

# Package ‘extremis’

October 28, 2018

**Version** 0.90

**Date** 2018-10-21

**Title** Statistics of Extremes

**Description** Conducts inference in statistical models for extreme values (de Carvalho et al (2012), <doi:10.1080/03610926.2012.709905>; de Carvalho and Davison (2014), <10.1080/01621459.2013.872651>; Einmahl et al (2016), <doi:10.1111/rssb.12099>).

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**Depends** R (>= 3.0.1)

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**License** GPL (>= 3)

**Repository** CRAN

**Imports** emplik

**NeedsCompilation** yes

**Date/Publication** 2018-10-28 22:30:03 UTC

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angcdf

*Empirical-Likelihood Based Inference for the Angular Measure*


---

### Description

This function computes empirical-likelihood based estimators for the angular distribution function of a bivariate extreme value distribution.

### Usage

```
angcdf(Y, tau = 0.95, method = "euclidean", raw = TRUE)
```

### Arguments

Y	data frame with two columns from which the estimate is to be computed.
tau	value used to threshold the data; by default it is set as the 0.95 quantile of the pseudo-radius $\tau = 0.95$ .
method	a character string setting the method to be used. By default <code>method = "euclidean"</code> , the other option being <code>method = "empirical"</code> . See details.
raw	logical; if TRUE, Y will be converted to unit Fréchet scale. If FALSE, Y will be understood as already in unit Fréchet scale.

### Details

`method = "euclidean"` implements the maximum Euclidean likelihood spectral distribution function as introduced by de Carvalho et al (2013). `method = "empirical"` implements the maximum Empirical likelihood spectral distribution function as introduced by Einmahl and Segers (2009).

### Value

H	angular distribution function.
w	pseudo-angles.
Y	data.

The `plot` method depicts the empirical likelihood-based angular distribution function.

### Author(s)

Miguel de Carvalho

### References

de Carvalho, M., Oumow, B., Segers, J. and Warchol, M. (2013) A Euclidean likelihood estimator for bivariate tail dependence. *Communications in Statistics—Theory and Methods*, 42, 1176–1192.

Einmahl, J. H. J., and Segers, J. (2009) Maximum empirical likelihood estimation of the spectral measure of an extreme-value distribution. *The Annals of Statistics*, 37, 2953–2989.

**Examples**

```
## de Carvalho et al (2013, Fig. 7)
data(beatenberg)
attach(beatenberg)
fit <- angcdf(beatenberg, tau = 0.98, raw = FALSE)
plot(fit)
rug(fit$w)
```

angdensity

*Empirical-Likelihood Based Inference for the Angular Density***Description**

This function computes empirical-likelihood based estimators for the angular distribution function of a bivariate extreme value distribution.

**Usage**

```
angdensity(Y, tau = 0.95, nu, grid = seq(0.01, 0.99, length = 2^8),
  method = "euclidean", raw = TRUE)
```

**Arguments**

<code>Y</code>	data frame with two columns from which the estimate is to be computed.
<code>tau</code>	value used to threshold the data; by default it is set as the 0.95 quantile of the pseudo-radius.
<code>nu</code>	concentration parameter of beta distribution which controls the amount of smoothing.
<code>grid</code>	grid with coordinates of the points where the angular density is estimated; by default <code>grid = seq(0.01, 0.99, length = 2^8)</code> .
<code>method</code>	a character string setting the method to be used. By default <code>method = "euclidean"</code> , the other option being <code>method = "empirical"</code> . See details.
<code>raw</code>	logical; if TRUE, <code>Y</code> will be converted to unit Fréchet scale. If FALSE, <code>Y</code> will be understood as already in unit Fréchet scale.

**Details**

The smooth angular density was introduced in by de Carvalho et al (2013). `method = "euclidean"` implements the version of the method based on Euclidean likelihood weights, whereas `method = "empirical"` uses Empirical likelihood weights.

**Value**

h                    the estimate angular density values.  
grid                grid with coordinates of the points where the angular density is estimated.  
w                    pseudo-angles.  
nu                   concentration parameter of the Beta-kernel.  
Y                    raw data.

The plot method depicts the smooth angular density.

**Author(s)**

Miguel de Carvalho

**References**

de Carvalho, M., Oumow, B., Segers, J. and Warchol, M. (2013) A Euclidean likelihood estimator for bivariate tail dependence. *Communications in Statistics—Theory and Methods*, 42, 1176–1192.

**Examples**

```
## de Carvalho et al (2013, Fig. 7)
data(beatenberg)
attach(beatenberg)
fit <- angdensity(beatenberg, tau = 0.98, nu = 163, raw = FALSE)
plot(fit)
rug(fit$w)
```

---

angscdf

*Smooth Empirical-Likelihood Based Inference for the Angular Measure*

---

**Description**

This function computes smooth empirical-likelihood based estimators for the angular distribution function of a bivariate extreme value distribution.

