

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.
Spline quantile regression (SQR) for regression coefficient estimation.

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>.
- [5] Li, T.-H. (2024) ``Spline autoregression method for estimation of quantile spectrum", <doi:10.48550/arXiv.2412.17163>.

Depends R (>= 3.5)

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doParallel,
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Contents

birthweight	2
engel	3
per	3
qacf	4
qcsar	4
qdft	5
qdft2qacf	6
qdft2qper	7
qdft2qser	7
qfa.plot	8
qkl.divergence	9
qper	9
qper2	10
qser	11
qser2ar	12
qser2qacf	12
qser2sar	13
qspec.ar	14
qspec.lw	15
qspec.sar	16
qspec2qcoh	18
sar.eq.bootstrap	18
sar.eq.test	19
sar.gc.bootstrap	20
sar.gc.coef	21
sar.gc.test	22
sqdft	23
sqdft.fit	24
sqr	25
sqr.fit	26
sqr.fit.optim	27
tqr.fit	29
tsqr.fit	29
yearssn	30
Index	32

birthweight

Birthweight data

Description

Infant birth weight data. Precare and Education should be treated as factors.

Usage

`data(birthweight)`

Format

An object of class `data.frame` with 50000 rows and 12 columns.

Source

nativity2022us.csv, <<https://www.nber.org/research/data/vital-statistics-nativity-birth-data>>

References

Koenker, R. (2005). Quantile Regression. Cambridge University Press.

engel	<i>Engel food expenditure data</i>
-------	------------------------------------

Description

The Engel food expenditure data from the R package quantreg.

Usage

```
data(engel)
```

Format

An object of class `data.frame` with 235 rows and 2 columns.

References

Koenker, R. (2005). Quantile Regression. Cambridge University Press.

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 or matrix of time series `s` (if matrix, `nrow(y)` = length of time series)

Value

vector or array 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	<i>Quantile Autocovariance Function (QACF)</i>
------	--

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	sequence 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

*Quantile Discrete Fourier Transform (QDFT)***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")

```


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	<i>Autoregression (AR) Estimator of Quantile Spectrum</i>
----------	---

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 <code>y.qacf = NULL</code>

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 = TRUE,
  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 FALSE, set RhpcBLASctl::blas_set_num_threads(1) (default = TRUE)
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

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

Value

a list with the following elements:

test	list of Wald test result containing wald and p.value
D.u	matrix of upper limits of 95% confidence band for A1 – A2
D.l	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

Bootstrap Simulation of SAR Coefficients for Granger-Causality Analysis

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 H0: (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 = TRUE,
  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
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 FALSE, set RhpcBLASctl::blas_set_num_threads(1) (default = TRUE)
seed	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

fit	object of SAR model from <code>qser2sar()</code> or <code>qspec.sar()</code> \$fit
index	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

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

```

sqdft

Spline Quantile Discrete Fourier Transform (SQDFT) of Time Series

Description

This function computes spline quantile discrete Fourier transform (SQDFT) for univariate or multivariate time series through trigonometric spline quantile regression.

Usage

```

sqdft(
  y,
  tau,
  spar = NULL,
  d = 1,
  weighted = FALSE,
  method = c("AIC", "BIC", "SIC"),
  ztol = 1e-05,
  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)
spar	smoothing parameter: if <code>spar=NULL</code> , smoothing parameter is selected by method
d	subsampling rate of quantile levels (default = 1)
weighted	if TRUE, penalty function is weighted (default = FALSE)
method	criterion for smoothing parameter selection when <code>spar=NULL</code> ("AIC", "BIC", or "SIC")
ztol	zero tolerance parameter used to determine the effective dimensionality of the fit
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:

coefficients	matrix of regression coefficients
qdft	matrix or array of the spline quantile discrete Fourier transform of y
crit	criteria for smoothing parameter selection: (AIC,BIC,SIC)

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.sqdft <- sqdft(y,tau,spar=NULL,d=4,metho="AIC")$qdft
```

sqdft.fit	<i>Spline Quantile Discrete Fourier Transform (SQDFT) of Time Series Given Smoothing Parameter</i>
-----------	--

Description

This function computes spline quantile discrete Fourier transform (SQDFT) for univariate or multivariate time series through trigonometric spline quantile regression with user-supplied spar.

