

# Package ‘qfa’

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**Type** Package

**Title** Quantile-Frequency Analysis (QFA) of Time Series

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## Description

Quantile-frequency analysis (QFA) of time series based on trigonometric quantile regression.

References:

- [1] Li, T.-H. (2012) ``Quantile periodograms'', Journal of the American Statistical Association, 107, 765–776, <doi:10.1080/01621459.2012.682815>.
- [2] Li, T.-H. (2014) Time Series with Mixed Spectra, CRC Press, <doi:10.1201/b15154>
- [3] Li, T.-H. (2022) ``Quantile Fourier transform, quantile series, and nonparametric estimation of quantile spectra'', <doi:10.48550/arXiv.2211.05844>.
- [4] Li, T.-H. (2024) ``Quantile crossing spectrum and spline autoregression estimation," <doi:10.48550/arXiv.2412.02513>.

**Depends** R (>= 3.5)

**Imports** RhpcBLASctl,

doParallel,  
fields,  
foreach,  
mgcv,  
nlme,  
parallel,  
quantreg,  
splines,  
stats,  
graphics,  
colorRamps,  
MASS

**License** GPL (>=2)

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|     |                          |
|-----|--------------------------|
| per | <i>Periodogram (PER)</i> |
|-----|--------------------------|

---

Description

This function computes the periodogram or periodogram matrix for univariate or multivariate time series.

Usage

per(y)

Arguments

y                      vector (n) or matrix (n x nc) of time series

Value

A vector (n) or array (nc x nc x n) of periodogram

## Examples

```
y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y.per <- per(y)
plot(y.per)
```

---

qacf

*Quantile Autocovariance Function (QACF)*


---

## Description

This function computes quantile autocovariance function (QACF) from time series or quantile discrete Fourier transform (QDFT).

## Usage

```
qacf(y, tau, y.qdft = NULL, n.cores = 1, cl = NULL)
```

## Arguments

|         |   |
|---------|---|
| y       | vector or matrix of time series (if matrix, nrow(y) = length of time series)  |
| tau     | sequence of quantile levels in (0,1)  |
| y.qdft  | matrix or array of pre-calculated QDFT (default = NULL: compute from y and tau); if y.qdft is supplied, y and tau can be left unspecified |
| n.cores | number of cores for parallel computing of QDFT if y.qdft = NULL (default = 1)   |
| cl      | pre-existing cluster for repeated parallel computing of QDFT (default = NULL)   |

## Value

matrix or array of quantile autocovariance function

## Examples

```
y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
# compute from time series
y.qacf <- qacf(y,tau)
# compute from QDFT
y.qdft <- qdft(y,tau)
y.qacf <- qacf(y.qdft=y.qdft)
```

---

|       |   |
|-------|---|
| qcser | <i>Quantile-Crossing Series (QCSER)</i> |
|-------|---|

---

**Description**

This function creates the quantile-crossing series (QCSER) for univariate or multivariate time series.

**Usage**

```
qcser(y, tau, normalize = FALSE)
```

**Arguments**

|           |  |
|-----------|--|
| y         | vector or matrix of time series                                |
| tau       | vector of quantile levels in (0,1)                             |
| normalize | TRUE or FALSE (default): normalize QCSER to have unit variance |

**Value**

A matrix or array of quantile-crossing series

**Examples**

```
y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y.qser <- qcser(y,tau)
dim(y.qser)
```

---

|      |   |
|------|---|
| qdft | <i>Quantile Discrete Fourier Transform (QDFT)</i> |
|------|---|

---

**Description**

This function computes quantile discrete Fourier transform (QDFT) for univariate or multivariate time series.

**Usage**

```
qdft(y, tau, n.cores = 1, cl = NULL)
```

**Arguments**

|         |  |
|---------|--|
| y       | vector or matrix of time series (if matrix, nrow(y) = length of time series) |
| tau     | sequence of quantile levels in (0,1)   |
| n.cores | number of cores for parallel computing (default = 1)                         |
| cl      | pre-existing cluster for repeated parallel computing (default = NULL)        |

**Value**

matrix or array of quantile discrete Fourier transform of y

## Examples

```
y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y.qdft <- qdft(y,tau)
# Make a cluster for repeated use
n.cores <- 2
cl <- parallel::makeCluster(n.cores)
parallel::clusterExport(cl, c("tqr.fit"))
doParallel::registerDoParallel(cl)
y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y.qdft <- qdft(y1,tau,n.cores=n.cores,cl=cl)
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y.qdft <- qdft(y2,tau,n.cores=n.cores,cl=cl)
parallel::stopCluster(cl)
```

qdft2qacf

*Quantile Autocovariance Function (QACF)*

## Description

This function computes quantile autocovariance function (QACF) from QDFT.