**Usage**

```
angscdf(Y, tau = 0.95, nu, grid = seq(0.01, 0.99, length = 2^8),
method = "euclidean", raw = TRUE)
```

**Arguments**

Y	data frame with two columns from which the estimate is to be computed.
tau	value used to threshold the data; by default it is set as the 0.95 quantile of the pseudo-radius $\tau = 0.95$ .
nu	concentration parameter of beta distribution which controls the amount of smoothing.
grid	grid with coordinates of the points where the angular measure is estimated; by default <code>grid = seq(0.01, 0.99, length = 2^8)</code> .
method	a character string setting the method to be used. By default <code>method = "euclidean"</code> , the other option being <code>method = "empirical"</code> . See details.
raw	logical; if TRUE, Y will be converted to unit Fréchet scale. If FALSE, Y will be understood as already in unit Fréchet scale.

**Details**

`method = "euclidean"` implements the maximum Euclidean likelihood spectral distribution function as introduced by de Carvalho et al (2013). `method = "empirical"` implements the maximum Empirical likelihood spectral distribution function as introduced by Einmahl and Segers (2009).

**Value**

H	the estimated angular distribution function values.
grid	grid with coordinates of the points where the angular measure is estimated.
w	pseudo-angles.
nu	concentration parameter of the Beta-kernel.
Y	raw data.

The `plot` method depicts the empirical likelihood-based angular distribution function.

**Author(s)**

Miguel de Carvalho

**References**

- de Carvalho, M., Oumow, B., Segers, J. and Warchol, M. (2013) A Euclidean likelihood estimator for bivariate tail dependence. *Communications in Statistics—Theory and Methods*, 42, 1176–1192.
- Einmahl, J. H. J., and Segers, J. (2009) Maximum empirical likelihood estimation of the spectral measure of an extreme-value distribution. *The Annals of Statistics*, 37, 2953–2989.

**Examples**

```
## de Carvalho et al (2013, Fig. 7)
data(beatenberg)
attach(beatenberg)
fit <- angscdf(beatenberg, tau = 0.98, nu = 163, raw = FALSE)
plot(fit)
rug(fit$w)
```

---

 beatenberg

*Beatenberg*


---

### Description

Preprocessed pairs of temperatures in unit Fréchet scale from Beatenberg forest, registered under forest cover and in the open field. Preprocessing is conducted as described in Ferrez et al (2011).

### Usage

beatenberg

### Format

The beatenberg data frame has 2839 rows and 2 columns.

### References

Ferrez, J., A. C. Davison, and Rebetez., M. (2011) Extreme temperature analysis under forest cover compared to an open field. *Agricultural and Forest Meteorology*, 151, 992–1001.

---

 cdensity

*Kernel Smoothed Scedasis Density*


---

### Description

This function computes a kernel scedasis density estimate.

### Usage

```
cdensity(Y, threshold = quantile(Y[, 2], 0.95), ...)
```

### Arguments

Y	data frame from which the estimate is to be computed; first column corresponds to time and the second to the variable of interest.
threshold	value used to threshold the data y; by default threshold = quantile(y, 0.95).
...	further arguments for density methods.

### Details

Kernel smoothing for the scedasis density was introduced by Einmahl et al (2016).

**Value**

c                   scedasis density estimator.  
k                   number of exceedances above the threshold.  
w                   standardized indices of exceedances.  
Y                   raw data.

The plot method depicts the smooth scedasis density.

**Author(s)**

Miguel de Carvalho

**References**

Einmahl, J. H., Haan, L., and Zhou, C. (2016) Statistics of heteroscedastic extremes. *Journal of the Royal Statistical Society: Ser. B*, 78(1), 31–51.

**Examples**

```
data(lse)
attach(lse)
Y <- data.frame(DATE[-1], -diff(log(ROYAL.DUTCH.SHELL.B)))
T <- dim(Y)[1]
k <- floor((0.4258597) * T / (log(T)))
fit <- cdensity(Y, kernel = "biweight", bw = 0.1 / sqrt(7),
               threshold = sort(Y[, 2])[T - k])
plot(fit)
plot(fit, original = FALSE)
```

---

cdf

*Empirical Scedasis Distribution Function*


---

**Description**

This function computes the empirical scedasis distribution function.

**Usage**

```
cdf(Y, threshold = quantile(Y[, 2], 0.95))
```

**Arguments**

Y                   data frame from which the estimate is to be computed; first column corresponds to time and the second to the variable of interest.  
threshold          value used to threshold the data y; by default threshold = quantile(Y[, 2], 0.95).

