUE)
DS.t2m.season.field(y,X,biascorrect=FALSE,
                  method="lm",swsm="step",m=5,
                  rmtrend=TRUE,eofs=1:7,area.mean.expl=FALSE,
                  verbose=FALSE,station=TRUE)
DS.precip.season.field(y,X,biascorrect=FALSE,
                  method="lm",swsm="step",
                  rmtrend=TRUE,eofs=1:7,area.mean.expl=FALSE,
                  verbose=FALSE,...)
DS.freq(y,X,threshold=1,biascorrect=FALSE,method="glm",
        family="gaussian",swsm="step",
        rmtrend=TRUE,eofs=1:7,verbose=FALSE,...)
DS.spell(y,X,threshold=1,biascorrect=FALSE,method="glm",
        family="gaussian",swsm="step",
        rmtrend=TRUE,eofs=1:7,verbose=FALSE,...)
DS.pca(y,X,biascorrect=FALSE,mon=NULL,
        method="MVR",swsm=NULL,rmtrend=TRUE,...)
DS.seasonalcycle(y,X,mon=NULL,FUN=wetmean,fun=mean,
                 method="lm",swsm="step",m=1,
                 eofs=1:7,area.mean.expl=FALSE,
                 verbose=FALSE,weighted=TRUE,...)

predict.ds.mon(x,...)
sametimescale(y,X,FUN=mean)
biasfix(x)

```

## Arguments

y	The predictand - the station series representing local climate parameter
X	The predictor - an <a href="#">EOF</a> object representing the large-scale situation.
mon	An integer or a vector of integers. The calendar month to downscale, e.g. '1' is all January months, '2' is February, and so on.
method	Model type, e.g. <a href="#">lm</a> or <a href="#">glm</a>
swsm	Stepwise screening, e.g. <a href="#">step</a> . NULL skips stepwise screening
rmtrend	TRUE for detrending the predicant and predictors (in the PCs) before calibrating the model
eofs	Which EOF modes to include in the model training.
plot	TRUE: plot the results
verbose	TRUE: suppress output to the terminal.
station	TRUE: convert monthly data to station class using <a href="#">combine.ds</a> , else return a list of different monthly DS-results.

area.mean.exp1	When TRUE, subtract the area mean for the domain and use as a the first co-variate before the PCs from the EOF analysis.
m	passed on to <a href="#">crossval</a> . A NULL value suppresses the cross-validation, e.g. for short data series.
weighted	TRUE: use the attribute 'error.estimate' as weight for the regresion analysis.
...	

## Value

The downscaling analysis returns a time series representing the local climate, patterns of large-scale anomalies associated with this, ANOVA, and analysis of residuals. Care must be taken when using this routine to infer local scenarios: check the R2 and p-values to check wether the calibration yielded an appropriate model. It is also important to examine the spatial structures of the large-scale anomalies assocaiated with the variations in the local climate: do these patterns make physical sense?

It is a good idea to check whether there are any structure in the residuals: if so, then a linear model for the relationship between the large and small-scale structures may not be appropriate. It is furthermore important to experiment with predictors covering different regions [ref: Benestad (2001), "A comparison between two empirical downscaling strategies", *Int. J. Climatology*, **vol 21**, Issue 13, pp.1645–1668. DOI 10.1002/joc.703].

There is a cautionary tale for how the results can be misleading if the predictor domain in not appropriate: domain for northern Europe used for sites in Greenland [ref: Benestad (2002), "Empirically downscaled temperature scenarios for northern Europe based on a multi-model ensemble", *Climate Research*, **vol 21** (2), pp.105–125. <http://www.int-res.com/abstracts/cr/v21/n2/index.html>]

## Author(s)

R.E. Benestad

## Examples

```
# One exampe doing a simple ESD analysis:
X <- t2m.ERA40(lon=c(-40,50),lat=c(40,75))
data(Oslo)
#X <- OptimalDomain(X,Oslo)
eof <- EOF(X,it=jan)
Y <- DS(Oslo,eof)
plot(Y)
str(Y)

# Look at the residual of the ESD analysis
y <- as.residual(Y)
plot(y)

# Check the residual: dependency to the global mean temperature?
T2m <- (t2m.ERA40())
yT2m <- merge.zoo(y,T2m)
plot(coredata(yT2m[,1]),coredata(yT2m[,2]))
```

```

# Example: downscale annual wet-day mean precipitation -calibrate over
# part of the record and use the other part for evaluation.
T2M <- as.annual(t2m.NCEP(lon=c(-10,30),lat=c(50,70)))
cal <- subset(T2M,it=c(1948,1980))
pre <- subset(T2M,it=c(1981,2013))
comb <- combine(cal,pre)
X <- EOF(comb)
data(bjornholt)
y <- as.annual(bjornholt,FUN="exceedance")
z <- DS(y,X)
plot(z)

lon <- c(-12,37)
lat <- c(52,72)
ylim <- c(-6,6)
t2m <- t2m.NCEP(lon=lon,lat=lat)
T2m <- t2m.NorESM.M(lon=lon,lat=lat)
data(Oslo)
X <- combine(t2m,T2m)

eof <- EOF(X,it=7)
ds <- DS(Oslo,eof)
plot(ds)

DS(Oslo,X,station=FALSE) -> y
plot(y)
##Y <- combine.ds.comb(y)
##plot(Y)

data(ferder)
t2m <- t2m.NCEP(lon=c(-30,50),lat=c(40,70))
slp <- slp.NCEP(lon=c(-30,50),lat=c(40,70))
T2m <- as.4seasons(t2m)
SLP <- as.4seasons(slp)
X <- EOF(T2m,it=1)
Z <- EOF(SLP,it=1)
y <- ferder
sametimescale(y,X) -> z
ym <- as.4seasons(y,FUN="mean")
ys <- as.4seasons(y,FUN="sd")
dsm <- DS(ym,X)
plot(dsm)
dss <- DS(ys,Z)
plot(dss)

## Example for downscaling with missing data
data(Oslo)
era40 <- t2m.ERA40(lon=c(-10,20),lat=c(55,65))
y <- subset(Oslo,it=jan)
X <- EOF(subset(era40,it=jan))
ds <- DS(y,X)
plot(ds) # Looks OK

```

```
# Now we replace some values of y with missing data:
y2 <- y
set2na <- order(rnorm(length(y)))[1:50]
y2[set2na] <- NA
ds2 <- DS(y2,X)
plot(ds2)

## Use downscale results to fill in missin data:
y3 <- predict(ds2,newdata=X)
## Plot a subset of y based on dates in predicted y3
plot(subset(y,it=range(index(y3))),col=grey80,lwd=4)
points(as.station(predict(ds2)))
# The downscaled
lines(y3,lty=2)
```

---

DSE	<i>Downscale monthly climate variables and parameters for several stations.</i>
-----	---

---

## Description

Performs a complete downscaling job based on [DSensemble](#) for several stations and save the down-scaled results locally. DSE can be used to downscale climate variables, parameters or statistics for various stations. Each station is downscaled separately for the whole rcp/gcm ensemble and stored the results in a separte rda file. DSE() will also generate two files: an inventory file containing the meta data and some statisitcs for the successfully downscaled stations and log file containing encountered errors and a list of stations that have not been downscaled.

## Usage

```
DSE.default(cntr, src, param, FUN, lon, lat, path, rcp, biascorrect,reanalysis,
  email, save, out.dir, force, update, ...)
DSE.metno(x,...) # Downscale METNO stations
DSE.ecad(x,...)
DSE.ghcnd(x,...)
meta.dse(path) # update the inventory file with additional downscaled results.
```

## Arguments

stid	A string of characters as an identifier of the weather/climate station.
param	Parameter or element type or variable identifier. There are several core parameters or elements as well as a number of additional parameters. The parameters or elements are: prcp, pr = Precipitation (mm) tas, tavg = 2m-surface temperature (in degrees Celcius) tmax, tasmax = Maximum temperature (in degrees Celcius) tmin, tasmin = Minimum temperature (in degrees Celcius)
FUN	A mathematical transformation of the parameter e.g. "mean", "sd", "r1", etc ...

rcp	A character string corresponding to the name of the RCP experiment. Possible values for CMIP5 are rcp20, rcp45, rcp65, rcp85. For the CMIP3, they are: sresa1b, sresa2.
src	Source: limit the downscaling to a specific data set ("NARP", "NACD", "NORD-KLIMA", "GHCNM", "METNOM", "ECAD", "GHCND" and "METNOD")
stid	A string of characters as an identifier of the weather/climate station.
lon	A numeric vector containing the range of longitude values in the form of c(lon.min,lon.max) which corresponds to the predictor longitude extension
lat	A numeric vector containing the range of latitude values in the form of c(lat.min,lat.max) which corresponds to the predictor latitude domain extension
biascorrect	Logical value. If TRUE, a bias correction method is applied
save	Logical value. If TRUE, the results are save in rda files and stored locally in directory specified by path.
out.dir	A character string of the full path specifying the name of the output directory. If the directory does not exist, it will be created automaitcally.
cntr	A string or a vector of strings of the full name of the country. Select the stations from a specified country or a set of countries.
force	Logical value. If TRUE, the DSE is performed from scratch and all existing DSE files will be re-written. If FALSE, DSE will skip already downscaled stations and do the downscaling only for non existing file stations.
path	A character string corresponding to the full path name of the CMIP3/5 results.
reanalysis	A character string with the full path file name of the reanalysis to be used. Only netcdf files are accepted. update=
email	A character string with an email address. If specified, a notification email is sent when the downscaling is complete. The email text message contains a list of non-downscaled stations because of errors. If any error, please forward the email to abdelkadem@met.no
update	A logical value. If TRUE, <a href="#">meta.dse</a> is called and the inventory file is updated if additional DSE files are copied to output directory.
...	Additional arguments to be passed in the function

**Value**

A "zoo" "DSensemble" object

**Author(s)**

A. Mezghani, MET Norway

**See Also**

[DSensemble](#)

## Examples

```
## Not run:
DSE.metno(cntr = "NORWAY", src = "METNO", param = "t2m", nmin=100,
  lon = c(-10, 10), lat = c(-10, 10), FUN = "mean", path = "CMIP5.monthly",
  rcp = "rcp45", biascorrect = TRUE, reanalysis = "ERA40_t2m_mon.nc",
  email = NULL, save = TRUE, force = FALSE, verbose = FALSE,
  out.dir = "dse.100", update = FALSE)

## End(Not run)
```

---

DSensemble

*Downscale ensemble runs*


---

## Description

Downscales an ensemble of climate model runs, e.g. CMIP5, taking the results to be seasonal climate statistics. For temperature, the result hold the seasonal mean and standard deviation, whereas for precipitation, the results hold the wet-day mean, the wet-day frequency, and the wet/dry-spell statistics. The call assumes that netCDF files containing the climate model ensemble runs are stores in a file structure, linked to the path argument and the rcp argument.

These methods are based on [DS](#), and DSensemble is designed to make a number of checks and evaluations in addition to performing the DS on an ensemble of models. It is based on a similar philosophy as the old R-package 'clim.pact', but there is a new default way of handling the predictors. In order to attempt to ensure a degree of consistency between the downscaled results and those of the GCMs, a fist covariate is introduced before the principal components (PCs) describing the [EOFs](#).

DSensemble.pca is used to downscale a predictor represented in terms of PCA, and can reduce the computation time significantly. See Benestad et al. (2015) <http://dx.doi.org/10.3402/tellusa.v67.28326>.

The argument non.stationarity.check is used to conduct an additional test, taking the GCM results as 'pseudo-reality' where the predictand is replaced by GCM results interpolated to the same location as the provided predictand. The time series with interpolated values are then used as predictor in calibrating the model, and used to predict future values. This set of prediction is then compared with the interpolated value itself to see if the dependency between the large and small scales in the model world is non-stationary.

Other chekch include cross-validation ([crossval](#)) and diagnostics comparing the sample of ensemble results with the observations: number of observations outside the predicted 90-percent conf. int and comparing trends for the past.

## Usage

```
DSensemble(y,...)
DSensemble.default(y,path=CMIP5.monthly/,rcp=rcp45,...)
DSensemble.t2m(y,plot=TRUE,path="CMIP5.monthly/",
  predictor="ERA40_t2m_mon.nc",
  rcp="rcp45",biascorrect=FALSE,
```



```

        non.stationarity.check=FALSE,type=ncdf,
eofs=1:6,lon=c(-20,20),lat=c(-10,10),it=NULL,rel.cord=TRUE,
select=NULL,FUN="mean",FUNX="mean",
pattern="tas_Amon_ens_",
path.ds=NULL,file.ds="DSensemble.rda",
nmin=NULL,verbose=FALSE)
DSensemble.precip(y,plot=TRUE,path="CMIP5.monthly/",
rcp="rcp45",biascorrect=FALSE,
predictor="ERA40_pr_mon.nc",
non.stationarity.check=FALSE,
type=ncdf,
eofs=1:6,lon=c(-10,10),lat=c(-10,10),it=NULL,rel.cord=TRUE,
select=NULL,FUN="wetmean",
FUNX="sum",threshold=1,
pattern="pr_Amon_ens_",verbose=FALSE,nmin=NULL)
DSensemble.annual(y,plot=TRUE,path="CMIP5.monthly/",
rcp="rcp45",biascorrect=FALSE,
predictor="ERA40_t2m_mon.nc",
non.stationarity.check=FALSE,type=ncdf,
eofs=1:6,lon=c(-10,10),lat=c(-10,10),it=NULL,rel.cord=TRUE,
abscoords=FALSE,select=NULL,FUN=NULL,
FUNX="mean",threshold=1,
pattern="tas_Amon_ens_",verbose=FALSE,nmin=NULL)
DSensemble.season(y,season="djf",plot=TRUE,path="CMIP5.monthly/",
predictor="slp_mon.mean.nc",
rcp="rcp45",biascorrect=FALSE,
non.stationarity.check=FALSE,type=ncdf,
eofs=1:6,lon=c(-20,20),lat=c(-10,10),it=NULL,rel.cord=TRUE,
select=NULL,FUN="mean",FUNX="mean",
pattern="psl_Amon_ens_",
path.ds=NULL,file.ds=NULL,
nmin=NULL,verbose=FALSE)
DSensemble.mu(y,plot=TRUE,path="CMIP5.monthly/",
rcp="rcp45",biascorrect=FALSE,
predictor=list(t2m="data/ncep/air_mon.mean.nc",
olr="data/ncep/OLR_mon.mean.nc",
slp="data/ncep/slp_mon.mean.nc"),
non.stationarity.check=FALSE,type=ncdf,
eofs=1:16,lon=c(-30,20),lat=c(-20,10),it=NULL,rel.cord=TRUE,
select=NULL,FUN="wetmean",threshold=1,
pattern=c("tas_Amon_ens_","slp_Amon_ens_"),verbose=FALSE,nmin=365)
DSensemble.mu.worstcase(y,plot=TRUE,path="CMIP5.monthly/",
predictor="ERA40_t2m_mon.nc",
rcp="rcp45",biascorrect=FALSE,
lon=c(-20,20),lat=c(-10,10),it=NULL,rel.cord=TRUE,
select=NULL,FUN="wetmean",type=ncdf,
pattern="tas_Amon_ens_",verbose=FALSE)
DSensemble.pca(y,plot=TRUE,path="CMIP5.monthly/",

```