Usage

```
sqdft.fit(
  y,
  tau,
  spar = 1,
  d = 1,
  weighted = FALSE,
  ztol = 1e-05,
  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)
spar	smoothing parameter
d	subsampling rate of quantile levels (default = 1)
weighted	if TRUE, penalty function is weighted (default = FALSE)
ztol	zero tolerance parameter used to determine the effective dimensionality of the fit
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:

coefficients	matrix of regression coefficients
qdft	matrix or array of the spline quantile discrete Fourier transform of y
crit	criteria for smoothing parameter selection: (AIC,BIC,SIC)

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.sqdft <- sqdft.fit(y,tau,spar=1,d=4)$qdft
```

sqr

*Spline Quantile Regression (SQR) by formula***Description**

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

Usage

```
sqr(
  formula,
  tau = seq(0.1, 0.9, 0.2),
  spar = NULL,
  d = 1,
  data,
  subset,
  na.action,
  model = TRUE,
  weighted = FALSE,
  mthreads = TRUE,
  method = c("AIC", "BIC", "SIC"),
  ztol = 1e-05
)
```

Arguments

formula	a formula object, with the response on the left of a ~ operator, and the terms, separated by + operators, on the right.
tau	sequence of quantile levels in (0,1)
spar	smoothing parameter: if spar=NULL, smoothing parameter is selected by method
d	subsampling rate of quantile levels (default = 1)
data	a data.frame in which to interpret the variables named in the formula
subset	an optional vector specifying a subset of observations to be used
na.action	a function to filter missing data (see <code>rq</code> in the 'quantreg' package)

model	if TRUE then the model frame is returned (needed for calling summary subsequently)
weighted	if TRUE, penalty function is weighted (default = FALSE)
mthreads	if FALSE, set RhpcBLASctl::blas_set_num_threads(1) (default = TRUE)
method	a criterion for smoothing parameter selection if spar=NULL ("AIC", "BIC", or "SIC")
ztol	a zero tolerance parameter used to determine the effective dimensionality of the fit

Value

object of SQR fit

Examples

```
library(quantreg)
data(engel)
engel$income <- engel$income - mean(engel$income)
tau <- seq(0.1, 0.9, 0.05)
fit <- rq(foodexp ~ income, tau=tau, data=engel)
fit.sqr <- sqr(foodexp ~ income, tau=tau, spar=0.5, data=engel)
par(mfrow=c(1,1), pty="m", lab=c(10,10,2), mar=c(4,4,2,1)+0.1, las=1)
plot(tau, fit$coef[2,], xlab="Quantile Level", ylab="Coeff1")
lines(tau, fit.sqr$coef[2,])
```

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 `rqf.nb.f` in the "quantreg" package with the kind permission of Dr. R. Koenker.

Usage

```
sqr.fit(
  X,
  y,
  tau,
  spar = 1,
  d = 1,
  weighted = FALSE,
  mthreads = TRUE,
  ztol = 1e-05
)
```

Arguments

X	design matrix (nrow(X) = length(y))
y	response vector
tau	sequence of quantile levels in (0,1)
spar	smoothing parameter
d	subsampling rate of quantile levels (default = 1)
weighted	if TRUE, penalty function is weighted (default = FALSE)
mthreads	if FALSE, set RhpcBLASctl::blas_set_num_threads(1) (default = TRUE)
ztol	zero tolerance parameter used to determine the effective dimensionality of the fit

Value

A list with the following elements:

coefficients	matrix of regression coefficients
crit	sequence criteria for smoothing parameter select: (AIC,BIC,SIC)
np	sequence of number of effective parameters
fid	sequence of fidelity measure as quasi-likelihood
nit	number of iterations

sqr.fit.optim

Spline Quantile Regression (SQR) by Gradient Algorithms

Description

This function computes spline quantile regression by a gradient algorithm BFGS, ADAM, or GRAD.

Usage

```
sqr.fit.optim(
  X,
  y,
  tau,
  spar = 0,
  d = 1,
  weighted = FALSE,
  method = c("BFGS", "ADAM", "GRAD"),
  beta.rq = NULL,
  theta0 = NULL,
  spar0 = NULL,
  sg.rate = c(1, 1),
  mthreads = TRUE,
  control = list(trace = 0)
)
```

Arguments

<code>X</code>	vector or matrix of explanatory variables (including intercept)
<code>y</code>	vector of dependent variable
<code>tau</code>	sequence of quantile levels in (0,1)
<code>spar</code>	smoothing parameter
<code>d</code>	subsampling rate of quantile levels (default = 1)
<code>weighted</code>	if TRUE, penalty function is weighted (default = FALSE)
<code>method</code>	optimization method: "BFGS" (default), "ADAM", or "GRAD"
<code>beta.rq</code>	matrix of regression coefficients from <code>quantreg::rq(y~X)</code> for initialization (default = NULL)
<code>theta0</code>	initial value of spline coefficients (default = NULL)
<code>spar0</code>	smoothing parameter for <code>stats::smooth.spline()</code> to smooth <code>beta.rq</code> for initialization (default = NULL)
<code>sg.rate</code>	vector of sampling rates for quantiles and observations in stochastic gradient version of GRAD and ADAM
<code>mthreads</code>	if FALSE, set <code>RhpcBLASctl::blas_set_num_threads(1)</code> (default = TRUE)
<code>control</code>	a list of control parameters <code>maxit</code> : max number of iterations (default = 100) <code>stepsize</code> : stepsize for ADAM and GRAD (default = 0.01) <code>warmup</code> : length of warmup phase for ADAM and GRAD (default = 70) <code>stepupdate</code> : frequency of update for ADAM and GRAD (default = 20) <code>stepredn</code> : stepsize discount factor for ADAM and GRAD (default = 0.2) <code>line.search.type</code> : line search option (1,2,3,4) for GRAD (default = 1) <code>line.search.max</code> : max number of line search trials for GRAD (default = 1) <code>seed</code> : seed for stochastic version of ADAM and GRAD (default = 1000) <code>trace</code> : -1 return results from all iterations, 0 (default) return final result