## Usage

```
qdft2qacf(y.qdft, return.qser = FALSE)
```

## Arguments

|             |  |
|-------------|--|
| y.qdft      | matrix or array of QDFT from qdft()                    |
| return.qser | if TRUE, return quantile series (QSER) along with QACF |

## Value

matrix or array of quantile autocovariance function if return.qser = FALSE (default), else a list with the following elements:

|      |   |
|------|---|
| qacf | matrix or array of quantile autocovariance function |
| qser | matrix or array of quantile series                  |

## Examples

```
# single time series
y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y.qdft <- qdft(y1,tau)
y.qacf <- qdft2qacf(y.qdft)
plot(c(0:9),y.qacf[c(1:10),1],type='h',xlab="LAG",ylab="QACF")
y.qser <- qdft2qacf(y.qdft,return.qser=TRUE)$qser
plot(y.qser[,1],type='l',xlab="TIME",ylab="QSER")
# multiple time series
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
y.qdft <- qdft(cbind(y1,y2),tau)
y.qacf <- qdft2qacf(y.qdft)
plot(c(0:9),y.qacf[1,2,c(1:10),1],type='h',xlab="LAG",ylab="QACF")
```

---

|           |                                    |
|-----------|------------------------------------|
| qdft2qper | <i>Quantile Periodogram (QPER)</i> |
|-----------|------------------------------------|

---

**Description**

This function computes quantile periodogram (QPER) from QDFT.

**Usage**

```
qdft2qper(y.qdft)
```

**Arguments**

`y.qdft`                      matrix or array of QDFT from `qdft()`

**Value**

matrix or array of quantile periodogram

**Examples**

```
# single time series
y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y.qdft <- qdft(y1,tau)
y.qper <- qdft2qper(y.qdft)
n <- length(y1)
ff <- c(0:(n-1))/n
sel.f <- which(ff > 0 & ff < 0.5)
qfa.plot(ff[sel.f],tau,Re(y.qper[sel.f,]))
# multiple time series
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
y.qdft <- qdft(cbind(y1,y2),tau)
y.qper <- qdft2qper(y.qdft)
qfa.plot(ff[sel.f],tau,Re(y.qper[1,1,sel.f,]))
qfa.plot(ff[sel.f],tau,Re(y.qper[1,2,sel.f,]))
```

---

|           |                               |
|-----------|-------------------------------|
| qdft2qser | <i>Quantile Series (QSER)</i> |
|-----------|-------------------------------|

---

**Description**

This function computes quantile series (QSER) from QDFT.

**Usage**

```
qdft2qser(y.qdft)
```

**Arguments**

`y.qdft`                      matrix or array of QDFT from `qdft()`

**Value**

matrix or array of quantile series

**Examples**

```
# single time series
y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y.qdft <- qdft(y1,tau)
y.qser <- qdft2qser(y.qdft)
plot(y.qser[,1],type='l',xlab="TIME",ylab="QSER")
# multiple time series
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
y.qdft <- qdft(cbind(y1,y2),tau)
y.qser <- qdft2qser(y.qdft)
plot(y.qser[1,,1],type='l',xlab="TIME",ylab="QSER")
```

qfa.plot

*Quantile-Frequency Plot***Description**

This function creates an image plot of quantile spectrum.

**Usage**

```
qfa.plot(
  freq,
  tau,
  rqper,
  rg.qper = range(rqper),
  rg.tau = range(tau),
  rg.freq = c(0, 0.5),
  color = colorRamps::matlab.like2(1024),
  ylab = "QUANTILE LEVEL",
  xlab = "FREQUENCY",
  tlab = NULL,
  set.par = TRUE,
  legend.plot = TRUE
)
```

**Arguments**

|         |  |
|---------|--|
| freq    | sequence of frequencies in (0,0.5) at which quantile spectrum is evaluated   |
| tau     | sequence of quantile levels in (0,1) at which quantile spectrum is evaluated |
| rqper   | real-valued matrix of quantile spectrum evaluated on the freq x tau grid     |
| rg.qper | zlim for qper (default = range(qper))  |
| rg.tau  | ylim for tau (default = range(tau))  |
| rg.freq | xlim for freq (default = c(0, 0.5))  |
| color   | colors (default = colorRamps::matlab.like2(1024))                            |

|             |   |
|-------------|---|
| ylab        | label of y-axis (default = "QUANTILE LEVEL")    |
| xlab        | label of x-axis (default = "FREQUENCY")         |
| tlab        | title of plot (default = NULL)                  |
| set.par     | if TRUE, par() is set internally (single image) |
| legend.plot | if TRUE, legend plot is added                   |

**Value**

no return value

---

|                |  |
|----------------|--|
| qkl.divergence | <i>Kullback-Leibler Divergence of Quantile Spectral Estimate</i> |
|----------------|--|

---

**Description**

This function computes Kullback-Leibler divergence (KLD) of quantile spectral estimate.