```
rcp="rcp45",biascorrect=FALSE,
predictor="ERA40_t2m_mon.nc",
non.stationarity.check=FALSE,
eofs=1:16,lon=c(-30,20),lat=c(-20,10), it=NULL,rel.cord=TRUE,
select=NULL,FUN="mean",rmtrend=TRUE,
FUNX="mean",threshold=1,type=ncdf,
pattern="tas_Amon_ens_",verbose=FALSE,
file.ds="DSensemble.rda",path.ds=NULL,nmin=NULL)
```

## Arguments

y	A station object.
plot	Plot intermediate results if TRUE.
path	The path where the GCM results are stored.
rcp	Which (RCP) scenario
biascorrect	TRUE, apply a bias adjustment using <a href="#">biasfix</a>
predictor	
non.stationarity.check	
eofs	Which EOFs to include in the step-wise multiple regression
rmtrend	TRUE: detrend before calibrating the regression model.
lon	Longitude range for predictor
lat	Latitude range for predictor
rel.cord	TRUE: use the range relative to predictand; FALSE use absolute range
it	Used to extract months or a time period. See <a href="#">subset</a> .
select	GCMs to select, e.g. <code>.subsample</code> the ensemble (1:3 selects the three first GCMs)
FUN	Function for aggregating the predictand (daily), e.g. <code>'mean'</code> , <code>'wetmean'</code>
threshold	Used together with FUN for some functions ( <code>'wetmean'</code> ).
nmin	Minimum number of day used in <a href="#">annual</a> used for aggregating the predictand/predictor
FUNX	Function for transforming the predictor, e.g. <code>'C.C.eq'</code> to estimate the saturation water vapour
type	Type of netCDF used in <a href="#">retrieve</a> for reading GCM data.
pattern	File name pattern for GCM data.
verbose	TRUE for checking and debugging the functions.
file.ds	Name of file saving the results.
path.ds	Path of file saving the results.

## Value

A `'dsensemble'` object - a list object holding DS-results.

Author(s)

R.E. Benestad and A. Mezghani

Examples

```
data(Oslo)
## Download NorESM1-M from climexp.knmi.nl in default directory
## (home directory for linux/mac users)
url <- "http://climexp.knmi.nl/CMIP5/monthly/tas/"
## Download NorESM1-ME from the rcp45
noresm <- "tas_Amon_NorESM1-M_rcp45_000.nc"
if (!file.exists(noresm))
  download.file(url=file.path(url,noresm), destfile=noresm,
               method = "auto",quiet=FALSE,mode = "w",cacheOK = TRUE)
fioesm <- "tas_Amon_FIO-ESM_rcp45_000.nc"
if (!file.exists(fioesm))
  download.file(url=file.path(url,fioesm), destfile=fioesm,
               method = "auto", quiet = FALSE, mode = "w",cacheOK = TRUE)
## Downscale 2m temperature
rcp4.5 <- DSensemble.t2m(Oslo,path=~ ,rcp=,pattern = "tas_Amon_",
biascorrect=TRUE, predictor = "~/air.2m.mon.mean.nc",plot=TRUE)

# Evaluation: (1) compare the past trend with downscaled trends for same
# interval by ranking and by fitting a Gaussian to the model ensemble;
# (2) estimate the probability for the counts outside the 90
# percent confidence interval according to a binomial distribution.

diagnose(rcp4.5, plot = TRUE, type = "target")
```

---

ele2param	<i>Dictionary and conversion tools between esd element identifier and variables names and specifications.</i>
-----------	---

---

Description

Converts between esd element/parameter identifier and variable names from different data sources.

Usage

```
ele2param(ele = 101, src = GHCND)
esd2ele(param=t2m,src=GHCND,verbose=FALSE)
```

Arguments

param,ele      Parameter or element identifier. There are several core parameters or elements as well as a number of additional parameters. The parameters or elements are :

PARAMETER	LONGNAME	ELE ID
auto	Automatic selection.	

prcp, pr, rr, precip	Precipitation (mm)	'601'
tas, tavg, t2m, t2	2m-surface temperature (in degrees Celcius)	'101'
tmax, tasmax	Maximum temperature (in degrees Celcius)	'111'
tmin, tasmin	Minimum temperature (in degrees Celcius)	'121'
slp, mslp	Mean sea level pressure (hPa)	'401'
cloud	Cloud cover (%)	'801'

**src** A character string for the acronym of the data source. The data sources are :

Data Source	Full name	Type
NACD	North Atlantic Climatological Dataset	Monthly
NARP	North Atlantic Research Programme ?	Monthly
NORDKLIM	Nordic co-operation within climate activities	Monthly
METNOM	MET Norway climate network	Monthly
GHCNM	Global historical climate network	Monthly
ECAD	European Climate Assessment & Dataset	Daily
GHCND	Global Historical Climate Network	Daily
METNOD	MET Norway climate network	Daily

### Value

A meta data matrix object with the glossary of the different variables or element identifiers as originally defined by each data source

### Author(s)

A. Mezghani, MET Norway

### Examples

```
# Eg.1 # Display the glossary of parameters or element identifiers for GHCND data source.
print(ele2param(ele=NULL,src=GHCND))
# Eg.2 # Display the glossary of parameters or element identifiers for all data sources.
print(ele2param())
# Eg.3 # Convert mean temperature parameter (param) to esd element (ele).
ele <- esd2ele(param=t2m)
print(ele)
```

## Description

Computes EOFs (a type of principal component analysis) for combinations of data sets, typically from gridded data, reanalysis and climate models results.

[ref: Benestad (2001), "A comparison between two empirical downscaling strategies", *Int. J. Climatology*, **vol 21**, Issue 13, pp.1645-1668. DOI 10.1002/joc.703]. and `mixFields` prepares for mixed-field EOF analysis [ref. Bretherton et al. (1992) "An Intercomparison of Methods for finding Coupled Patterns in Climate Data", *J. Climate*, **vol 5**, 541-560; Benestad et al. (2002), "Empirically downscaled temperature scenarios for Svalbard", *Atm. Sci. Lett.*, doi.10.1006/asle.2002.0051].

Uncertainty estimates are computed according to North et al. (1982), "Sampling Errors in the Estimation of Empirical Orthogonal Functions", *Mon. Weather Rev.*, **vol 110**, 699-706.

The EOFs are based on `svd`.

See the course notes from Environmental statistics for climate researchers <http://www.gfi.uib.no/~nilsg/kurs/notes/course.html> for a discussion on EOF analysis.

The method PCA is similar to EOF, but designed for parallel station series (e.g. grouped together with `merge`). PCA does not assume gridded values and hence does not weigh according to grid area. PCA is useful for downscaling where the spatial covariance/coherence is important, e.u involving different variables from same site, same variable from different sites, or a mix between these. For instance, PCA can be applied to the two wind components from a specific site and hence extract the most important wind directions/speeds.

## Usage

```
EOF(X,it=NULL,n=20,lon=NULL,lat=NULL,verbose=FALSE,...)
EOF.default(X,it=NULL,n=20,lon=NULL,lat=NULL,
            area.mean.expl=TRUE,verbose=FALSE,...)
EOF.field(X,it=NULL,n=20,lon=NULL,lat=NULL,
          area.mean.expl=TRUE,verbose=FALSE)
EOF.comb(X,it=NULL,n=20,lon=NULL,lat=NULL,area.mean.expl=TRUE,verbose=FALSE)
eof2field(x,it=NULL,is=NULL,anomaly=FALSE,verbose=FALSE)
PCA(X,...)
EOF.default(X,...)
PCA.station(X,na.action=fill,verbose=FALSE)
pca2station(X,lon=NULL,lat=NULL,anomaly=FALSE,what=pca,verbose=FALSE)
```

## Arguments

<code>X</code>	a 'field' or 'pca' object
<code>it</code>	see <code>subset</code>
<code>n</code>	number of EOFs
<code>is</code>	Spatial subsetting - see <code>subset.eof</code>
<code>lon</code>	set longitude range - see <code>t2m.ERAINT</code>
<code>lat</code>	set latitude range
<code>verbose</code>	TRUE - clutter the screen with messages
<code>area.mean.expl</code>	When TRUE, subtract the area mean for the domain and use as a the first co-variate before the PCs from the EOF analysis.

na.action	'fill' uses <a href="#">approx</a> to interpolate the NA-values before the PCA.
what	Default set to 'pca' for convertin the PCA-results, but also works on downscaled PCA results. Option 'xval' returns the station values for cross-validation results and 'test' returns the station values used in the cross-validation (same interval as for 'xval').

## Value

File containing an 'eof' object which is based on the 'zoo' class.

## Author(s)

R.E. Benestad

## Examples

```
# Simple EOF for annual mean SST:
sst <- sst.NCEP(lon=c(-90,20),lat=c(0,70))
SST <- aggregate(sst, year, mean, na.rm = FALSE)
eof.sst <- EOF(SST)
plot(eof.sst)

# EOF of July SST:
eof.sst7 <- EOF(sst,it=7)
plot(eof.sst7)

# common EOF for model
# Get some sample data, extract regions:
GCM <- t2m.NorESM.M()
gcm <- subset(GCM,is=list(lon=c(-50,60),lat=c(30,70)))
t2m.eraint <- t2m.ERAINT()
eraint <- subset(t2m.eraint,is=list(lon=c(-50,60),lat=c(30,70)))

OBS <- aggregate(eraint, by=year, mean, na.rm = FALSE)
GCM <- aggregate(gcm, by=year, mean, na.rm = FALSE)
OBSGCM <- combine(OBS,GCM,dimension=time)

ceof <- EOF(OBSGCM)
plot(ceof)

# Example for using PCA in downscaling
## nacd <- station(src=nacd)
## X <- annual(nacd)
X <- station(src=nacd)
nv <- function(x) sum(is.finite(x))
ok <- (1:dim(X)[2])[apply(X,2,nv) == dim(X)[1]]
X <- subset(X,is=ok)
pca <- PCA(X)
map(pca)

slp <- slp.NCEP(lon=c(-20,30),lat=c(30,70))
eof <- EOF(slp,it=1)
```

```

ds <- DS(pca,eof)
# ds is a PCA-object
plot(ds)

# Recover the station data:
Z <- pca2station(pca)
plot(Z,plot.type=multiple)

```

---

hotsummerdays

*Projection of hot and cold day statistics*


---

## Description

The functions `hotsummerdays`, `heatwavespells`, `coldwinterdays`, and `coldspells` estimate statistics for heatwaves/hot days or cold spells based on seasonal mean temperatures. The estimations are based on a regression analysis (GLM) between observed number of events or spell lengths and seasonal mean from station data. `nwetdays` estimates the number of days per year with precipitation amount exceeding a threshold values.

## Usage

```

coldwinterdays(x,y=NULL,dse=NULL,it=djf,threshold=0,
               verbose=FALSE,plot=TRUE,nmin=90,...)
coldspells(x,y=NULL,dse=NULL,it=jja,threshold=0,
           verbose=FALSE,plot=TRUE,nmin=90,...)
hotsummerdays(x, y=NULL,dse = NULL, it = "jja", threshold = 30,
               verbose = FALSE, plot = TRUE)
heatwavespells(x,y=NULL,dse=NULL,it=jja,threshold=30,
               verbose=FALSE,plot=TRUE,...)
nwetdays(x,y=NULL,dse=NULL,threshold=10,
          FUN=mu,verbose=FALSE,plot=TRUE,...)

```

## Arguments

<code>x</code>	station object, e.g. the temperature. Matches the element used in the <code>dsensemble</code> object 'dse'
<code>y</code>	station object which may be some statistics with dependency to <code>x</code> , e.g. snow depth.
<code>dse</code>	a <code>dsensemble</code> object. If <code>NULL</code> , then run <code>DSensemble</code>
<code>it</code>	Default season set for northern hemisphere.
<code>threshold</code>	Temperature threshold
<code>verbose</code>	TRUE for trouble shooting, debugging etc.
<code>plot</code>	TRUE - produce graphics

## Details

The estimation of these statistics makes use of general linear models (GLMs) and take the counts to follow the 'Poisson family' whereas the spall lengths belong to the geometric distribution. The seasonal mean temperature or annual wet-mean precipitation are used as independent variable.

## Author(s)

R.E. Benestad

## Examples

```
data(ferder)
data(dse.ferder)
hw <- hotsummerdays(ferder,dse.ferder,threshold=20)
```

---

iid.test

*iid test*

---

## Description

Test for whether a variable is independent and identically distributed (iid). Used in [daily.station.records](#).

Reference:

Benestad, R.E., 2003: How often can we expect a record-event? Climate Research. 23, 3-13 (pdf)

Benestad, R.E., 2004: Record values, nonstationarity tests and extreme value distributions, Global and Planetary Change, vol 44, p. 11-26

The papers are available in the pdf format from [http://regclim.met.no/results\\_iii\\_artref.html](http://regclim.met.no/results_iii_artref.html).

Note, gaps of missing data (NA) can bias the results and produce an under-count. The sign of non-iid behaviour is when the 'forward' analysis indicated higher number of record-events than the confidence region and the backward analysis gives lower than the confidence region.

Version 0.7: Added a test checking for dependencies based on an expected number from a binomial distribution and given the probability  $p1(n) = 1/n$ . This test is applied to the parallel series for one respective time (realisation), and is then repeated for all observation times. The check uses [qbinom](#) to compute a theoretical 95% confidence interval, and a number outside this range is marked with red in the 'ball diagram' (first plot). [pbinom](#) is used to estimate the p-value for the

## Usage