Value

A list with the following elements:

<code>beta</code>	matrix of regression coefficients
<code>all.beta</code>	coefficients from all iterations for GRAD and ADAM
<code>spars</code>	smoothing parameters from <code>stats::smooth.spline()</code> for initialization
<code>fit</code>	object from the optimization algorithm

Examples

```
data(engel)
y <- engel$foodexp
X <- cbind(rep(1,length(y)),engel$income-mean(engel$income))
tau <- seq(0.1,0.9,0.05)
fit.rq <- quantreg::rq(y ~ X[,2],tau)
fit.sqr <- sqr(y ~ X[,2],tau,d=2,spar=0.2)
fit <- sqr.fit.optim(X,y,tau,spar=0.2,d=2,method="BFGS",beta.rq=fit.rq$coef)
fit <- sqr.fit.optim(X,y,tau,spar=0.2,d=2,method="BFGS",beta.rq=fit.rq$coef)
par(mfrow=c(1,2),pty="m",lab=c(10,10,2),mar=c(4,4,2,1)+0.1,las=1)
```

```

for(j in c(1:2)) {
  plot(tau,fit.rq$coef[j,],type="n",xlab="QUANTILE LEVEL",ylab=paste0("COEFF",j))
  points(tau,fit.rq$coef[j,],pch=1,cex=0.5)
  lines(tau,fit.sqr$coef[j,],lty=1); lines(tau,fit$beta[j,],lty=2,col=2)
}

```

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 $f_0 = 0.5$

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')

```

tsqr.fit

*Trigonometric Spline Quantile Regression (TSQR) of Time Series***Description**

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

Usage

```
tsqr.fit(  
  y,  
  f0,  
  tau,  
  spar = 1,  
  d = 1,  
  weighted = FALSE,  
  mthreads = TRUE,  
  prepared = TRUE,  
  ztol = 1e-05  
)
```

Arguments

y	time series
f0	frequency in [0,1)
tau	sequence of quantile levels in (0,1)
spar	smoothing parameter
d	subsampling rate of quantile levels (default = 1)
weighted	if TRUE, penalty function is weighted (default = FALSE)
mthreads	if FALSE, set RnpcBLASctl::blas_set_num_threads(1) (default = TRUE)
prepared	if TRUE, intercept is removed and coef of cosine is doubled when f0 = 0.5
ztol	zero tolerance parameter used to determine the effective dimensionality of the fit

Value

object of `sqr.fit()` (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)  
fit.sqr <- tsqr.fit(y,f0=0.1,tau=tau,spar=1,d=4)  
plot(tau,fit$coef[1,],type='p',xlab='QUANTILE LEVEL',ylab='TQR COEF')  
lines(tau,fit.sqr$coef[1,],type='l')
```

yearssn	<i>Yearly sunspot numbers</i>
---------	-------------------------------

Description

Sunspot numbers from 1700 to 2007.

Usage

```
data(yearssn)
```

Format

An object of class `data.frame` with 308 rows and 2 columns.

References

Li, T.-H. (2014). Time Series with Mixed Spectra. CRC Press.

Index

* datasets

birthweight, [2](#)

engel, [3](#)

yearssn, [30](#)

birthweight, [2](#)

engel, [3](#)

per, [3](#)

qacf, [4](#)

qcser, [4](#)

qdft, [5](#)

qdft2qacf, [6](#)

qdft2qper, [7](#)

qdft2qser, [7](#)

qfa.plot, [8](#)

qkl.divergence, [9](#)

qper, [9](#)

qper2, [10](#)

qser, [11](#)

qser2ar, [12](#)

qser2qacf, [12](#)

qser2sar, [13](#)

qspec.ar, [14](#)

qspec.lw, [15](#)

qspec.sar, [16](#)

qspec2qcoh, [18](#)

sar.eq.bootstrap, [18](#)

sar.eq.test, [19](#)

sar.gc.bootstrap, [20](#)

sar.gc.coef, [21](#)

sar.gc.test, [22](#)

sqdft, [23](#)

sqdft.fit, [24](#)

sqr, [25](#)

sqr.fit, [26](#)

sqr.fit.optim, [27](#)

tqr.fit, [29](#)

tsqr.fit, [29](#)

yearssn, [30](#)