**Usage**

```
qkl.divergence(y.qper, qspec, sel.f = NULL, sel.tau = NULL)
```

**Arguments**

|         |   |
|---------|---|
| y.qper  | matrix or array of quantile spectral estimate from, e.g., qspec.lw()                    |
| qspec   | matrix of array of true quantile spectrum (same dimension as y.qper)                    |
| sel.f   | index of selected frequencies for computation (default = NULL: all frequencies)         |
| sel.tau | index of selected quantile levels for computation (default = NULL: all quantile levels) |

**Value**

real number of Kullback-Leibler divergence

---

|      |                                    |
|------|------------------------------------|
| qper | <i>Quantile Periodogram (QPER)</i> |
|------|------------------------------------|

---

**Description**

This function computes quantile periodogram (QPER) from time series or quantile discrete Fourier transform (QDFT).

**Usage**

```
qper(y, tau, y.qdft = NULL, n.cores = 1, cl = NULL)
```



**Arguments**

|         |   |
|---------|---|
| y       | vector or matrix of time series (if matrix, <code>nrow(y)</code> = length of time series)   |
| tau     | sequence of quantile levels in (0,1)  |
| y.qdft  | matrix or array of pre-calculated QDFT (default = NULL: compute from y and tau); if y.qdft is supplied, y and tau can be left unspecified |
| n.cores | number of cores for parallel computing of QDFT if y.qdft = NULL (default = 1)   |
| cl      | pre-existing cluster for repeated parallel computing of QDFT (default = NULL)   |

**Value**

matrix or array of quantile periodogram

**Examples**

```
y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
# compute from time series
y.qper <- qper(y,tau)
# compute from QDFT
y.qdft <- qdft(y,tau)
y.qper <- qper(y.qdft=y.qdft)
```

qper2

*Quantile Periodogram Type II (QPER2)***Description**

This function computes type-II quantile periodogram for univariate time series.

**Usage**

```
qper2(y, freq, tau, weights = NULL, n.cores = 1, cl = NULL)
```

**Arguments**

|         |   |
|---------|---|
| y       | univariate time series  |
| freq    | sequence of frequencies in [0,1)  |
| tau     | sequence of quantile levels in (0,1)  |
| weights | sequence of weights in quantile regression (default = NULL: weights equal to 1) |
| n.cores | number of cores for parallel computing (default = 1)                            |
| cl      | pre-existing cluster for repeated parallel computing (default = NULL)           |

**Value**

matrix of quantile periodogram evaluated on `freq * tau` grid

**Examples**

```

y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
n <- length(y)
ff <- c(0:(n-1))/n
sel.f <- which(ff > 0 & ff < 0.5)
y.qper2 <- qper2(y,ff,tau)
qfa.plot(ff[sel.f],tau,Re(y.qper2[sel.f,]))

```

qser

*Quantile Series (QSER)***Description**

This function computes quantile series (QSER) from time series or quantile discrete Fourier transform (QDFT).

**Usage**

```
qser(y, tau, y.qdft = NULL, n.cores = 1, cl = NULL)
```

**Arguments**

|         |   |
|---------|---|
| y       | vector or matrix of time series (if matrix, nrow(y) = length of time series)  |
| tau     | sequence of quantile levels in (0,1)  |
| y.qdft  | matrix or array of pre-calculated QDFT (default = NULL: compute from y and tau); if y.qdft is supplied, y and tau can be left unspecified |
| n.cores | number of cores for parallel computing of QDFT if y.qdft = NULL (default = 1)   |
| cl      | pre-existing cluster for repeated parallel computing of QDFT (default = NULL)   |

**Value**

matrix or array of quantile series

**Examples**

```

y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
# compute from time series
y.qser <- qser(y,tau)
# compute from QDFT
y.qdft <- qdft(y,tau)
y.qser <- qser(y.qdft=y.qdft)

```

---

|         |   |
|---------|---|
| qser2ar | <i>Autoregression (AR) Model of Quantile Series</i> |
|---------|---|

---

**Description**

This function fits an autoregression (AR) model to quantile series (QSER) separately for each quantile level using `stats::ar()`.