```
iid.test(x,plot=TRUE,Monte.Carlo=TRUE,N.test=200,reverse.plot.reverse=TRUE)
n.records(x)
```



**Arguments**

<code>x</code>	A data matrix or a vector.
<code>plot</code>	Flag: plot the diagnostics.
<code>Monte.Carlo</code>	Flag: for estimating confidence limits.
<code>N.test</code>	Number of Monre-Carlo runs.
<code>reverse.plot.reverse</code>	TRUE: plots reverse from right to left, else left to right..

**Value**

list: 'record.density' and 'record.density.rev' for the reverse analysis. The variables CI.95, p.val, and i.cluster (and their reverse equivalents '.rev') return the estimated 95% conf. int, p-value, and the location of the clusters (binomial).

**Author(s)**

R.E. Benestad

**Examples**

```
dat <- rnorm(100*30)
dim(dat) <- c(100,30)
iid.test(dat)
```

---

InfoGraphics

*InfoGraphics.*

---

**Description**

Wheel

Risk

prob - boxes with forseen outcomes - area proportional to probability

conf - confidence intervals and uncertainty - clouds...

vis

diagram

cumugram

prob

graph

**Usage**

```

vis(x,...)
diagram(x,...)
diagram.dsensemble(x,it=0,...)
diagram.station(x,...)
wheel(x,...)
wheel.station(x,y=NULL,new=TRUE,lwd=2,col=NULL,bg="grey90",verbose=FALSE,...)
wheel.spell(x,y=NULL,new=TRUE,lwd=2,col=NULL,verbose=FALSE,...)
cumugram(x,it=NULL,prog=FALSE,...)
climvar(x,FUN=sd,plot=TRUE,...)
colscal(n=30,col="bwr",alpha=NULL,test=FALSE)
seasevol(x,...)
seasevol.station(x,it=NULL,nv=25,...)
prob(x,...)
prob.default(x,...)
prob.station(x,y=NULL,is=1,...)
prob.station.precip(x,y=NULL,is=1,threshold=1,...)
graph(x,...)
graph.dsensemble(x,img=NULL,it=0,col=rgb(1,0.7,0.7,0.1),
                  lwd=5,xlim=NULL,ylim=NULL,add=FALSE,new=TRUE)
graph.list(x,img=NULL,it=0,
            col=c(rgb(1,1,0.5,0.05),rgb(1,0.5,0.5,0.05),rgb(0.5,1,0.5,0.05)),
            lwd=5,xlim=NULL,ylim=NULL,add=FALSE,new=TRUE)
graph.zoo(x,img=NULL,it=NULL,col=rgb(1,0.7,0.7,0.1),
          lwd=5,xlim=NULL,ylim=NULL,xlab=,ylab=,add=FALSE,new=TRUE)

```

**Arguments**

x                      a data object

**Value**

A field object

**Author(s)**

R.E. Benestad and A. Mezghanil

**See Also**

[map](#), [plot.station](#), [hist.spell](#)

**Examples**

```

data(bjornholt)
wheel(bjornholt)

z <- spell(bjornholt,threshold=1)
wheel(z)

```

---

is	<i>Test for .</i>
----	-------------------

---

**Description**

Computes different formulas

**Usage**

```
is.T(x)
is.precip(x)
is.field(x)
is.station(x)
is.eof(x)
is.pca(x)
is.cca(x)
is.trajectory(x)
is.daily(x)
is.monthly(x)
is.seasonal(x)
is.annual(x)
is.model(model, verbose=FALSE)
```

**Arguments**

x                      a data object

**Value**

Boolean

**Author(s)**

R. Benestad, MET Norway

**Examples**

```
data(ferder)
is.T(ferder)
```

map

*Plot maps for esd objects***Description**

Make map of geophysical data. These plot functions are S3 methods for esd objects.

**Usage**

```
<<<<<<< HEAD
map(x,it=NULL,is=NULL,new=TRUE,...)

map.default(x,FUN="mean",it=NULL,is=NULL,new=FALSE,
             projection="lonlat",xlim=NULL,ylim=NULL,zlim=NULL,n = 15,
             colbar=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
             type="p",cex=2,show=TRUE,h=0.6,v=1,pos=0.05),
             type=c("fill","contour"),gridlines=FALSE,lonR=NULL,
             latR=NULL,axiR=NULL,verbose=FALSE, ...)

map.matrix(x,new=TRUE,projection="lonlat",...)

map.station(x,it=NULL,is=NULL,new=TRUE,projection = "lonlat",
            xlim = NULL, ylim = NULL, zlim = NULL,
            col = "darkred", bg = "orange",
            type = NULL,gridlines= TRUE,
            lonR = NULL,latR=45,axiR=NULL,verbose=FALSE,
            FUN = NULL,
            colbar = list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
                          type="p",cex=2,h=0.6,v=1),
            cex=0.8,zexpr = "alt",
            cex.subset = 1, add.text.subset = FALSE, showall = FALSE,
            add.text = FALSE, height = NULL, width = NULL, cex.axis = 1,
            cex.lab = 0.6, pch = 21, from = NULL, to = NULL,
            showaxis = FALSE, border = FALSE,
            full.names = FALSE, full.names.subset = FALSE,
            text = FALSE, fancy = FALSE, na.rm = TRUE, show.val = FALSE,
            colorbar = TRUE, legend.shrink = 1, ...)

map.comb(x,it=NULL,is=NULL,new=TRUE,projection = "lonlat",
          xlim=NULL,ylim=NULL,zlim=NULL,n=15,
          colbar=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
                      type="p",cex=2,h=0.6,v=1),
          type=c("fill", "contour"),gridlines=FALSE,
          lonR=NULL,latR=NULL,axiR=NULL,verbose=FALSE,pattern=1,...)

map.eof(x,it=NULL,is=NULL,new=TRUE,projection="lonlat",
        xlim= NULL,ylim=NULL,zlim=NULL,n=15,
```

```

colbar=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
            type="p",cex=2,h=0.6,v=1),
type=c("fill","contour"),gridlines=FALSE,
lonR=NULL,latR=NULL,axiR=NULL,verbose=FALSE,pattern=1,...)

map.ds(x,it=NULL,is=NULL,new=TRUE,projection="lonlat",
      xlim=NULL,ylim=NULL,zlim=NULL,n=15,
      colbar=list(pal = NULL,rev=FALSE,n=10,breaks=NULL,
                  type="p",cex = 2,h=0.6,v=1,pos=0.05),
      type= c("fill", "contour"),gridlines=FALSE,
      lonR=NULL,latR=NULL,axiR = NULL,verbose = FALSE,...)

map.field(x,FUN="mean",it=NULL,is=NULL,new=TRUE,projection = "lonlat",
          xlim=NULL,ylim=NULL,zlim=NULL,
          colbar= list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
                      type="p",cex=2,h=0.6,v=1),
          type = c("fill", "contour"), gridlines = FALSE,
          lonR = NULL,latR = NULL, axiR = NULL, verbose = FALSE,
          na.rm = TRUE, ...)

map.corfield(x,it=NULL,is=NULL,new=TRUE,projection= "lonlat",
             xlim=NULL,ylim=NULL,zlim=NULL,n = 15,
             colbar=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
                         type="p",cex=2,h=0.6,v=1),
             type=c("fill", "contour"),gridlines=FALSE,
             lonR=NULL,latR=NULL,axiR = NULL,verbose = FALSE, ...)

map.trend(x,it=NULL,is=NULL,new=TRUE,projection="lonlat",
          xlim=NULL,ylim=NULL,zlim=NULL,n=15,
          colbar=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
                      type="p",cex=2,h=0.6,v=1),
          type=c("fill", "contour"),gridlines=FALSE,
          lonR=NULL,latR=NULL,axiR=NULL,verbose=FALSE,...)

map.pca(x,it=NULL,is=NULL,pattern=1,new=TRUE,projection="lonlat",
        xlim=NULL,ylim=NULL,zlim=NULL,n=15,
        colbar=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
                    type="p",cex=2,h=0.6,v=1),
        type=c("fill","contour"),gridlines=FALSE,verbose = FALSE, ...)

map.mvr(x,it=NULL,is=NULL,new=TRUE,projection="lonlat",
        xlim=NULL,ylim=NULL,zlim=NULL,n=15,
        colbar=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,pos=0.05,
                    type="p",cex=2,h=0.6,v=1),
        type=c("fill", "contour"),gridlines = FALSE, verbose = FALSE, ...)

```

```

map.cca(x, icca=1, it=NULL, is=NULL, new=FALSE, projection="lonlat",
        xlim=NULL, ylim=NULL, zlim=NULL, ##n=15,
        colbar1=list(pal=NULL, rev=FALSE, n=10, breaks=NULL, type="p",
                     cex=2, show=TRUE, h=0.6, v=1, pos=0.05), colbar2= NULL,
        type=c("fill", "contour"), gridlines=FALSE,
        lonR=NULL, latR=NULL, axiR=NULL, verbose=FALSE, cex=2, ...)

map.trajectory(x, it=NULL, is=NULL, type="paths",
               projection="sphere", verbose=TRUE, ...)

map.events(x, Y=NULL, it=NULL, is=NULL, xlim=NULL, ylim=NULL,
            param=NA, alpha=0.5, col="black", pch=13, lwd=4, cex=2,
            colbar=list(pal="budrd", rev=FALSE, n=10, breaks=NULL,
                        pos=0.05, show=TRUE, type="p", cex=2, h=0.6, v=1),
            projection="sphere", latR=NULL, lonR=NULL, new=TRUE,
            verbose=FALSE, ...)

lonlatprojection(x, it=NULL, is=NULL, new=FALSE, projection="lonlat",
                 xlim=NULL, ylim=NULL, zlim=NULL, n =15,
                 colbar=list(pal=NULL, rev=FALSE, n=10, breaks=NULL, pos=0.05,
                             type="p", cex=2, h=0.6, v=1),
                 type = c("fill", "contour"), geography=TRUE,
                 gridlines = FALSE, verbose = FALSE, fancy=FALSE, ...)

map.googleearth(x)

rotM(x=0, y=0, z=0)

gridbox(x, col, density = NULL, angle = 45)

map2sphere(x, it=NULL, is=NULL, new = TRUE,
            xlim=NULL, ylim=NULL, zlim = NULL,
            colbar= list(pal=NULL, rev=FALSE, n=10, breaks=NULL,
                         type="p", cex=2, h=0.6, v=1, pos=0.05),
            lonR = NULL, latR = NULL, axiR = 0,
            type = c("fill", "contour"), colorbar=TRUE,
            gridlines = TRUE, fancy = FALSE, verbose = FALSE, ...)

vec(x, y, it=10, a=1, r=1, ix=NULL, iy=NULL,
    projection=lonlat, lonR=NULL, latR=NULL, axiR=0, ...)
mask(x, land=FALSE)

```

### Arguments

x	the object to be plotted; in rotM, x holds a vector of x-coordinates.
FUN	The function to be applied on x before mapping (e.g. mean)
pattern	Which EOF pattern (mode) to plot
colbar	The colour scales defined through colscal. Users can specify the colour 'pal'*ette

	(‘pal’), the number of breaks (‘n’), values of ‘breaks’, and the position of the color bar from the main plot (‘pos’). The ‘rev’ argument, will produce a reversed color bar if set to TRUE. The other arguments (‘type’, ‘h’ and ‘v’) are more specific to <code>col.bar</code> and are used only if argument ‘fancy’ is set to TRUE (not yet finished).
<code>it</code>	see <a href="#">subset</a>
<code>is</code>	see <a href="#">subset</a>
<code>new</code>	TRUE: create a new graphic device.
<code>projection</code>	Projections: <code>c("lonlat", "sphere", "np", "sp")</code> - the latter gives stereographic views from the North and south poles.
<code>xlim</code>	see <a href="#">plot</a> - only used for ‘lonlat’ projection.
<code>ylim</code>	see <a href="#">plot</a> - only used for ‘lonlat’ projection.
<code>n</code>	The number of colour breaks in the color bar
<code>breaks</code>	graphics setting - see <a href="#">image</a>
<code>type</code>	graphics setting - colour shading or contour
<code>gridlines</code>	Only for the lon-lat projection
<code>lonR</code>	Only for the spherical projection - see <a href="#">map2sphere</a>
<code>latR</code>	Only for the spherical projection - see <a href="#">map2sphere</a>
<code>axiR</code>	Only for the spherical projection - see <a href="#">map2sphere</a>
<code>density</code>	
<code>y</code>	a vector of y coordinates
<code>z</code>	a vector of z coordinates
<code>pattern</code>	Selects which pattern (see <a href="#">EOF</a> , <a href="#">CCA</a> ) to plot
<code>geography</code>	TRUE: plot geographical features
<code>angle</code>	for hatching
<code>a</code>	used in <a href="#">vec</a> to scale the length of the arrows
<code>r</code>	used in <a href="#">vec</a> to make a 3D effect of plotting the arrows up in the air.
<code>ix</code>	used to subset points for plotting errors
<code>iy</code>	used to subset points for plotting errors
<code>colorbar</code>	Show the color bar in the map (default TRUE). If FALSE, the colorbar is not added into the map (ignored).
<code>cex.subset</code>	...
<code>add.text</code>	Add abbreviated location names.
<code>full.names</code>	Show the full name of the location.
<code>showall</code>	Default is set to FALSE
<code>showaxis</code>	If set to FALSE, the axis are not displayed in the plot.
<code>fancy</code>	If set to true, will use <a href="#">col.bar</a> instead of <code>image</code> to produce the colour bar
<code>text</code>	If TRUE, display text info on the map. The default is set to FALSE
<code>show.val</code>	Display the values of ‘x’ or ‘FUN(x)’ on top of the coloured map.
<code>legend.shrink</code>	If set, the size of the color bar is shrunk (values between 0 and 1)
<code>...</code>	further arguments passed to or from other methods.
<code>land</code>	if TRUE mask land, else mask ocean

**Value**

A field object

**Author(s)**

R.E. Benestad

**See Also**

[plot.station](#)

**Examples**

```
# Select stations in ss and map the geographical location of the selected stations with a zoom on Norway.
ss <- select.station(cntr="NORWAY",param="precip",src="GHCND")
map(ss, col="blue",bg="lightblue",xlim = c(-10,30) , ylim = c(50,70))