**Details**

The empirical scedasis distribution function was introduced by Einmahl et al (2016).

**Value**

C                    empirical scedasis distribution function.  
w                    standardized indices of exceedances.  
k                    number of exceedances above a threshold.  
Y                    raw data.

The plot method depicts the empirical cumulative scedasis function, and the reference line for the case of constant frequency of extremes over time (if `uniform = TRUE`).

**Author(s)**

Miguel de Carvalho

**References**

Einmahl, J. H., Haan, L., and Zhou, C. (2016) Statistics of heteroscedastic extremes. *Journal of the Royal Statistical Society: Ser. B*, 78(1), 31–51.

**Examples**

```
data(sp500)
attach(sp500)
Y <- data.frame(date[-1], -diff(log(close)))
fit <- cdf(Y)
plot(fit)
plot(fit, original = FALSE)
```

---

cmodes

*Mode Mass Function*

---

**Description**

This function computes the mode mass function.

**Usage**

```
cmodes(Y, thresholds = apply(Y[, -1], 2, quantile, probs =
  0.95), nu = 100, ...)
```



**Arguments**

Y	data frame from which the estimate is to be computed; first column corresponds to time and the second to the variable of interest.
thresholds	values used to threshold the data y; by default threshold = quantile(y, 0.95).
nu	concentration parameter of beta kernel used to smooth mode mass function.
...	further arguments for density methods.

**Details**

The scedasis functions on which the mode mass function is based are computed using the default "nrd0" option for bandwidth.

**Value**

c	scedasis density estimators.
k	number of exceedances above the threshold.
w	standardized indices of exceedances.
Y	raw data.

The plot method depicts the smooth mode mass function along with the smooth scedasis densities.

**Author(s)**

Miguel de Carvalho

**References**

Rubio, R., de Carvalho, M., and Huser, R. (2018) Similarity-Based Clustering of Extreme Losses from the London Stock Exchange. Submitted.

**Examples**

```
data(lse)
attach(lse)
nlr <- -apply(log(lse[, -1]), 2, diff)
Y <- data.frame(DATE[-1], nlr)
T <- dim(Y)[1]
k <- floor((0.4258597) * T / (log(T)))
fit <- cmodes(Y, thresholds = as.numeric(apply(nlr, 2, sort)[T - k, ]),
             kernel = "biweight", bw = 0.1 / sqrt(7), nu = 100)
plot(fit)
```

kgvar

*K-Geometric Means Algorithm for Value-at-Risk***Description**

This function performs k-geometric means for time-varying value-at-risk.

**Usage**

```
kgvar(y, centers, iter.max = 10, conf.level = 0.95)
```

**Arguments**

<code>y</code>	data frame from which the estimate is to be computed; first column corresponds to time and the second to the remainder of interest.
<code>centers</code>	the number of clusters or a set of initial (distinct) cluster centres. If a number, a random set of (distinct) rows in <code>y</code> is chosen as the initial centers.
<code>iter.max</code>	the maximum number of iterations allowed. The default is 10.
<code>conf.level</code>	the confidence level. The default is 0.95.

**Details**

The intermediate sequence  $\kappa_T$  is chosen proportional to  $T/\log T$ .

**Value**

`kgvar` returns an object of class "kgvar" which has a fitted method. It is a list with at least the following components:

<code>var.new</code>	cluster center value-at-risk function.
<code>clusters</code>	cluster allocation.
<code>Y</code>	raw data.
<code>n.clust</code>	number of clusters.
<code>scale.param</code>	the scale parameters in the Pareto-like tail specification.
<code>conf.level</code>	the confidence level.
<code>hill</code>	hill estimator of extreme value index.

The `plot` method depicts the k-geometric means algorithm for time-varying value-at-risk. If `c.c` is TRUE, the method displays the cluster means.

**Author(s)**

Miguel de Carvalho, Rodrigo Rubio.

## References

Rubio, R., de Carvalho, M. and Huser, R. (2018) Similarity-Based Clustering of Extreme Losses from the London Stock Exchange. Submitted.

## Examples

```
## Not run:
## Example (Overlapping version of Fig. 8 in Supplementary Materials)
data(lse)
attach(lse)
y <- -apply(log(lse[, -1]), 2, diff)
fit <- kgvar(y, centers = 3)
plot(fit, c.c = TRUE, ylim = c(0, 0.1))

## End(Not run)
```

---

khetmeans

*K-Means Clustering for Heteroscedastic Extremes*


---

## Description

This function performs k-means clustering for heteroscedastic extremes.