**Usage**

```
qser2ar(y.qser, p = NULL, order.max = NULL, method = c("none", "gamm", "sp"))
```

**Arguments**

|                        |  |
|------------------------|--|
| <code>y.qser</code>    | matrix or array of pre-calculated QSER, e.g., using <code>qser()</code>                                  |
| <code>p</code>         | order of AR model (default = NULL: selected by AIC)  |
| <code>order.max</code> | maximum order for AIC if <code>p = NULL</code> (default = NULL: determined by <code>stats::ar()</code> ) |
| <code>method</code>    | quantile smoothing method: "gamm", "sp", or "NA" (default)   |

**Value**

a list with the following elements:

|                        |   |
|------------------------|---|
| <code>A</code>         | matrix or array of AR coefficients      |
| <code>V</code>         | vector or matrix of residual covariance |
| <code>p</code>         | order of AR model                       |
| <code>n</code>         | length of time series                   |
| <code>residuals</code> | matrix or array of residuals            |

---

|           |  |
|-----------|--|
| qser2qacf | <i>ACF of Quantile Series (QSER) or Quantile-Crossing Series (QCACF)</i> |
|-----------|--|

---

**Description**

This function creates the ACF of quantile series or quantile-crossing series

**Usage**

```
qser2qacf(y.qser)
```

**Arguments**

|                     |   |
|---------------------|---|
| <code>y.qser</code> | matrix or array of quantile-crossing series |
|---------------------|---|

**Value**

A matrix or array of ACF

## Examples

```
y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y.qser <- qcser(y,tau)
y.qacf <- qser2qacf(y.qser)
dim(y.qacf)
```

---

qser2sar

*Spline Autoregression (SAR) Model of Quantile Series*


---

## Description

This function fits spline autoregression (SAR) model to quantile series (QSER).

## Usage

```
qser2sar(
  y.qser,
  tau,
  d = 1,
  p = NULL,
  order.max = NULL,
  spar = NULL,
  method = c("GCV", "AIC", "BIC"),
  weighted = FALSE
)
```

## Arguments

|           |   |
|-----------|---|
| y.qser    | matrix or array of pre-calculated QSER, e.g., using <code>qser()</code>                     |
| tau       | sequence of quantile levels where y.qser is calculated                                      |
| d         | subsampling rate of quantile levels (default = 1)   |
| p         | order of SAR model (default = NULL: automatically selected by AIC)                          |
| order.max | maximum order for AIC if p = NULL (default = NULL: determined by <code>stats::ar()</code> ) |
| spar      | penalty parameter alla <code>smooth.spline</code> (default = NULL: automatically selected)  |
| method    | criterion for penalty parameter selection: "AIC" (default), "BIC", or "GCV"                 |
| weighted  | if TRUE, penalty function is weighted (default = FALSE)                                     |

## Value

a list with the following elements:

|      |   |
|------|---|
| A    | matrix or array of SAR coefficients         |
| V    | vector or matrix of SAR residual covariance |
| p    | order of SAR model                          |
| spar | penalty parameter                           |
| tau  | sequence of quantile levels                 |
| n    | length of time series                       |

|          |                                      |
|----------|--------------------------------------|
| d        | subsampling rate of quantile levels  |
| weighted | option for weighted penalty function |
| fit      | object containing details of SAR fit |

qspec.ar

*Autoregression (AR) Estimator of Quantile Spectrum*

## Description

This function computes autoregression (AR) estimate of quantile spectrum from time series or quantile series (QSER).

## Usage

```
qspec.ar(
  y,
  tau,
  y.qser = NULL,
  p = NULL,
  order.max = NULL,
  freq = NULL,
  method = c("none", "gamm", "sp"),
  n.cores = 1,
  cl = NULL
)
```

## Arguments

|           |   |
|-----------|---|
| y         | vector or matrix of time series (if matrix, nrow(y) = length of time series)  |
| tau       | sequence of quantile levels in (0,1)  |
| y.qser    | matrix or array of pre-calculated QSER (default = NULL: compute from y and tau);  |
| p         | order of AR model (default = NULL: automatically selected by AIC)   |
| order.max | maximum order for AIC if p = NULL (default = NULL: determined by stats::ar())   |
| freq      | sequence of frequencies in [0,1) (default = NULL: all Fourier frequencies)  |
| method    | quantile smoothing method: "gamm" for mgcv::gamm(), "sp" for stats::smooth.spline(), or "none" (default) if y.qser is supplied, y and tau can be left unspecified |
| n.cores   | number of cores for parallel computing of QDFT if y.qser = NULL (default = 1)   |
| cl        | pre-existing cluster for repeated parallel computing of QDFT (default = NULL)   |

