

# Package ‘extremis’

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

**Author** Miguel de Carvalho [aut, cre],  
Rodrigo Rubio [aut],  
Vianey Palacios [aut]

**Depends** R (>= 3.0.1)

**Maintainer** Miguel de Carvalho <[miguel.decarvalho@ed.ac.uk](mailto:miguel.decarvalho@ed.ac.uk)>

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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**

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.
nu	concentration parameter of beta distribution which controls the amount of smoothing.
grid	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> .
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**

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)
```

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beatenberg	<i>Beatenberg</i>
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---

**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.

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cdensity	<i>Kernel Smoothed Scedasis Density</i>
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---

**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 <code>threshold = quantile(y, 0.95)</code> .
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.

---

plotFréchet

*Unit Fréchet Scatterplot in Log-log Scale*

---

### Description

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

### Usage

```
plotFréchet(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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