## Get NACD data and map the mean values
y <- station.nacd()
map(y,FUN=mean,colbar=list(pal=varid(y),n=10),cex=2)
```

---

map.trajectory

*Plot trajectory maps*


---

**Description**

Make different types of trajectory maps. Individual trajectories are mapped with `map.trajectory`. The number density can be visualised with `map.hexbin.trajectory` and `map.sunflower.trajectory` which are versions of [scatter.hexbin](#) and [scatter.sunflower](#) adapted to show trajectory density.

**Usage**

```
map.trajectory(x,it=NULL,is=NULL,type="paths",projection="sphere",
              lonR=10,latR=90,
              col=red,colmap=rainbow,alpha=0.3,pfit=FALSE,
              main=NULL,xlim=NULL,ylim=NULL,new=TRUE)
lonlat.trajectory <- function(x,xlim=NULL,ylim=NULL,col=blue,alpha=0.1,
                              lty=1,lwd=1,main=NULL,new=TRUE)
sphere.trajectory <- function(x,xlim=NULL,ylim=NULL,col=blue,alpha=0.1,
                              lty=1,lwd=1,lonR=0,latR=90,main=NULL,new=TRUE)
map.density.trajectory(x,dx=4,dy=2,it=NULL,is=NULL,
                      colbar=list(pal=precip,rev=TRUE,breaks=NULL,cex=2,h=0.6,v=1),
                      projection=sphere,latR=90,lonR=10,gridlines=FALSE,...)
```



**Arguments**

x	the trajectory object to be plotted.
it	A list or data.frame providing time index, e.g. month
is	A list or data.frame providing space index, e.g. station record
type	type of map: 'paths' shows trajectories; 'density' shows the spatial density of the trajectories)
col	color of trajectories
colmap	Colour scales, either as an output from <a href="#">rbg</a> or a single character string 'bwr' (blue-white-red) or 'rwb' ('red-white-blue')
new	TRUE: create a new graphics device
projection	Projections: c("lonlat", "sphere", "np", "sp") - the latter gives stereographic views from the North and south poles.
xlim	see <a href="#">plot</a> - only used for 'lonlat' projection
ylim	see <a href="#">plot</a> - only used for 'lonlat' projection
main	an overall title for the plot
lonR	Only for the spherical projection - see <a href="#">map2sphere</a>
latR	Only for the spherical projection - see <a href="#">map2sphere</a>
leg	logical. If TRUE, legend is shown.
alpha	factor modifying the opacity alpha; typically in [0,1]

**Author(s)**

K. Parding

**See Also**

[map](#) [map.events](#)

**Examples**

```
# plot storm tracks zoomed in on the north Atlantic and northern Europe
data(imilast.M03)
map.trajectory(imilast.M03,col="blue",alpha=0.1,
               projection=latlon,xlim=c(-60,60),ylim=c(30,90))

# spherical projection
map.trajectory(imilast.M03,col="blue",alpha=0.1,projection=sphere)

# plot number density for grid boxes of width 2 degrees and height 1 degree
map.hexbin.trajectory(imilast.M03,xlim=c(-60,60),ylim=c(30,90),dx=2,dy=1)
map.sunflower.trajectory(imilast.M03,xlim=c(-60,60),ylim=c(30,90),dx=2,dy=1)
```

Models	<i>Calibrated models.</i>
<div><div><b>Description</b></div><div><p>mu.eq.f.tx contains a regression model <math>\mu = f(tx)</math>, relating the wet-day mean (mu) to the mean daily maximum temperature - however, the input used in this model is the saturation vapour pressure according to the Clausius-Clapeyron equation, and 'f()' here is 'beta * C.C.eq()'.</p></div><div><b>Usage</b></div><div><pre>data(mu.eq.f.tx)</pre></div><div><b>Value</b></div><div><p>lm() object.</p></div><div><b>Author(s)</b></div><div><p>R.E. Benestad</p></div><div><b>Examples</b></div><div><pre>## Retrieve the model data(mu.eq.f.tx) ## Sample data - temperature from Ferder Lighthouse data(ferder) pre &lt;- data.frame(x=mean(C.C.eq(ferder),na.rm=TRUE)) ## Predict the wet-day mean based on mean temperatures predict(mu.eq.f.tx,newdata=pre)</pre></div></div>	
MVR	<i>Multi-variate regression</i>

**Description**

MVR solves the equation

$$Y = \Psi X$$

and estimates

$$\Psi$$

by inverting the equation. Predictions give the varlue of Y, given this matrix and some input. MVR is useful for data where Y contains several time series where the spatial coherence/covariance is important to reproduce. For instance, Y may be a combination of stations, the two wind components from one station, or a set of different elements from a group of stations.

**Usage**

```

MVR(Y,X,...)
MVR.default(Y,X,...)
MVR.field(Y,X,SVD=TRUE,LINPACK=FALSE)
MVR.pca(Y,X,SVD=TRUE,LINPACK=FALSE)
MVR.eof(Y,X,SVD=TRUE,LINPACK=FALSE)
predict.MVR(object, newdata=NULL, ...)

```

**Arguments**

Y	An object with climate data: field, eof, or pca.
X	Same as Y or any zoo object.
SVD	Use a singular value decomposition as a basis for the PCA.
i.eofs	Which EOFs to include in the CCA.
LINPACK	an option for <a href="#">svd</a> .
object	The result from CCA.
newdata	The same as X.

**Value**

A CCA object: a list containing a.m, b.m, u.k, v.k, and r, describing the Canonical Correlation variates, patterns and correlations. a.m and b.m are the patterns and u.k and v.k the vectors (time evolution).

**Author(s)**

R.E. Benestad

**Examples**

```

## Not run:
# Example for using EOF and MVR
slp <- slp.NCEP(lat=c(-40,40),anomaly=TRUE)
sst <- sst.NCEP(lat=c(-40,40),anomaly=TRUE)
eof.1 <- EOF(slp,mon=1)
eof.2 <- EOF(sst,mon=1)
mvr <- MVR(eof.1,eof.2)
plot(mvr)

# Example for using PCA and MVR
oslo <- station(src="NACD",loc="Oslo")
bergen <- station.nacd("Bergen")
stockholm <- station.nacd("Stockholm")
copenhagen <- station.nacd("Koebenhavn")
helsinki <- station.nacd("Helsinki")
reykjavik <- station.nacd("Stykkisholmur")
edinburgh <- station.nacd("Edinburgh")
debilt <- station.nacd("De_Bilt")
uccle <- station.nacd("Uccle")

```

```

tromso <- station.nacd("Tromsoe")
falun <- station.nacd("Falun")
stensele <- station.nacd("Stensele")
kuopio <- station.nacd("Kuopio")
valentia <- station.nacd("Valentia")
X <- combine(oslo,bergen,stockholm,copenhagen,helsinki,reykjavik,
            edinburgh,debilt,uccle,tromso,falun,stensele,kuopio,valentia)
pca <- PCA(X)
slp <- slp.NCEP(lon=c(-20,30),lat=c(30,70))
eof <- EOF(slp)
mvr <- MVR(pca,eof)
plot(mvr)

# Find the teleconnection pattern to the NAO
data("NAOI")
data("sunspots")
data("NINO3.4")
X <- merge(NAOI,sunspots,NINO3.4,all=FALSE)

mvr <- MVR(pca,X)

# Find the pattern for NAOI:
teleconnection <- predict(mvr,newdata= c(1,0,0))
map(teleconnection,cex=2)

## End(Not run)

```

---

Oslo

*Oslo monthly mean temperature time series*


---

## Description

Oslo temperature monthly record from 1837 up to now.

## Usage

```
data(Oslo)
```

## Format

The format is: 'zoo' series from 1837-01-01 to 2014-02-01  
 Data: atomic [1:2126] NaN NaN NaN 3.1 8.9 13.5 16 15.7 11.2 7.6 ...  
 - attr(\*, "location")= chr "Oslo"  
 - attr(\*, "variable")= chr "T2m"  
 - attr(\*, "unit")= chr "deg C"  
 - attr(\*, "longitude")= num 10.7  
 - attr(\*, "latitude")= num 59.9  
 - attr(\*, "altitude")= num 94  
 - attr(\*, "country")= chr "Norway"

```

- attr(*, "longname")= chr "temperature at 2m"
- attr(*, "station_id")= num 18700
- attr(*, "quality")= chr "homogenised"
- attr(*, "calendar")= chr "gregorian"
- attr(*, "source")= chr "Dr. Nordli, 2013, met.no"
- attr(*, "URL")= logi NA
- attr(*, "type")= chr "observation"
- attr(*, "aspect")= chr "original"
- attr(*, "reference")= chr "Nordli et al. (in progress). 'The Oslo Temperature series 1837-2012: Homogeneity testing and Climate Analysis'"
- attr(*, "info")= logi NA
- attr(*, "method")= chr "Blended reconstruction (1877-1936) and instrumental data (1937-)"
- attr(*, "history")=List of 3
..$ call :length 19 as.station.data.frame(oslo, loc = "Oslo", param = "T2m", unit = "deg C", lon =
10.7, lat = 59.9, alt = 94, cntr = "Norway", longname = "temperature at 2m", ...
.. .. attr(*, "srcref")=Class 'srcref' atomic [1:8] 85 1 90 104 1 104 85 90
.. .. .. attr(*, "srcfile")=Classes 'srcfilecopy', 'srcfile' <environment: 0x23b1980>
..$ timestamp : chr "Tue Dec 10 14:53:51 2013"
..$ sessioninfo:List of 3
.. ..$ R.version : chr "R version 3.0.2 (2013-09-25)"
.. ..$ esd.version: chr "esd_0.2-1"
.. ..$ platform : chr "x86_64-pc-linux-gnu (64-bit)"
Index: Date[1:2126], format: "1837-01-01" "1837-02-01" "1837-03-01" "1837-04-01" ...

```

## Details

Oslo surface temperature recorded on a monthly basis from 1837 up to 2012. It corresponds to a blended reconstruction (1837-1936) and instrumental data (1937-2012). An homogenisation procedure has been carried out on the data by [yvind Nordli](#) at MET Norway.

## Source

MET Norway

## References

Nordli et al. (in progress). 'The Oslo Temperature series 1837-2012: Homogeneity testing and Climate Analysis'

## Examples

```

data(Oslo)
## maybe str(Oslo) ; plot(Oslo) ...

```

---

PCA.trajectory	<i>Principle component analysis of trajectory objects.</i>
----------------	--

---

## Description

Computes principal component analysis for trajectory data, e.g., storm tracks. Add some reference and details about the method. The PCA is based on [svd](#).

## Usage

```
PCA.trajectory(X,neofs=20,param=c(lon,lat),anomaly=TRUE,verbose=FALSE)
pca2trajectory(X)
```

## Arguments

X	a 'trajectory' object
verbose	TRUE - clutter the screen with messages
anomaly	logical. If TRUE, subtract the first latitude/longitude from each trajectory.
param	parameters to include in principle component analysis.

## Author(s)

K. Parding

## Examples

```
# Simple EOF for annual mean SST:
data(imilast.M03)
x <- subset(imilast.M03,is=list(lon=c(-20,20),lat=c(50,70)))
# PCA of longitude and latitude
pca <- PCA.trajectory(x,param=c(lon,lat))
plot.pca.trajectory(pca)
map.pca.trajectory(pca,projection=latlon)

# latitude only
pca <- PCA.trajectory(x,param=c(lat))
plot.pca.trajectory(pca)
```

---

pcafill	<i>PCA-based missing-value filling</i>
---------	--

---

**Description**

Fills missing (station) values by predicting their values using multiple regression. The regression uses as input principal components from PCA from the same (group of station) data, but where series with missing data have been excluded. This makes sense for (station) data where most of the variability is accounted for by a few leading modes. This method is not expected to be useful when there are many large data gaps.

**Usage**

```
pcafill(X, insertmiss = 0, eofs = 1:4, test = FALSE, verbose = FALSE)
pcafill.test(X,N=100,max.miss=100, eofs = 1:4, verbose=FALSE)
fitpc(y,x,eofs=4)
```

**Arguments**

X	station data (group of stations)
insertmiss	Used for testing and evaluating. Missing data are introduced to test the predictive capability
eofs	Number of EOFs/PCAs to include in filling in. In many cases, it may be useful to keep this to a small set of values.
test	Extra test - debugging
verbose	Print diagnostics - debugging
N	Number of runs in Monte-Carlo simulation
max.miss	Maximum NAs to insert (insertmiss) in Monte-Carlo simulations
x	time series for calibrating regression analysis
y	PC input for regression analysis

**Details**

This function is handy for the downscaling of PCAs. See Benestad, R.E., D. Chen, A. Mezghani, L. Fan, K. Parding, On using principal components to represent stations in empirical-statistical downscaling, Tellus A 28326, accepted.

**Value**

The same as the input - station object with filled-in values

**Author(s)**

Rasmus Benestad

**See Also**

[PCA](#), [allgood](#)

**Examples**

```
download.file(http://files.figshare.com/2073466/Norway.Tx.rda,
              Norway.Tx.rda)
load(Norway.Tx.rda)
X <- annual(Tx,FUN=mean,nmin=200)
ok<- apply(X,1,nv)
X <- subset(X,it=ok > 0)
Y <- pcafill(X)

plot(PCA(Y))
plot(c(coredata(Y)),c(coredata(X)))

## Monte-Carlo test with random selection of data points set to NA:
Y.test <- pcafill.test(X,max.miss=10,eofs=1:3)
cor(Y.test)
```

---

plot

*Plot esd objects*

---

**Description**

These plot functions are S3 methods for esd objects, based on plot.

**Usage**