## Value

a list with the following elements:

|      |   |
|------|---|
| spec | matrix or array of AR quantile spectrum             |
| freq | sequence of frequencies                             |
| fit  | object of AR model                                  |
| qser | matrix or array of quantile series if y.qser = NULL |

## Examples

```

y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
y <- cbind(y1,y2)
tau <- seq(0.1,0.9,0.05)
n <- length(y1)
ff <- c(0:(n-1))/n
sel.f <- which(ff > 0 & ff < 0.5)
y.qspec.ar <- qspec.ar(y,tau,p=1)$spec
qfa.plot(ff[sel.f],tau,Re(y.qspec.ar[1,1,sel.f,]))
y.qser <- qcser(y1,tau)
y.qspec.ar <- qspec.ar(y.qser=y.qser,p=1)$spec
qfa.plot(ff[sel.f],tau,Re(y.qspec.ar[sel.f,]))
y.qspec.arqs <- qspec.ar(y.qser=y.qser,p=1,method="sp")$spec
qfa.plot(ff[sel.f],tau,Re(y.qspec.arqs[sel.f,]))

```

---

qspec.lw

*Lag-Window (LW) Estimator of Quantile Spectrum*


---

## Description

This function computes lag-window (LW) estimate of quantile spectrum with or without quantile smoothing from time series or quantile autocovariance function (QACF).

## Usage

```

qspec.lw(
  y,
  tau,
  y.qacf = NULL,
  M = NULL,
  method = c("none", "gamm", "sp"),
  spar = "GCV",
  n.cores = 1,
  cl = NULL
)

```

## Arguments

|         |   |
|---------|---|
| y       | vector or matrix of time series (if matrix, nrow(y) = length of time series)  |
| tau     | sequence of quantile levels in (0,1)  |
| y.qacf  | matrix or array of pre-calculated QACF (default = NULL: compute from y and tau); if y.qacf is supplied, y and tau can be left unspecified |
| M       | bandwidth parameter of lag window (default = NULL: quantile periodogram)  |
| method  | quantile smoothing method: "gamm" for mgcv::gamm(), "sp" for stats::smooth.spline(), or "none" (default)                                  |
| spar    | smoothing parameter in smooth.spline() if method = "sp" (default = "GCV")   |
| n.cores | number of cores for parallel computing (default = 1)  |
| cl      | pre-existing cluster for repeated parallel computing (default = NULL)   |

**Value**

A list with the following elements:

|         |  |
|---------|--|
| spec    | matrix or array of spectral estimate                                 |
| spec.lw | matrix or array of spectral estimate without quantile smoothing      |
| lw      | lag-window sequence  |
| qacf    | matrix or array of quantile autocovariance function if y.qacf = NULL |

**Examples**

```

y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
tau <- seq(0.1,0.9,0.05)
n <- length(y1)
ff <- c(0:(n-1))/n
sel.f <- which(ff > 0 & ff < 0.5)
y.qacf <- qacf(cbind(y1,y2),tau)
y.qper.lw <- qspec.lw(y.qacf=y.qacf,M=5)$spec
qfa.plot(ff[sel.f],tau,Re(y.qper.lw[1,1,sel.f,]))
y.qper.lwqs <- qspec.lw(y.qacf=y.qacf,M=5,method="sp",spar=0.9)$spec
qfa.plot(ff[sel.f],tau,Re(y.qper.lwqs[1,1,sel.f,]))

```

---

qspec.sar

*Spline Autoregression (SAR) Estimator of Quantile Spectrum*


---

**Description**

This function computes spline autoregression (SAR) estimate of quantile spectrum.

**Usage**

```

qspec.sar(
  y,
  y.qser = NULL,
  tau,
  d = 1,
  p = NULL,
  order.max = NULL,
  spar = NULL,
  method = c("GCV", "AIC", "BIC"),
  weighted = FALSE,
  freq = NULL,
  n.cores = 1,
  cl = NULL
)

```

**Arguments**

|                        |  |
|------------------------|--|
| <code>y</code>         | vector or matrix of time series (if matrix, <code>nrow(y)</code> = length of time series)  |
| <code>y.qser</code>    | matrix or array of pre-calculated QSER (default = NULL: compute from <code>y</code> and <code>tau</code> ); if <code>y.qser</code> is supplied, <code>y</code> can be left unspecified |
| <code>tau</code>       | sequence of quantile levels in (0,1)   |
| <code>d</code>         | subsampling rate of quantile levels (default = 1)  |
| <code>p</code>         | order of SAR model (default = NULL: automatically selected by AIC)   |
| <code>order.max</code> | maximum order for AIC if <code>p</code> = NULL (default = NULL: determined by <code>stats::ar()</code> )   |
| <code>spar</code>      | penalty parameter alla <code>smooth.spline</code> (default = NULL: automatically selected)   |
| <code>method</code>    | criterion for penalty parameter selection: "GCV", "AIC" (default), or "BIC"  |
| <code>weighted</code>  | if TRUE, penalty function is weighted (default = FALSE)  |
| <code>freq</code>      | sequence of frequencies in [0,1) (default = NULL: all Fourier frequencies)   |
| <code>n.cores</code>   | number of cores for parallel computing of QDFT if <code>y.qser</code> = NULL (default = 1)   |
| <code>cl</code>        | pre-existing cluster for repeated parallel computing of QDFT (default = NULL)  |