## Usage

```
khetmeans(y, centers, iter.max = 10, alpha = 0.5)
```

## Arguments

y	data frame from which the estimate is to be computed; first column corresponds to time and the second to the remainder of interest.
centers	the number of clusters or a set of initial (distinct) cluster centres. If a number, a random set of (distinct) rows in y is chosen as the initial centers.
iter.max	the maximum number of iterations allowed. The default is 10.
alpha	the tuning parameter. The default is 0.5.

## Details

The intermediate sequence  $\kappa_T$  is chosen proportional to  $T/\log T$ .

## Value

khetmeans returns an object of class "khetmeans" which has a fitted method. It is a list with at least the following components:

mus.new	cluster center scedasis density.
mugamma.new	cluster center extreme value index.

clusters            cluster allocation.  
 Y                    raw data.  
 n.clust            number of clusters.

The plot method depicts the k-means clustering for heteroscedastic extremes. If c.c is TRUE, the method displays the cluster means.

### Author(s)

Miguel de Carvalho, Rodrigo Rubio.

### References

Rubio, R., de Carvalho, M. and Huser, R. (2018) Similarity-Based Clustering of Extreme Losses from the London Stock Exchange. Submitted.

### Examples

```
## Not run:
## Example 1 (Scenario B, T = 5000)
## This example requires package evd
require(evd)
set.seed(12)
T <- 5000
n <- 30
b <- 0.1
gamma1 <- 0.7
gamma2 <- 1
grid <- seq(0, 1, length = 100)
c2 <- function(s)
  dbeta(s, 2, 5)
c3 <- function(s)
  dbeta(s, 5, 2)
X <- matrix(0, ncol = T, nrow = n)
for(i in 1:5)
  for(j in 1:T)
    X[i, j] <- rgev(1, c2(j / T), c2(j / T), gamma1)
for(i in 6:15)
  for(j in 1:T)
    X[i, j] <- rgev(1, c2(j / T), c2(j / T), gamma2)
for(i in 16:20)
  for(j in 1:T)
    X[i, j] <- rgev(1, c3(j / T), c3(j / T), gamma1)
for(i in 21:30)
  for(j in 1:T)
    X[i, j] <- rgev(1, c3(j / T), c3(j / T), gamma2)
Y <- t(X)
fit <- khetmeans(Y, centers = 4)
plot(fit, c.c = TRUE)
lines(grid, c2(grid), type = 'l', lwd = 8, col = 'black')
lines(grid, c3(grid), type = 'l', lwd = 8, col = 'black')
```

```
## End(Not run)

## Not run:
## Example 2 (Overlapping version of Fig. 5 in Supplementary Materials)
data(lse)
attach(lse)
y <- -apply(log(lse[, -1]), 2, diff)
fit <- khetmeans(y, centers = 3)
plot(fit, c.c = TRUE, ylim = c(0, 3))

## End(Not run)
```

---

lse

*Selected Stocks from the London Stock Exchange*

---

### Description

Prices at close from 26 selected stocks from the London stock exchange from 1989 till 2016.

### Usage

```
lse
```

### Format

The lse data frame has 6894 rows and 27 columns.

### References

Rubio, R., de Carvalho, M., and Huser (2018) Similarity-based clustering of extreme losses from the London stock exchange.

---

plotFrechet

*Unit Fréchet Scatterplot in Log-log Scale*

---

### Description

This function depicts a scatterplot of bivariate data transformed to unit Fréchet scale.

### Usage

```
plotFrechet(Y, tau = 0.95, raw = TRUE, ...)
```

**Arguments**

Y	list with data from which the estimates are to be computed.
tau	value used to threshold the data y; by default <code>treshold = quantile(y, 0.95)</code> .
raw	logical; if TRUE, Y will be converted to unit Fréchet scale. If FALSE, Y will be understood as already in unit Fréchet scale.
...	other arguments to be passed to <code>plot</code> .

**Details**

The solid line corresponds to the boundary threshold in the log-log scale, with both axes being logarithmic.

**Author(s)**

Miguel de Carvalho

**Examples**

```
## de Carvalho et al (2013, Fig. 5)
data(beatenberg)
plotFrechet(beatenberg, xlab = "Forest Cover", ylab = "Open Field",
            raw = FALSE)
```

---

sp500

*Standard & Poor 500*

---

**Description**

Daily Standard and Poor's index at close from 1988 till 2007.

**Usage**

`sp500`

**Format**

The `sp500` data frame has 5043 rows and 2 columns.

**References**

- de Carvalho, M. (2016) Statistics of extremes: Challenges and opportunities. In: *Handbook of EVT and its Applications to Finance and Insurance*. Eds F. Longin. Hoboken: Wiley.
- Einmahl, J. H., Haan, L., and Zhou, C. (2016) Statistics of heteroscedastic extremes. *Journal of the Royal Statistical Society: Ser. B*, 78(1), 31–51.

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