```
nam2expr(x)
plot.station(x, plot.type = "single", new = TRUE, lwd = 3, type = "l",
             pch = 0, main = NULL, col = NULL, xlim = NULL, ylim = NULL,
             xlab = "", ylab = NULL, errorbar = TRUE, legend.show = FALSE,
             map.show = TRUE, map.type = "points", mar = c(4.5, 4.5, 0.75,
                 0.5), alpha = 0.3, verbose = FALSE, ...)
plot.eof(x,new=TRUE,xlim=NULL,ylim=NULL,pattern=1,what=c("pc","eof","var"),...)
plot.eof.field(x,new=TRUE,xlim=NULL,ylim=NULL,pattern=1,what=c("pc","eof","var"),...)
plot.eof.comb(x,new=TRUE,xlim=NULL,ylim=NULL,
              pattern=1,col=c("red"),what=c("pc","eof","var"),...)
plot.eof.var(x,new=TRUE,xlim=NULL,ylim=NULL,pattern=20,...)
plot.ds(x,plot.type="single",what=c("map","ts"),new=TRUE,
        lwd=3,type=b,pch=0,main=NULL,col=NULL,
        xlim=NULL,ylim=NULL,xlab="",ylab=NULL,...)
plot.ds.pca(y,pattern=1,verbose=FALSE,
            colbar1=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,type="p",cex=2,show=TRUE,
                h=0.6, v=1,pos=0.05),colbar2=NULL,...)
plot.ds.eof(y,pattern=1,
```



```

colbar1=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,type="p",cex=2,show=TRUE,
             h=0.6, v=1,pos=0.05),colbar2=NULL,verbose=FALSE,...)
plot.cca(x,icca=1,colbar1=list(pal=NULL,rev=FALSE,n=10,breaks=NULL,type="p",cex=2,show=TRUE,
                               h=0.6, v=1,pos=0.05),colbar2=NULL,verbose=FALSE,...)
plot.field(x,is=NULL,it=NULL,FUN="mean",...)
plot.spell(x)
plot.diagnose(x,...)
plot.diagnose.comb.eof(x,xlim=NULL,ylim=NULL,verbose=FALSE,add=FALSE,...)
plot.diagnose.matrix(x,xlim=NULL,ylim=NULL,verbose=FALSE,...)
plot.diagnose.dsensemble(x,map=TRUE,verbose=FALSE,new=TRUE,cex=NULL)
plot.xval(x,...)
plot.dsensemble(x,pts=FALSE,showci=TRUE,showtrend=TRUE,legend=TRUE,it=0,
                envcol=c(1,0,0,0.2),...)
plot.nevents(x,verbose=FALSE,...)
vis.map(x,col=red,map.type=points,add.text=FALSE)

```

### Arguments

x	the object to be plotted
pattern	Which EOF pattern (mode) to plot
col	Colour
icca	Which CCA pattern to plot
is	For subsetting in space - See <code>link{subset}</code> , but can also be a station value and if provided, the plotting will involve an interpolation to the same coordinates as defined by <code>is</code> .
it	For subsetting in time - See <code>link{subset}</code> . 'it=0' returns the annual means (mean of DJF + MAM + JJA + SON)
FUN	function
map.show	Show a small map of the location
map.type	

### Value

A field object

### Author(s)

R.E. Benestad

### See Also

[plot](#)

**Examples**

```

# Example: use aggregate to compute annual mean temperature for Svalbard:
data(Svalbard)
year <- as.numeric( format(index(Svalbard), %Y) )
y <- aggregate(Svalbard, by=year, FUN=mean, na.rm = FALSE)
plot(y)

# Example with downscaling:
lon <- c(-12,37)
lat <- c(52,72)
t2m <- t2m.ERA40(lon=lon,lat=lat)
data(Oslo)
ds <- DS(Oslo,t2m)

# Plot the results for January month
# plot(subset(ds,it=Jan))

# Plot the residuals:
residual <- as.residual(ds)
obs <- as.anomaly(as.calibrationdata(ds))

plot.zoo(obs,lwd=2)
lines(residual,col="red")

print("Global climate model simulation NorESM")
T2m <- t2m.NorESM.M(lon=lon,lat=lat)

# Plot the global mean of the field:
plot(T2m)
# Plot area mean of a sub region
plot(T2m,is=list(lon=c(0,10),lat=c(60,70)))

# Plot interpolated results corresponding to ferder
data(ferder)
plot(T2m,ferder)

# Plot Hovmuller diagram: Not working ...
## plot(T2m,is=list(lon=0))

print("Extract a subset - the January month")
x <- subset(t2m,it=1)
X <- subset(T2m,it=1)

print("Combine the fields for computing common EOFs:")
XX <- combine(x,X)

print("Compute common EOFs")
eofxx <- EOF(XX)
plot(eofxx)

print("Downscale the January mean temperature")

```

```
ds.jan <- DS(Oslo,eofxx)
plot(ds.jan)
```

---

predict.ds

*Prediction based on DS model*


---

## Usage

```
predict.ds(x, newdata = NULL, addnoise = FALSE, n = 100)
predict.ds.eof(x, newdata = NULL, addnoise = FALSE, n = 100)
predict.ds.comb(x, newdata = NULL, addnoise = FALSE, n = 100)

predict.mvr(object, newdata = NULL, ...)

project.ds(x, newdata = NULL, addnoise = FALSE, n = 100) # Not yet finished
```

## Arguments

x	A ds object
newdata	An eof object containing the new data sets on which the prediction is made.
addnoise	If TRUE, will add an attribute called "noise" to the ouput based on WG
n	Number of runs to be generated, used only if addnoise is set to TRUE

## Details

'predict' is similar to the predict function in R  
 'project' returns projection of climate

## Value

Predicted ds values.

## Author(s)

A. Mezghani

## See Also

[DS](#)

## Examples

```
# Get predictor
## Get reanalysis
X <- t2m.ERA40(lon=c(-40,50),lat=c(40,75))
## Get Gcm output
Y <- t2m.NorESM.M(lon=c(-40,50),lat=c(40,75))
## Combine
XY <- combine(X,Y)
# Compute common eof for January
ceof <- EOF(XY,it=jan)
# Get predictand
data(Oslo)
# Do the downscaling
ds <- DS(Oslo,ceof)
# Plot ds results
plot(ds)
# Do the prediction
ds.pre <- predict.ds(ds)
#Plot predicted results based on ds object
plot(ds.pre)
# Display the attribute "aspect"
attr(ds.pre, "aspect")
```

---

regrid

Regrid

---

## Description

Fast transform data from one longitude-latitude grid to another through bi-linear interpolation. The regridding is done by first calculating a set of weights. This is a "QUICK & DIRTY" way of getting approximate results. More sophisticated methods exist (e.g. Kriging - LatticeKrig).

Let  $X(i,j)$  be a  $i$ - $j$  matrix containing the data on a grid with  $i$  longitudes and  $j$  latitudes. We want to transform this to a different grid with  $k$  longitudes and  $l$  latitudes:

$X(i,j) \rightarrow Y(k,l)$

First the routine computes a set of weight, then performs a matrix multiplication to map the original data onto the new grid. The weights are based on the distance between points, taking longitude & latitude and use `distAB()` to estimate the geographical distance in km.

The matrix operation is:  $Y = \text{beta} X$

beta is a matrix with dimensions  $(i*j,k*l)$

$(Y(1,1) \text{ } (beta(1,1), beta(2,1), beta(3,1), \dots) \text{ } (X(1,1) \text{ } (Y(1,2) \text{ } (beta(1,2), beta(2,2), beta(3,2), \dots) \text{ } (X(1,2) \text{ } (\dots) \text{ } (beta(1,3), beta(2,3), beta(3,3), \dots) \text{ } (X(1,3) \text{ } )$

Most of the elements in Beta are zero!

**Usage**

```

regrid.weights(xo,yo,xn,yn,verbose)
sparseMproduct(beta,x)
regrid(x,is,...)
regrid.default(x,is,verbose=FALSE,...)
regrid.field(x,is,verbose=FALSE,...)
regrid.matrix(x,is,verbose=FALSE,...)
regrid.eof(x,is,verbose=FALSE)
nearest(x,is,...)
nearest.station(x,is)
nearest.field(x,is)

```

**Arguments**

xo	Old x-coordinates (longitudes)
yo	Old y-coordinates (latitudes)
xn	New x-coordinates (longitudes)
yn	New y-coordinates (latitudes)
beta	The matrix of interpolation weights
x	a field object.
is	A list holding the coordinates xn and yn, a field object, an eof object, or a station object - for the latter three, the field x is interpolated to the longitude/latitude held by is.
verbose	Clutter the screen.

**Value**

A field object

**Author(s)**

R.E. Benestad and A. Mezghanil

**Examples**

```

# Use regrid to interpolate to station location:
t2m <- t2m.ERAINT()
data(Oslo)
z.oslo <- regrid(t2m,is=Oslo)
plot(Oslo)
lines(z.oslo)

# Regrid t2m onto the grid of the gcm
gcm <- t2m.NorESM.M()
Z <- regrid(t2m,gcm)
map(Z)

# Example using regrid on a matrix object:

```

```

t2m.mean <- as.pattern(t2m,FUN=mean)
z <- regrid(t2m.mean,is=list(seq(min(lon(t2m)),max(lon(t2m)),by=0.5),
                             seq(min(lat(t2m)),max(lat(t2m),by=0.5))))
image(lon(z),lat(z),z)
# Add land borders on top
data(geoborders)
lines(geoborders)

## Regrid station data using weights defined by the distance of the 4
## nearest stations: quick and dirty method

if (!file.exists("stationsVALUE_exp1a.rda")) {
  download.file("http://files.figshare.com/2085591/value_predictands4exp1a.R", "value_predictands4exp1a.R")
  source("value_predictands4exp1a.R")
}

load("stationsVALUE_exp1a.rda")
TX <- regrid(Tx,is=list(lon=seq(-8,30,by=1),lat=seq(40,60,by=0.5)))
map(TX)

```

---

retrieve

---

*Retrieve field data from a netcdf file.*


---

## Description

Retrieve data from a netcdf file and return a zoo field object with attributes. `retrieve` assumes data on a regular lon-lat grid and `retrieve.rcm` reads data on irregular (rotated) grid. typically output from RCMs.

## Usage

```

retrieve.default(ncfile,type="ncdf",verbose=FALSE,...)
retrieve.ncdf(ncfile = ncfile, path = path , param = "auto",
              lon = NULL, lat = NULL, lev = NULL, it = NULL,
              miss2na = TRUE, greenwich = TRUE,
              verbose = FALSE , plot = TRUE)
retrieve.ncdf4(ncfile = ncfile, path = path , param = "auto",
               lon = NULL, lat = NULL, lev = NULL, it = NULL,
               miss2na = TRUE, greenwich = TRUE,
               verbose = FALSE , plot = TRUE)
retrieve.rcm(ncfile,param=NULL,is=NULL,it=NULL,verbose=FALSE)

summary.ncdf4(ncfile, verbose=FALSE)
check.ncdf4(ncid = ncid, param = "auto" , verbose = FALSE)
check.ncdf(ncid = ncid, param = "auto" , verbose = FALSE)

```

**Arguments**

<code>ncfile</code>	A character string of full path netcdf file name (include the path if necessary) or any object of class 'ncdf' or 'ncdf4'.
<code>type</code>	The type of the NetCDF format to be returned as "ncdf" or "ncdf4".
<code>ncid</code>	An object of class 'ncdf' or 'ncdf4'.
<code>lon</code>	Numeric value of longitude for the reference point (in decimal degrees East) or a vector containing the range of longitude values in the form of <code>c(lon.min,lon.max)</code>
<code>lat</code>	Numeric value of latitude for the reference point (in decimal degrees North) or a vector containing the range of latitude values in the form of <code>c(lat.min,lat.max)</code>
<code>lev</code>	Numeric value of pressure levels or a vector containing the range of pressure level values in the form of <code>c(lev.min,lev.max)</code>
<code>it</code>	Numerical or date values of time or a vector containing the range of values in the form of <code>c(start,end)</code> . Date format should be in the form of "YYYY-MM-DD".
<code>param</code>	Parameter or element type. There are several core parameters or elements as well as a number of additional parameters. The parameters or elements are: <code>auto</code> = automatic selection. <code>precip</code> , <code>prcp</code> , <code>pr</code> = Precipitation (mm) <code>tas</code> , <code>tavg</code> = 2m-surface temperature (in degrees Celcius) <code>tmax</code> , <code>tasmax</code> = Maximum temperature (in degrees Celcius) <code>tmin</code> , <code>tasmin</code> = Minimum temperature (in degrees Celcius)
<code>plot</code>	Logical value. if, TRUE provides a map.
<code>greenwich</code>	Logical value. If FALSE, convert longitudes to -180E/180E or centre maps on Greenwich meridian (0 deg E).
<code>ncdf.check</code>	Logical value. If TRUE, performs a quick check of the ncfile contents
<code>miss2na</code>	Logical value. If TRUE missing values are converted to "NA"
<code>verbose</code>	Logical value defaulting to FALSE. If FALSE, do not display comments (silent mode). If TRUE, displays extra information on progress.

**Value**

A "zoo" "field" object with additional attributes used for further processing.

**Author(s)**

A. Mezghani

**See Also**

[test.retrieve.ncdf4](#).

**Examples**

```
#\dontrun{
# Download air surface temperature (tas) for the NorESM1-ME model
# output prepared for CMIP5 RCP4.5 and for run r1i1p1 from the climate
# explorer web portal (http://climexp.knmi.nl) and store the file into the
# local machine, e.g. temporary folder /tmp (Size ~96Mb) using the following
# command if needed. Otherwise, specify a netcdf file to retrieve data from.
```

```

url <- "http://climexp.knmi.nl/CMIP5/monthly/tas"
noresm <- "tas_Amon_NorESM1-ME_rcp45_000.nc"
# download.file(url=file.path(url,noresm), destfile=noresm, method = "auto", quiet = FALSE,mode = "w",cacheOK = TRUE)
# Retrieve the data into "gcm" object
gcm <- retrieve(ncfile=file.path(~,noresm),param="tas",lon=c(-20,30),lat=c(40,90),plot=TRUE)
# Download the air surface temperature (tas) for RCP 4.5 scenarios and
# NorESM1-ME model from the climate explorer and store it in destfile.
# Compute the anomalies
gcm.a <- as.anomaly(gcm,ref=c(1960:2001))
map(gcm.a,projection="sphere")
#}

```

---

rtools	<i>Simple and handy functions. lag.station and lag.station are wrap-around functions for lag.zoo that maintains all the attributes. attrcp(x,y) passes on attributes from x to y and returns the y with new attributes.</i>
--------	---

---

## Usage