**Value**

a list with the following elements:

|                   |  |
|-------------------|--|
| <code>spec</code> | matrix or array of SAR quantile spectrum                         |
| <code>freq</code> | sequence of frequencies  |
| <code>fit</code>  | object of SAR model  |
| <code>qser</code> | matrix or array of quantile series if <code>y.qser</code> = NULL |

**Examples**

```

y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
tau <- seq(0.1,0.9,0.05)
n <- length(y1)
ff <- c(0:(n-1))/n
sel.f <- which(ff > 0 & ff < 0.5)
# compute from time series
y.sar <- qspec.sar(cbind(y1,y2),tau=tau,p=1)
qfa.plot(ff[sel.f],tau,Re(y.sar$spec[1,1,sel.f,]))
# compute from quantile series
y.qser <- qser(cbind(y1,y2),tau)
y.sar <- qspec.sar(y.qser=y.qser,tau=tau,p=1)
qfa.plot(ff[sel.f],tau,Re(y.sar$spec[1,1,sel.f,]))

```



---

|            |                                    |
|------------|------------------------------------|
| qspec2qcoh | <i>Quantile Coherence Spectrum</i> |
|------------|------------------------------------|

---

### Description

This function computes quantile coherence spectrum (QCOH) from quantile spectrum of multiple time series.

### Usage

```
qspec2qcoh(qspec, k = 1, kk = 2)
```

### Arguments

|       |                                      |
|-------|--------------------------------------|
| qspec | array of quantile spectrum           |
| k     | index of first series (default = 1)  |
| kk    | index of second series (default = 2) |

### Value

matrix of quantile coherence evaluated at Fourier frequencies in (0,0.5)

### Examples

```
y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
tau <- seq(0.1,0.9,0.05)
n <- length(y1)
ff <- c(0:(n-1))/n
sel.f <- which(ff > 0 & ff < 0.5)
y.qacf <- qacf(cbind(y1,y2),tau)
y.qper.lw <- qspec.lw(y.qacf=y.qacf,M=5)$spec
y.qcoh <- qspec2qcoh(y.qper.lw,k=1,kk=2)
qfa.plot(ff[sel.f],tau,y.qcoh)
```

---

|                  |  |
|------------------|--|
| sar.eq.bootstrap | <i>Bootstrap Simulation of SAR Coefficients for Testing Equality of Granger-Causality in Two Samples</i> |
|------------------|--|

---

### Description

This function simulates bootstrap samples of selected spline autoregression (SAR) coefficients for testing equality of Granger-causality in two samples based on their SAR models under H0: effect in each sample equals the average effect.

**Usage**

```
sar.eq.bootstrap(
  y.qser,
  fit,
  fit2,
  index = c(1, 2),
  nsim = 1000,
  method = c("ar", "sar"),
  n.cores = 1,
  mthreads = FALSE,
  seed = 1234567
)
```

**Arguments**

|          |   |
|----------|---|
| y.qser   | matrix or array of QSER from qser() or qspec.sar()\$qser  |
| fit      | object of SAR model from qser2sar() or qspec.sar()\$fit   |
| fit2     | object of SAR model for the other sample  |
| index    | a pair of component indices for multiple time series or a sequence of lags for single time series (default = c(1, 2)) |
| nsim     | number of bootstrap samples (default = 1000)  |
| method   | method of residual calculation: "ar" (default) or "sar"   |
| n.cores  | number of cores for parallel computing (default = 1)  |
| mthreads | if TRUE, multithread BLAS is enabled when available (default = FALSE, required for parallel computing)                |
| seed     | seed for random sampling (default = 1234567)  |

**Value**

array of simulated bootstrap samples of selected SAR coefficients

**Examples**

```
y11 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y21 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
y12 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y22 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
tau <- seq(0.1, 0.9, 0.05)
y1.sar <- qspec.sar(cbind(y11, y21), tau=tau, p=1)
y2.sar <- qspec.sar(cbind(y12, y22), tau=tau, p=1)
A1.sim <- sar.eq.bootstrap(y1.sar$qser, y1.sar$fit, y2.sar$fit, index=c(1, 2), nsim=5)
A2.sim <- sar.eq.bootstrap(y2.sar$qser, y2.sar$fit, y1.sar$fit, index=c(1, 2), nsim=5)
```

sar.eq.test

---

*Wald Test and Confidence Band for Equality of Granger-Causality in Two Samples*


---

## Description

This function computes Wald test and confidence band for equality of Granger-causality in two samples using bootstrap samples generated by `sar.eq.bootstrap()` based on the spline autoregression (SAR) models of quantile series (QSER).

## Usage

```
sar.eq.test(A1, A1.sim, A2, A2.sim, sel.lag = NULL, sel.tau = NULL)
```

## Arguments

|                      |   |
|----------------------|---|
| <code>A1</code>      | matrix of selected SAR coefficients for sample 1                              |
| <code>A1.sim</code>  | simulated bootstrap samples from <code>sar.eq.bootstrap()</code> for sample 1 |
| <code>A2</code>      | matrix of selected SAR coefficients for sample 2                              |
| <code>A2.sim</code>  | simulated bootstrap samples from <code>sar.eq.bootstrap()</code> for sample 2 |
| <code>sel.lag</code> | indices of time lags for Wald test (default = NULL: all lags)                 |
| <code>sel.tau</code> | indices of quantile levels for Wald test (default = NULL: all quantiles)      |

## Value

a list with the following elements:

|                   |   |
|-------------------|---|
| <code>test</code> | list of Wald test result containing wald and p.value        |
| <code>D.u</code>  | matrix of upper limits of 95% confidence band for $A1 - A2$ |
| <code>D.l</code>  | matrix of lower limits of 95% confidence band for $A1 - A2$ |

## Examples

```
y11 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y21 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
y12 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y22 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y1.sar <- qspec.sar(cbind(y11,y21),tau=tau,p=1)
y2.sar <- qspec.sar(cbind(y12,y22),tau=tau,p=1)
A1.sim <- sar.eq.bootstrap(y1.sar$qser,y1.sar$fit,y2.sar$fit,index=c(1,2),nsim=5)
A2.sim <- sar.eq.bootstrap(y2.sar$qser,y2.sar$fit,y1.sar$fit,index=c(1,2),nsim=5)
A1 <- sar.gc.coef(y1.sar$fit,index=c(1,2))
A2 <- sar.gc.coef(y2.sar$fit,index=c(1,2))
test <- sar.eq.test(A1,A1.sim,A2,A2.sim,sel.lag=NULL,sel.tau=NULL)
```

---

|                  |  |
|------------------|--|
| sar.gc.bootstrap | <i>Bootstrap Simulation of SAR Coefficients for Granger-Causality Analysis</i> |
|------------------|--|

---

## Description

This function simulates bootstrap samples of selected spline autoregression (SAR) coefficients for Granger-causality analysis based on the SAR model of quantile series (QSER) under  $H_0$ : (a) for multiple time series, the second series specified in `index` is not causal for the first series specified in `index`; (b) for single time series, the series is not causal at the lags specified in `index`.

## Usage

```
sar.gc.bootstrap(
  y.qser,
  fit,
  index = c(1, 2),
  nsim = 1000,
  method = c("ar", "sar"),
  n.cores = 1,
  mthreads = FALSE,
  seed = 1234567
)
```

## Arguments

|                       |   |
|-----------------------|---|
| <code>y.qser</code>   | matrix or array of QSER from <code>qser()</code> or <code>qspec.sar()</code> \$ <code>qser</code>                                   |
| <code>fit</code>      | object of SAR model from <code>qser2sar()</code> or <code>qspec.sar()</code> \$ <code>fit</code>                                    |
| <code>index</code>    | a pair of component indices for multiple time series or a sequence of lags for single time series (default = <code>c(1, 2)</code> ) |
| <code>nsim</code>     | number of bootstrap samples (default = 1000)  |
| <code>method</code>   | method of residual calculation: "ar" (default) or "sar"   |
| <code>n.cores</code>  | number of cores for parallel computing (default = 1)  |
| <code>mthreads</code> | if TRUE, multithread BLAS is enabled when available (default = FALSE, required for parallel computing)                              |
| <code>seed</code>     | seed for random sampling (default = 1234567)  |

## Value

array of simulated bootstrap samples of selected SAR coefficients

## Examples

```
y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
tau <- seq(0.1, 0.9, 0.05)
y.sar <- qspec.sar(cbind(y1, y2), tau=tau, p=1)
A.sim <- sar.gc.bootstrap(y.sar$qser, y.sar$fit, index=c(1, 2), nsim=5)
```

sar.gc.coef

*Extraction of SAR Coefficients for Granger-Causality Analysis***Description**

This function extracts the spline autoregression (SAR) coefficients from an SAR model for Granger-causality analysis. See `sar.gc.bootstrap` for more details regarding the use of `index`.