```

as.decimal(x = NULL)
cv(x,na.rm=TRUE)
nv(x)
q5(x)
q95(x)
q995(x)
trend.coef(x)
trend.pval(x)
lag.station(x,...)
lag.field(x,...)
exit()
ndig(x)
filt(x,...)
filt.default(x,n,type=ma,lowpass=TRUE)
figlab(x,xpos=0.001,ypos=0.001)
attrcp(x, y, ignore = c("name", "model", "n.apps", "appendix","dimnames"))

```

## Arguments

x	A data.frame or a coredata zoo object.
na.rm	If TRUE, remove NA's from data
type	'ma' for moving average (box-car), 'gauss' for Gaussian, 'binom' for binomial filter, 'parzen' for Parzen filter, 'hanning' for Hanning filter, or 'welch' for Welch filter.
lowpass	True for smoothing, otherwise the highpass results is returned



**Details**

'as.decimal' converts between degree-minute-second into decimal value.

'cv' computes the coefficient of variation.

'nv' count the number of valid data points.

'q5','q95' and 'q995' are shortcuts to the 5%, 95%, and 99.5% percentiles.

'trend.coef' and 'trend.pval' return the coefficient and the p-value of the linear trend.

'exit' is a handy function for exiting the R session without saving.

'figlab' is a handy function for labelling figures (e.g. 'Figure 1')

'ndig' estimates the number of digits for round(x,ndig), e.g. in scales for plotting.

**Value**

as.decimal	Decimal value
trend.coef	Linear trend per decade

**Author(s)**

A. Mezghani

**Examples**

```
## Monthly mean temperature at Oslo - Blindern station
data(Oslo)
## Compute the linear trend and the p-value on annual aggregated values
tr <- trend.coef(coredata(annual(Oslo)))
pval <- trend.pval(coredata(annual(Oslo)))
```

---

scatter.hexbin	<i>Produce a binned scatter plot with a hexagon grid</i>
----------------	--

---

**Description**

Multiple points are plotted as hexagons of different sizes or colors such that overplotting is visualized instead of accidental and invisible.

**Usage**

```
scatter.hexbin(x, y, new = TRUE, scale.col=TRUE, scale.size=TRUE, Nmax = NULL, dx = NULL, dy = NULL, xgr1
```

**Arguments**

<code>x, y</code>	the <code>x</code> and <code>y</code> arguments provide the x and y coordinates for the plot. Any reasonable way of defining the coordinates is acceptable. See the function <code>xy.coords</code> for details.
<code>scale.col</code>	logical. If TRUE, a color scale represents the count in each grid point. If FALSE, all markers are the same color.
<code>scale.size</code>	logical. If TRUE, the marker size represent the count in each grid point. If FALSE, all markers are the same size.
<code>colmap</code>	color scale of markers if <code>scale.col</code> is TRUE
<code>col</code>	face color of markers if <code>scale.col</code> is FALSE
<code>border</code>	border color of markers if <code>scale.col</code> is FALSE
<code>Nmax</code>	minimum count in grid points with the largest hexagonal marker. used only if <code>scale.size</code> is TRUE.
<code>dx, dy</code>	circumradius of the hexagonal markers in the x and y direction.
<code>xgrid, ygrid</code>	the first row and column of the hexagonal grid. If 'xgrid' is of length 2, it is interpreted as the edges of a grid of hexagons of circumradius 'dx' and 'dy'.
<code>leg</code>	logical. If TRUE, shows legend.
<code>xlim</code>	limits of the x-axis
<code>ylim</code>	limits of the y-axis
<code>xlab</code>	a label for the x-axis
<code>ylab</code>	a label for the y-axis

**Author(s)**

Kajsa Parding

**See Also**[scatter.sunflower](#)**Examples**

```

x <- sample(seq(1,5,1e-2),size=5000,replace=TRUE)
y <- sample(seq(5,10,1e-2),size=5000,replace=TRUE)
scatter.hexbin(x,y)

x <- station.metnod(stid=39100,param="Tmax")
y <- station.metnod(stid=39100,param="Tmin")
OK <- (!is.na(x) & !is.na(y))
x <- x[OK]; y <- y[OK]
scatter.hexbin(x,y,
  scale.size=TRUE,scale.col=TRUE,Nmax=500,colmap="heat.colors",
  dx=2,dy=1.5,xlim=c(-20,35),ylim=c(-25,25),xlab="Tmax",ylab="Tmin")

```

---

scatter.sunflower	<i>Produce a sunflower scatter plot with a hexagonal grid</i>
-------------------	---

---

## Description

Multiple points are plotted as `<e2><80><98>sunflowers<e2><80><99>` with multiple leaves (`<e2><80><98>petals<e2><80><99>`) such that overplotting is visualized instead of accidental and invisible.

## Usage

```
scatter.sunflower(x, y, petalsize = 7, dx = NULL, dy = NULL, xgrid = NULL, ygrid = NULL, xlim = NULL, ylim = NULL)
```

## Arguments

<code>x, y</code>	the <code>&lt;e2&gt;&lt;80&gt;&lt;98&gt;x&lt;e2&gt;&lt;80&gt;&lt;99&gt;</code> and <code>&lt;e2&gt;&lt;80&gt;&lt;98&gt;y&lt;e2&gt;&lt;80&gt;&lt;99&gt;</code> arguments provide the x and y coordinates for the plot. Any reasonable way of defining the coordinates is acceptable. See the function <code>&lt;e2&gt;&lt;80&gt;&lt;98&gt;xy.coords&lt;e2&gt;&lt;80&gt;&lt;99&gt;</code> for details.
<code>petalsize</code>	counts per petal of the larger sunflowers
<code>xgrid, ygrid</code>	the first row and column of the hexagonal grid. If 'xgrid' is of length 2, it is interpreted as the first and last point of a grid with distance 'dx' between the intermediate points.
<code>xlim</code>	limits of the x-axis
<code>ylim</code>	limits of the y-axis
<code>xlab</code>	a label for the x-axis
<code>ylab</code>	a label for the y-axis
<code>leg</code>	logical. If TRUE, shows legend.
<code>rotate</code>	logical. If TRUE, randomly rotates petals in sunflowers
<code>alpha</code>	factor modifying the opacity alpha; typically in [0,1]
<code>leg.loc</code>	location of legend; "upper left", "upper right", "lower left", or "lower right"

## Author(s)

Kajsa Parding

## See Also

[scatter.smooth](#), [scatter.hexbin](#)

**Examples**

```

x <- sample(seq(1,5,1e-2),5000,replace=TRUE)
y <- sample(seq(5,10,1e-2),5000,replace=TRUE)
scatter.sunflower(x,y)

x <- station.metnod(stid=39100,param="Tmax")
y <- station.metnod(stid=39100,param="Tmin")
OK <- (!is.na(x) & !is.na(y))
x <- x[OK]; y <- y[OK]
scatter.sunflower(x,y,petalsize=40,
  dx=2,dy=2,xlim=c(-20,35),ylim=c(-25,25),
  xlab="Tmax",ylab="Tmin")

```

---

spatial.avg.field	<i>Spatial Average of a Field Object.</i>
-------------------	---

---

**Description**

Computes the spatial average of a field object and return a zoo time series object.

**Usage**

```
spatial.avg.field(x,...)
```

**Arguments**

x	A zoo field object with two (longitude, latitude) or three dimensions (longitude, latitude, time)
---	---

**Value**

A "zoo" "time series" object

**Author(s)**

A. Mezghani, MET Norway

**See Also**

[retrieve.ncdf4](#)

## Examples

```
## Not run:
# Consider the "gcm" object from the e.g. in \link{retrieve.ncdf4}
# Compute the spatial average along lon and lat in gcm2
gcm2 <- spatial.avg.field(gcm)
# keep all attributes in gcm2
gcm2 <- attrcp(gcm,gcm2)
# Compute the annual mean
gcm.am <- as.annual(gcm2,FUN=mean,na.rm=TRUE)
# keep all attributes in gcm.am
gcm.am <- attrcp(gcm2,gcm.am)
year <- index(gcm.am)
# Compute the anomalies relative to the period 1986-2005
agcm.ave <- gcm.am-mean(gcm.am[is.element(as.numeric(year),c(1986:2005))])
agcm.ave <- attrcp(gcm.am,agcm.ave)
# plot anomaly time series of the global mean temperature
frame.metno(agcm.ave,col="black",cex.lab=0.75,cex.axis=0.75) ! Should be
  updated !
# Add vertical margin text with y-label
mtext("Anomaly values relative to 1986-2005",side=2,line=2,cex=0.75,las=3)

## End(Not run)
```

---

spell

*Spell statistics*


---

## Description

Statistics of spell durations (consecutive wet and dry days), e.g. dry and wet periods or duration of extremes.

exceedance estimates statistics for peak-over-threshold, and nevents returns the number of events with exceeding values (e.g. the number of rainy days  $X > 1$  mm/day). wetfreq returns the wet-day frequency (a fraction) and wetmean wet-day mean.

## Usage

```
spell(x, threshold, ...)
spell.default(x, threshold, upper=150, ...)
spell.station(x, threshold, upper=150, ...)
hist.spell(x, ...)
count(x, threshold=1, fraction=FALSE)
exceedance(x, threshold=1, fun=mean, ...)
exceedance.default(x, threshold=1, fun=mean, ...)
exceedance.station(x, threshold=1, fun=mean, ...)
exceedance.field(x, threshold=1, fun=mean, ...)
nevents(x, threshold=1)
wetfreq(x, threshold=1)
wetmean(x, threshold=1)
```

```
HDD(x,x0=18,na.rm=TRUE)
CDD(x,x0=22,na.rm=TRUE)
GDD(x,x0=10,na.rm=TRUE)
coldwinterdays(x,dse=NULL,threshold=0,verbose=FALSE,plot=TRUE)
```

### Arguments

x	station or field object
threshold	threshold value
upper	upper limit for maximum length - ignore any above this because they are likely erroneous
fraction	TRUE: divide the number of counts by number of samples
fun	function

### Value

Station or field objects

### Author(s)

R.E. Benestad and A. Mezghanil

### See Also

plot

### Examples

```
# Example 1 :
precip <- station.metnod(stid="18700",param="precip")
x <- spell(precip,threshold=.1)
x.ann <- annual(x,FUN="max")
plot(x.ann,plot.type="multiple")
# Example 2 :
x11() ; plot(x)

# Growing degree days:
data(ferder)
plot(as.seasons(ferder,FUN=GDD))

# Mild winter days - number of days in the winter season with
# above freezing temperatures
data(ferder)
try(coldwinterdays(ferder))
```

**Description**

After von Storch & Zwiers (1999), Statistical Analysis in Climate Research, p. 312

**Usage**

```
SSA(x,m,plot=TRUE,main="SSA analysis",sub="",LINPACK=TRUE,
    param = "t2m", anom = TRUE,i.eof=1)
plotSSA(ssa,main="SSA analysis",sub="")
```

**Arguments**

x	A station or eof object.
m	Window length.
plot	Flag: plot the diagnostics.
LINPACK	'TRUE': svd; 'FALSE':La.svd
main	main title (see link{plot}).
sub	subtitle (see link{plot}).
ssa	An 'SSA' object returned by SSA().
param	Which parameter ("daily.station.record") to use: "precip", "t2m" or other.
anom	TRUE if analysis on anomalies
i.eof	If x is an eof-object, which PC to use.

**Value**

A SSA object: An link{svd} object with additional parameters: m (window length), nt (original length of series), Nm (effective length of series= nt - m), anom (FLAG for use of anomaly), param (name of parameter, typically 'precip' or 't2m'), station (the station object to which SSA is applied).

**Author(s)**

R.E. Benestad

**Examples**

```
## Not run:
data(DNMI.t2m)
eof.1 <- EOF(DNMI.t2m,mon=1)
pop <- POP(eof.1)

## End(Not run)
```

---

station

---

*Retrieve meta data and data from observational weather stations.*


---

## Description

Retrieve station record from a given data source.

`allgood` and `clean.station` provide two filters for extracting stations with good data (discarding missing values). `allgood` will not leave any NA's whereas `clean.station` provides a more 'gentle' filtering.

`station.sonel`, `station.gloss`, and `station.newlyn` read sea level from tidal gauges in France (SONEL), on a global scale (GLOSS) and for a single station (sub-daily data) in the UK (Newlyn).

## Usage

```
# Retrieve meta data
select.station(loc=NULL , param = NULL,  stid = NULL ,lon = NULL,
               lat = NULL, alt = NULL, cntr = NULL, src = NULL ,
               it = NULL , nmin = NULL , verbose=FALSE)

# Retrieve data
station(stid=NULL,...)

station.default(loc=NULL, param="t2m", src = c("GHCNM", "ECAD", "GHCND",
        "NACD","NARP","NORDKLIMA"), stid=NULL, lon=NULL, lat=NULL,
        alt=NULL, cntr=NULL, it= NULL,nmin=30,
        path=NULL, plot=FALSE,verbose=FALSE)

# Monthly weather stations
station.nordklim(...)
station.narp(...)
station.nacd(...)
station.ghcnm(...,path=~ /data.GHCNM")
station.metnom(...)