**Usage**

```
sar.gc.coef(fit, index = c(1, 2))
```

**Arguments**

|                    |   |
|--------------------|---|
| <code>fit</code>   | object of SAR model from <code>qser2sar()</code> or <code>qspec.sar()</code> \$fit  |
| <code>index</code> | a pair of component indices for multiple time series or a sequence of lags for single time series (default = <code>c(1, 2)</code> ) |

**Value**

matrix of selected SAR coefficients (number of lags by number of quantiles)

**Examples**

```
y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y.sar <- qspec.sar(cbind(y1,y2),tau=tau,p=1)
A <- sar.gc.coef(y.sar$fit,index=c(1,2))
```

sar.gc.test

*Wald Test and Confidence Band for Granger-Causality Analysis***Description**

This function computes Wald test and confidence band for Granger-causality using bootstrap samples generated by `sar.gc.bootstrap()` based the spline autoregression (SAR) model of quantile series (QSER).

**Usage**

```
sar.gc.test(A, A.sim, sel.lag = NULL, sel.tau = NULL)
```

**Arguments**

|                      |  |
|----------------------|--|
| <code>A</code>       | matrix of selected SAR coefficients  |
| <code>A.sim</code>   | simulated bootstrap samples from <code>sar.gc.bootstrap()</code>                       |
| <code>sel.lag</code> | indices of time lags for Wald test (default = <code>NULL</code> : all lags)            |
| <code>sel.tau</code> | indices of quantile levels for Wald test (default = <code>NULL</code> : all quantiles) |

**Value**

a list with the following elements:

|      |  |
|------|--|
| test | list of Wald test result containing wald and p.value |
| A.u  | matrix of upper limits of 95% confidence band of A   |
| A.l  | matrix of lower limits of 95% confidence band of A   |

**Examples**

```

y1 <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
y2 <- stats::arima.sim(list(order=c(1,0,0), ar=-0.5), n=64)
tau <- seq(0.1,0.9,0.05)
y.sar <- qspec.sar(cbind(y1,y2),tau=tau,p=1)
A <- sar.gc.coef(y.sar$fit,index=c(1,2))
A.sim <- sar.gc.bootstrap(y.sar$qser,y.sar$fit,index=c(1,2),nsim=5)
y.gc <- sar.gc.test(A,A.sim)

```

---

sqr.fit

*Spline Quantile Regression (SQR)*


---

**Description**

This function computes spline quantile regression (SQR) solution from response vector and design matrix. It uses the FORTRAN code `rqfmb.f` in the "quantreg" package with the kind permission of Dr. R. Koenker.

**Usage**

```
sqr.fit(y, X, tau, c0, d = 1, weighted = FALSE, mthreads = FALSE)
```

**Arguments**

|          |  |
|----------|--|
| y        | response vector  |
| X        | design matrix ( <code>nrow(X) = length(y)</code> )   |
| tau      | sequence of quantile levels in (0,1)   |
| c0       | penalty parameter  |
| d        | subsampling rate of quantile levels (default = 1)  |
| weighted | if TRUE, penalty function is weighted (default = FALSE)  |
| mthreads | if TRUE, multithread BLAS is enabled when available (default = FALSE, required for parallel computing) |

**Value**

A list with the following elements:

|              |                                   |
|--------------|-----------------------------------|
| coefficients | matrix of regression coefficients |
| nit          | number of iterations              |

tqr.fit

*Trigonometric Quantile Regression (TQR)***Description**

This function computes trigonometric quantile regression (TQR) for univariate time series at a single frequency.

**Usage**

```
tqr.fit(y, f0, tau, prepared = TRUE)
```

**Arguments**

|          |   |
|----------|---|
| y        | vector of time series   |
| f0       | frequency in [0,1)  |
| tau      | sequence of quantile levels in (0,1)  |
| prepared | if TRUE, intercept is removed and coef of cosine is doubled when $f0 = 0.5$ |

**Value**

object of `rq()` (coefficients in `$coef`)

**Examples**

```
y <- stats::arima.sim(list(order=c(1,0,0), ar=0.5), n=64)
tau <- seq(0.1,0.9,0.05)
fit <- tqr.fit(y,f0=0.1,tau=tau)
plot(tau,fit$coef[1,],type='o',pch=0.75,xlab='QUANTILE LEVEL',ylab='TQR COEF')
```

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