#Daily weather stations
station.ecad(..., path=~ /data.ECAD")
station.ghcnd(..., path=~ /data.GHCND")
station.metnod(...)
station.sonel(urls=c(http://www.sonel.org/msl/Demerliac/VALIDATED/dCHERB.slv,
                     http://www.sonel.org/msl/Demerliac/VALIDATED/dRSCOF.slv,
                     http://www.sonel.org/msl/Demerliac/VALIDATED/dLCONQ.slv,
                     http://www.sonel.org/msl/Demerliac/VALIDATED/dBREST.slv,
                     http://www.sonel.org/msl/Demerliac/VALIDATED/dHAVRE.slv,
                     http://www.sonel.org/msl/Demerliac/VALIDATED/dDIEPP.slv,
                     http://www.sonel.org/msl/Demerliac/VALIDATED/dBOULO.slv,
```



```

http://www.sonel.org/msl/Demerliac/VALIDATED/dCALAI.slv,
http://www.sonel.org/msl/Demerliac/VALIDATED/dDUNKE.slv,
http://www.sonel.org/msl/Demerliac/VALIDATED/dCONCA.slv,
http://www.sonel.org/msl/Demerliac/VALIDATED/dPTUDY.slv,
http://www.sonel.org/msl/Demerliac/VALIDATED/dSNAZA.slv,
http://www.sonel.org/msl/Demerliac/VALIDATED/dBOUCA.slv,
http://www.sonel.org/msl/Demerliac/VALIDATED/dSJLUZ.slv,
http://www.sonel.org/msl/Demerliac/VALIDATED/dPBLOC.slv,
http://www.sonel.org/msl/Demerliac/VALIDATED/dLROCH.slv),
verbose=FALSE)
station.gloss(url=http://browse.ceda.ac.uk/browse/badc/CDs/gloss/data)
station.newlyn(path=data/gloss-241_Newlyn,verbose=TRUE)
allgood(x,miss=.1,verbose=TRUE)
clean.station(x,miss=.1,verbose=TRUE)

```

### Arguments

loc	A string of characters as the name of the location (weather/climate station) or an object of class "stationmeta".
param	Parameter or element type or variable identifier. There are several core parameters or elements as well as a number of additional parameters. The parameters or elements are: precip = Precipitation (mm) tas, tavg = 2m-surface temperature (in degrees Celcius) tmax, tasmax = Maximum temperature (in degrees Celcius) tmin, tasmin = Minimum temperature (in degrees Celcius)
src	Source: limit the downscaling to a specific data set ("NARP", "NACD", "NORD-KLIMA", "GHCM", "METNOM", "ECAD", "GHCND" and "METNOD")
stid	A string of characters as an identifier of the weather/climate station.
lon	Numeric value of longitude (in decimal degrees East) for the reference point (e.g. weather station) as a single value or a vector containing the range of longitude values in the form of c(lon.min,lon.max)
lat	Numeric value of latitude for the reference point (in decimal degrees North) or a vector containing the range of latitude values in the form of c(lat.min,lat.max)
alt	Numeric value of altitude (in meters a.s.l.) used for selection. Positive value, select all stations above this altitude; for negative values, select all stations below this altitude.
cntr	A string or a vector of strings of the full name of the country: Select the stations from a specified country or a set of countries.
it	A single integer or a vector of integers or Dates. An integer in the range of [1:12] for months, an integer of 4 digits for years (e.g. 2014), or a vector of Dates in the form "2014-01-01").
nmin	Select only stations with at least nmin number of years, months or days depending on the class of object x (e.g. 30 years).
plot	Logical value. If, TRUE provides a plot.
verbose	Logical value defaulting to FALSE. If FALSE, do not display comments (silent mode). If TRUE, displays extra information on progress.
path	The path where the data are stored. Can be a symbolic link.

**Value**

A time series of "zoo" "station" class with additional attributes used for further processing.

**Author(s)**

A. Mezghani

**See Also**

[station.meta](#) and [map.station](#).

**Examples**

```
# \dontrun{
# Get daily and monthly mean temperature for "Oslo" station ("18700") from METNO data source
t2m.dly <- station.metnod(stid="18700",param="t2m")
t2m.mon <- station.metnom(stid="18700",param="t2m")

# Get daily data from the ECA&D data source:
# If called for the first time, the script will download a huge chunk of
# data and store it locally.
# select meta for "De Bilt" station into ss,
ss <- select.station(loc = "de bilt",param="t2m",src="ECAD")
# Retrieve the data from the local directory specified in path based on
# previous selected station
t2m.dly <- station.ecad(loc=ss,path=~ /data.ECAD")
# or directly retrieve the data without a prior selection
t2m.dly <- station.ecad(loc = "oslo - blindern",param="t2m",path=~ /data.ECAD")
plot(t2m.dly)
# Aggregate to monthly and annual mean temperature values and plot the results
t2m.mon <- as.monthly(t2m.dly, FUN="mean") ; plot(t2m.mon)
t2m.ann <- as.annual(t2m.mon, FUN = "mean") ; plot(t2m.ann)
# specify one station from ECAD, and this time get daily mean precipitation
precip.dly <- station.ecad(loc="Oxford",param="precip") ; plot(precip.dly)
# Aggregate to annual accumulated precipitation values and plot the result
precip.ann <- as.annual(precip.dly,FUN="sum") ; plot(precip.ann)

# Get daily data from the GHCND data source
# Select a subset of stations across Norway with a minimum number of
# 130 years using "GHCND" as a data source, retrieve the data and show its
# structure.
ss <- select.station(cntr="NORWAY",param="precip",src="GHCND",nmin=130)
y <- station.ghcnd(loc=ss , path=~ /data.GHCND",plot=TRUE)
str(y)
# Subselect one station and display the geographical location of both selected stations and highlight the subselected
y1 <- subset(y,is=2)
map(y, xlim = c(-10,30), ylim = c(50,70), cex=1, select=y1, cex.select=2, showall=TRUE)
#}
```

station.meta

*Weather station metadata***Description**

Meta datasets based on different data sources or datasets included in esd package. Mainly weather stations' coordinates and other meta data.

**Usage**

```
data(station.meta)
```

**Format**

The format is:

List of 12

```
$ station_id: chr [1:284239] "6447" "6193" "21100" "25140" ...
$ location : chr [1:284239] "UCCLE" "HAMMERODDE_FYR" "VESTERVIG" "NORDBY" ...
$ country : chr [1:284239] "BELGIA" "DENMARK" "DENMARK" "DENMARK" ...
$ longitude : num [1:284239] 4.35 14.78 8.32 8.4 10.6 ...
$ latitude : num [1:284239] 50.8 55.3 56.8 55.4 55.9 ...
$ altitude : num [1:284239] 100 11 18 5 11 9 4 51 85 105 ...
$ element : chr [1:284239] "101" "101" "101" "101" ...
$ start : chr [1:284239] "1833" "1853" "1873" "1872" ...
$ end : chr [1:284239] NA NA NA NA ...
$ source : chr [1:284239] "NACD" "NACD" "NACD" "NACD" ...
$ wmo : num [1:284239] 6447 6193 -999 -999 -999 ...
$ quality : int [1:284239] 2 2 2 2 2 1 1 2 5 5 ...
- attr(*, "history")= chr [1:7]
"meta2esd.R - data taken from the clim.pact package and consolidated for NACD and NARP"
"nordklim.meta.rda" "ecad.meta.rda" "ghcnd.meta.rda" ...
- attr(*, "date")=function ()
- attr(*, "call")= language meta2esd(save = TRUE)
- attr(*, "author")= chr "R.E. Benestad & A. Mezghani"
- attr(*, "URLs")= chr [1:6] "www.dmi.dk/dmi/sr96-1.pdf"
"http://www.norden.org/en/publications/publikationer/2005-450"
"http://www.smhi.se/hfa_coord/nordklim/" "http://eca.knmi.nl/" ...
```

**See Also**

[select.station,station](#)

**Examples**

```
data(station.meta)
str(station.meta)
map(station.meta)
```

subset

*Subsetting esd objects***Description**

The subset method tries to be 'intelligent', and if the list has no names, then the list contains two vectors of length 2, then this is interpreted as a region, e.g. argument `is = list( c(lon.min, lon.max), c(lat.min, lat.max) )`. If, on the other hand, `is = list( lon=1:50, lat=55:65 )`, then the function picks the longitudes and latitudes which match these. This makes it flexible so that one can pick any irregular sequence.

**Usage**

```
subset(x, it=NULL, is=NULL, ...)
subset.station(x, it = NULL, is = NULL, verbose=FALSE)
subset.eof(x, it=NULL, is=NULL, verbose=FALSE)
subset.cca(x, it=NULL, is=NULL)
subset.mvr(x, it=NULL, is=NULL)
subset.pca(x, pattern=NULL, it=NULL, is=NULL, verbose=FALSE)
subset.trend(x, it=NULL, is=NULL)
subset.corfield(x, it=NULL, is=NULL)
subset.comb(x, it=NULL, is=NULL)
subset.events(x, it=NULL, is=NULL)
subset.trajectory(x, it=NULL, is=NULL)
subset.field(x, it=NULL, is=NULL)
subset.spell(x, is=NULL, it=NULL)
subset.ds(x, it=NULL, is=NULL)
subset.trend(x, it=NULL, is=NULL)
subset.dsensemble(x, it=NULL, is=NULL)
subset.zoo(x, it=NULL)
subset.trajectory(x, it=NULL, is=NULL)
matchdate(x, it)
sort.station(x, is=NULL, decreasing=TRUE)
```

**Arguments**

<code>x</code>	Data object from which the subset is taken
<code>it</code>	A list or data.frame providing time index, e.g. month
<code>is</code>	A list or data.frame providing space index, e.g. station record
<code>pattern</code>	selection of patterns in PCA or EOF (used for e.g. filtering the data)
<code>verbose</code>	Dump diagnostics to the screen

**Value**

A field object

**Author(s)**

R.E. Benestad and A. Mezghanil

**Examples**

```

data(Oslo)
# January months:
jan <- subset(Oslo,1)
# The last 10 years:
recent <- subset(Oslo,2003:2012)
# JJA season
jja <- subset(Oslo,it="jja")
# Seasonl values for MAM
mam <- subset(as.4seasons(Oslo),it=2)

data(ferder)
# Aggregated values for May (it=5)
may <- subset(as.monthly(Oslo),it=5)
# The last 10 aggregated annual values
recent.ann <- subset(as.annual(Oslo),it=2004:2013)

gcm <- t2m.NorESM.M()
# Extract July months from a field:
gcm.jul <- subset(gcm,it=7)

# Extract a period from a field:
gcm.short <- subset(gcm.jul,it=1950:2030)

# Extract data for the region 0-50E/55-65N
X <- subset(gcm,is=list(c(0,50),c(55,65)))

# Extract data for a specific set of longitudes and latitudes
Z <- subset(gcm,is=list(lon=1:30,lat=58:63))

t2m <- t2m.NCEP(lon=c(-10,30),lat=c(50,70))
cal <- subset(t2m,it=c("1948-01-01","1980-12-31"))

# Example on how to split the data into two parts for
# split-sample test...

T2M <- as.annual(t2m.NCEP(lon=c(-10,30),lat=c(50,70)))
cal <- subset(T2M,it=c(1948,1980))
pre <- subset(T2M,it=c(1981,2012))
comb <- combine(cal,pre)
X <- EOF(comb)
plot(X)

data(ferder)
y <- as.annual(ferder)
z <- DS(y,X)
plot(z)

```

```

# Test of subset the commutative property of subset and combine:
T2M <- as.4seasons(t2m.NCEP(lon=c(-10,30),lat=c(50,70)))
GCM <- as.4seasons(t2m.NorESM.M(lon = range(lon(T2M))+c(-2,2), lat = range(lat(T2M))+c(-2,2)))
XY <- combine(T2M,GCM)
X1 <- subset(XY,it=3)
X2 <- combine(subset(T2M,it=3),subset(GCM,it=3))
eof1 <- EOF(X1)
eof2 <- EOF(X2)
eof3 <- biasfix(eof2)
plot(merge(eof1[,1],eof2[,1],eof3[,1]),plot.type=single,
      col=c(red,blue,green),lty=c(1,1,2),lwd=c(4,2,2))
# OK - identical results

# Extract storm tracks for specific periods and regions
# from the example trajectory object imilast.M03
data(imilast.M03)
# spring season (march, april, may)
x1 <- subset(imilast.M03,it=mam)
plot(x1)

# region 10W-10E/55-65N
x2 <- subset(imilast.M03,is=list(lat=c(55,65),lon=c(-10,10)))
map(x2)

# storms longer than 3 days (12x6 hours)
x3 <- subset(imilast.M03,it=(param.trajectory(imilast.M03,param=n)>12))
map(x3)

```

---

summary.dsensemble	<i>Summary showing summary of objects</i>
--------------------	---

---

## Usage

```

summary.dsensemble(x, years = seq(1990, 2090, by = 20))
summary.station(x)
summary.ds(x)
summary.eof(x)
summary.cca(x)

```

## Arguments

x  
years

## Examples

```

##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,

```

```

##--or do  help(data=index)  for the standard data sets.

## The function is currently defined as
function (x, years = seq(1990, 2090, by = 20))
{
  x0 <- subset(x, it = 0)
  djf <- subset(x, it = "djf")
  mam <- subset(x, it = "mam")
  jja <- subset(x, it = "jja")
  son <- subset(x, it = "son")
  tab <- rep("", length(years) + 1)
  tab[1] <- paste(loc(x), "  Annual, DFJ, MAM, JJA, SON")
  i <- 1
  for (yr in years) {
    i <- i + 1
    tab[i] <- paste(years[i - 1], ":", " ", round(mean(coredata(subset(x0,
      it = years[i - 1]))), 2), " [", round(quantile(subset(x0,
      it = years[i - 1]), 0.05), 2), " ", round(quantile(subset(x0,
      it = years[i - 1]), 0.95), 2), "]", " ", round(mean(coredata(subset(djf,
      it = years[i - 1]))), 2), " [", round(quantile(subset(djf,
      it = years[i - 1]), 0.05), 2), " ", round(quantile(subset(djf,
      it = years[i - 1]), 0.95), 2), "]", " ", round(mean(coredata(subset(mam,
      it = years[i - 1]))), 2), " [", round(quantile(subset(mam,
      it = years[i - 1]), 0.05), 2), " ", round(quantile(subset(mam,
      it = years[i - 1]), 0.95), 2), "]", " ", round(mean(coredata(subset(jja,
      it = years[i - 1]))), 2), " [", round(quantile(subset(jja,
      it = years[i - 1]), 0.05), 2), " ", round(quantile(subset(jja,
      it = years[i - 1]), 0.95), 2), "]", " ", round(mean(coredata(subset(son,
      it = years[i - 1]))), 2), " [", round(quantile(subset(son,
      it = years[i - 1]), 0.05), 2), " ", round(quantile(subset(son,
      it = years[i - 1]), 0.95), 2), "]", sep = "")
  }
  tab
}

```

---

test.retrieve

*Test functions*


---

## Description

This routine contains a series of test functions which compute the global mean 2m-temperature anomalies and other predefined regions based on both CMIP3 and CMIP5 experiments. The main function is [test.retrieve.ncdf4](#). The others are specific cases for the "Scandinavian" and "Arctic" regions for both CMIP3 and CMIP5 projects

## Usage

```

test.retrieve(path=~"/CMIP3.monthly/20c3m-sresa1b",param="auto",
  cntr= "Global",lon=NULL,lat=NULL,time=NULL,lev=NULL,
  experiment="CMIP3",climatology=c(1986,2005),fig=TRUE,

```

```

saveinfile=TRUE,verbose=FALSE)

test.cmip3.arctic(...)
test.cmip5.arctic(...)
test.cmip3.global(...)
test.cmip5.global(...)
test.cmip3.scandinavia(...)
test.cmip5.scandinavia

test.cmip35.global()

```

### Arguments

path	Character vector of the full path to the CMIP data files.
param	Parameter or element type. There are several core parameters or elements as well as a number of additional parameters. The parameters or elements are # (Abdelkader : We need to update this list): auto = automatic selection. prep, pr = Precipitation (mm) tas, tavg = 2m-surface temperature (in degrees Celcius) tmax, tasmax = Maximum temperature (in degrees Celcius) tmin, tasmin = Minimum temperature (in degrees Celcius)
lon	Numeric value of longitude for the reference point (in decimal degrees East) or a vector containing the range of longitude values in the form of c(lon.min,lon.max)
lat	Numeric value of latitude for the reference point (in decimal degrees North) or a vector containing the range of latitude values in the form of c(lat.min,lat.max)
lev	Numeric value of pressure levels or a vector containing the range of pressure level values in the form of c(lev.min,lev.max)
time	Numerical year values or date values of time or a vector containing the range of values in the form of c(time.min,time.max). Date format should be in the form of "YYYY-MM-DD".
saveinfile	Logical or a character value. The output filename is set automatically if not specified. Default value is "TRUE".
verbose	Logical value. If TRUE, do not display comments (silent mode)

### Value

A field object

### Author(s)

A. Mezghani

### Examples

```

# Eg.1 :
# Compute the global mean surface temperature anomalies from CMIP3 experiment
test.cmip3.global(...)
# Compute the global mean surface temperature anomalies from CMIP5 experiment
test.cmip5.global(...)

```



```
# Plot the Global t2m-temperature anomalies from both CMIP3 and CMIP5 experiments.
plot.cmip35.global()

# Eg.2 Compute the mean surface temperature anomalies from CMIP3 experiment over Arctic
test.cmip3.arctic(...)

# Eg.3 Compute the mean surface temperature anomalies from CMIP5 experiment over Scandinavia.
test.cmip5.scandinavia(...)
```

track

*3-step cyclone tracking algorithm.*

## Description

Applies a tracking algorithm to a set of cyclones ([CCI](#)).

The algorithm connects events in three subsequent time steps, choosing the path that minimizes the change in angle and displacement between them. The analysis can be applied to 'events' objects.

The maximum displacement ('dmax') between two time steps depends on the angle of direction. The circle of maximum displacement around a cyclone can be shifted in the east-west ('dE') and north-south ('dN') direction. For example, 'dE' = 0.3 means that the maximum displacement is 1.3\*dmax in the eastward direction and only 0.7\*dmax in the westward direction, in line with the general westerly flow in the storm track regions.

Note: The algorithm has been developed for tracking midlatitude cyclones in the northern hemisphere and may not work as well for other regions or 'events' of different types, e.g., anti-cyclones.

## Usage

```
track(X, x0=NULL, it=NULL, is=NULL, dmax=1.2E6, amax=90,
      nmax=124, nmin=5, dmin=5E5, dE=0.3, dN=0.2,
      lplot=FALSE, progress=TRUE, verbose=FALSE)
track.events(X, ...)
```

## Arguments

X	An 'events' object containing temporal and spatial information about a set of cyclones or anticyclones.
x0	A tracked 'events' object from previous time steps, used as a starting point for the tracking of X so that trajectories can continue from x0 to X.
it	A list providing time index, e.g. month.
is	A list providing space index, lon and/or lat.
dmax	Maximum displacement of events between two time steps in a trajectory. Unit: m.
amax	Maximum change in the angle of direction. Unit: degrees.

nmax	Maximum lifetime of a trajectory. Unit: number of time steps.
nmin	Minimum lifetime of a trajectory. Unit: number of time steps.
dmin	Minimum length of a trajectory. Unit: m.
dE	Shift of the maximum displacement in the eastward direction.
dN	Shift of the maximum displacement in the northward direction.
lplot	TRUE: Produce plots of trajectories for selected time steps.
progress	TRUE: Show progress bar.
verbose	TRUE: Print out diagnostics.

### Value

An 'events' object containing the original information as well as the trajectory number ('trajectory') of each event and statistical properties of the trajectories ('trackcount' - number of events in path; 'tracklen' - distance between start and end point of path').

### Author(s)

K. Parding

---

transforms

*Various formulas, equations and transforms.*

---

### Description

Computes different formulas.

C.C.eq: Clapeyron-Clausius equation (saturation evaporation pressure) where x is a data object holding the temperature.

precip.vul: and index for the vulnerability to precipitation defined as  $\text{wetmean}(x)/\text{wetfreq}(x)$ . High when the mean intensity is high and/or the frequency is low (it rains seldom, but when it rains, it really pours down).

t2m.vul: and index for the vulnerability to temperature defined as the mean spell length for heat waves with temperatures exceeding 30C (default).

precip.rv: a rough estimate of the return value for precipitation under the assumption that it is exponentially distributed. Gives approximate answers for low return levels (less than 20 years). Advantage, can be predicted given wet-day mean and frequency.

nv: number of valid data points.

precip.Pr: rough estimate of the probability of more than x0 of rain based on an exponential distribution.

t2m.Pr: rough estimate of the probability of more than x0 of rain based on a normal distribution.

NE: predicts the number of events given the probability Pr.

**Usage**

```

C.C.eq(x)
precip.vul(x)
t2m.vul(x,x0=30,is=1)
precip.rv(x,tau=10)
nv(x)
precip.Pr(x,x0=10)
t2m.Pr(x,x0=10,na.rm=TRUE)
NE(p)

```

**Arguments**

x	a data object
p	a probability
x0	a threshold value
tau	time scale (years)
is	which of the spell results [1,2]
na.rm	See <a href="#">mean</a> .

**Value**

The right hand side of the equation

**Author(s)**

R. Benestad, MET Norway

**Examples**

```

t2m <- t2m.ERAINT(lon=c(-70,-10),lat=c(20,60))
es <- C.C.eq(t2m)
map(es)

```

---

trend

*Trending and detrending data*


---

**Description**

Trend analysis and de-trending of data.ls

**Usage**

```

trend(x,result="trend",model="y ~ t",...)
trend.default(x,result="trend",model="y ~ t",...)
trend.one.station(x,result="trend",model="y ~ t",...)
trend.station(x,result="trend",model="y ~ t",...)
trend.eof(x,result="trend",model="y ~ t",...)
trend.field(x,result="trend",model="y ~ t",...)
trend.zoo(x,result="trend",model="y ~ t",...)

```

**Arguments**

x	The data object
result	"trend" returns the trend; "residual" returns the residual
model	The trend model used by <a href="#">lm</a> .

**Value**

Similar type object as the input object

**Author(s)**

R.E. Benestad

**See Also**

[link{climatology}](#), [link{anomaly}](#)

**Examples**

```

data(ferder)
tr <- trend(annual(ferder,max))
attr(tr,coefficients)

```

---

vis.trends

---

*Visualise trends for multiple overlapping periods*


---

**Description**

Produce a plot showing trends for multiple periods within a time series. The strength of the trend is represented by the color scale and significant trends are marked with black borders.

**Usage**

```

vis.trends(x, unitlabel = "unit", varlabel = "", pmax = 0.01, minlen =
15, lwd = NA, vmax = NA, new = TRUE, show.significance = TRUE, verbose =
FALSE)

```

**Arguments**

<code>x</code>	the 'x' argument provides the time series for which the trend analysis is performed. Only zoo objects are accepted.
<code>minlen</code>	minimum time interval to calculate trends for in units of years.
<code>unitlabel</code>	unit of x.
<code>varlabel</code>	name of x.
<code>vmax</code>	upper limit of trend scale.
<code>show.significance</code>	TRUE to mark statistically significant trends.
<code>pmax</code>	maximum p-value of trends marked as significant.
<code>verbose</code>	TRUE or FALSE.

**Author(s)**

Kajsa Parding

**Examples**

```
t <- seq(as.Date("1955-01-01"), as.Date("2004-12-31"), by=1)
x <- zoo(sample(seq(-30, 30, 1e-1), length(t), rep=TRUE), order.by=t)
vis.trends(x, show.significance=FALSE)

data(Oslo)
vis.trends(Oslo, unitlabel="oC", varlabel = "Temperature",
  pmax = 1e-2, minlen = 40)
vis.trends(subset(Oslo, it=jja), unitlabel="oC",
  varlabel = "Temperature JJA",
  pmax = 1e-3, vmax=0.5, minlen = 40)
vis.trends(subset(Oslo, it=mam), unitlabel="oC",
  varlabel = "Temperature MAM",
  pmax = 1e-3, vmax=0.5, minlen = 40)
```

**Description**

Weather generators for conditional simulation of daily temperature and/or precipitation, given mean and/or standard deviation. The family of WG functions produce stochastic time series with similar characteristics as the station series provided (if none is provided, it will use either ferder or bjornholt provided by the esd-package). Here characteristics means similar mean value, standard deviation, and spectral properties. FTscramble takes the Fourier components (doing a Fourier Transform - FT) of a series and reassigns random phase to each frequency and then returns a new series through an inverse FT. The FT scrambling is used for temperature, but not for precipitation that is non-Gaussian and involves sporadic events with rain. For precipitation, a different approach is used,

taking the wet-day frequency of each year and using the wet-day mean and randomly generated exponentially distributed numbers to provide similar aggregated annual statistics as the station or predicted through downscaling. The precipitation WG can also take into account the number of consecutive number-of-dry-days statistics, using either a Poisson or a geometric distribution.

The weather generator produces a series with similar length as the provided sample data, but with shifted dates according to specified scenarios for annual mean mean/standard deviation/wet-day mean/wet-day frequency.

WG.FT.day.t2m generates daily temperature from seasonal means and standard deviations. It is given a sample station series, and uses FTscramble to generate a series with random phase but similar (or predicted - in the future) spectral characteristics. It then uses a quantile transform to prescribe predicted mean and standard deviation, assuming the distributions are normal. The temporal structure (power spectrum) is therefore similar as the sample provided.

WG.fw.day.precip uses the annual wet-day mean and the wet-day frequency as input, and takes a sample station of daily values to stochastically simulate number consecutive wet days based on its annual mean number. If not specified, it is taken from the sample data after being phase scrambled (FTscramble). The number of wet-days per year is estimated from the wet-day frequency, it too taken to be phase scrambled estimates from the sample data unless specifically specified. The daily amount is taken from stochastic values generated with [rexp](#). The number of consecutive wet days can be approximated by a geometric distribution ([rgeom](#)), and the annual mean number was estimated from the sample series.

## Usage

```
FTscramble(x, interval=NULL, spell.stats=FALSE, wetfreq.pred=FALSE)
WG(x, ...)
WG.station(x, option=default, ...)
WG.FT.day.t2m(x=NULL, amean=NULL, asd=NULL, t=NULL, eofs=1:4,
              select=NULL, lon=c(-20, 20), lat=c(-20, 20),
              plot=TRUE, biascorrect=TRUE, verbose=FALSE)
WG.fw.day.precip(x=NULL, mu=NULL, fw=NULL, ndd=NULL, t=NULL,
                 threshold=1, select=NULL,
                 lon=c(-10, 10), lat=c(-10, 10),
                 plot=TRUE, biascorrect=TRUE,
                 method=rgeom)
WG.pca.day.t2m.precip(t2m=NULL, precip=NULL, threshold=1, select=NULL)
```

## Arguments

x	station object
option	Define the type of WG
amean	annual mean values. If NULL, use those estimated from x; if NA, estimate using <a href="#">DSensemble.t2m</a> , or if provided, assume a 'dsensemble' object.
asd	annual standard deviation. If NULL, use those estimated from x; if NA, estimate using <a href="#">DSensemble.t2m</a> , or if provided, assume a 'dsensemble' object.
t	Time axis. If null, use the same as x or the last interval of same length as x from downscaled results.
eofs	passed on to <a href="#">DSensemble.t2m</a>

select	passed on to <a href="#">DSensemble.t2m</a>
lon	passed on to <a href="#">DSensemble.t2m</a>
lat	passed on to <a href="#">DSensemble.t2m</a>
plot	
biascorrect	passed on to <a href="#">DSensemble.t2m</a>
verbose	passed on to <a href="#">DSensemble.t2m</a>
mu	annual wet-mean values. If NULL, use those estimated from x; if NA, estimate using <a href="#">DSensemble.t2m</a> , or if provided, assume a 'dsensemble' object.
fw	annual wet-day frequency. If NULL, use those estimated from x; if NA, estimate using <a href="#">DSensemble.t2m</a> , or if provided, assume a 'dsensemble' object.
ndd	annual mean dry spell length. If NULL, use those estimated from x; if NA, estimate using <a href="#">DSensemble.t2m</a> , or if provided, assume a 'dsensemble' object.
threshold	Definition of a rainy day.
method	Assume a gemoetric or a poisson distribution. Can also define ownth methods.
t2m	station object with temperature
precip	station object with precipitation.

**Author(s)**

R.E. Benestad

**Examples**

```
data(ferder)
t2m <- WG(ferder)
data(bjornholt)
pr <- WG(bjornholt)
```

---

write2ncdf	<i>Saves climate data as netCDF.</i>
------------	--------------------------------------

---

**Description**

Method to save station data as netCDF, making sure to include the data structure and meta-data (attributes). The code tries to follow the netCDF 'CF' convention. The method is built on the [ncdf4](#) package.

**Usage**

```
write2ncdf(x,file,...)
write2ncdf.station(x,file,prec=short,missval=-99.9,offs=0,scalf=0.1,
                  torq=1899-12-31,verbose=FALSE)
```

**Arguments**

x	data object
file	file name
prec	Precision: see <a href="#">ncvar_def</a>
missval	Missing value: see <a href="#">ncvar_def</a>
offs	Sets the attribute 'add_offset' which is added to the values stored (to save space may be represented as 'short').
scalf	Sets the attribute 'scale_factor' which is used to scale (multiply) the values stored (to save space may be represented as 'short').
torg	Time origin
verbose	TRUE - clutter the screen.

**Value**

A "zoo" "field" object with additional attributes used for further processing.

**Author(s)**

R.E. Benestad

**See Also**

[test.retrieve.ncdf4](#).

**Examples**

```
nacd <- station(src=nacd)
X <- annual(nacd)
write2ncdf(X, file=test.nc)
```



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