Package 'bsplinePsd'

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Description Implementation of a Metropolis-within-Gibbs MCMC algorithm to flexibly estimate the spectral density of a stationary time series. The algorithm updates a nonparametric B-spline prior using the Whittle likelihood to produce pseudo-posterior samples and is based on the work presented in Edwards, M.C., Meyer, R. and Christensen, N., Statitics and Computing (2018). <doi.org 10.1007="" s11222-017-9796-9="">.</doi.org>
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bsplinePsd-package	Bayesian Nonparametric Spectral Density Estimation Using B-Spline Priors
	Priors

Description

Implementation of a Metropolis-within-Gibbs MCMC algorithm to flexibly estimate the spectral density of a stationary time series. The algorithm updates a nonparametric B-spline prior using the Whittle likelihood to produce pseudo-posterior samples.

Details

The function gibbs_bspline is an implementation of the (serial version of the) MCMC algorithm presented in Edwards et al. (2018). This algorithm uses a nonparametric B-spline prior to estimate the spectral density of a stationary time series and can be considered a generalisation of the algorithm of Choudhuri et al. (2004), which used the Bernstein polynomial prior. A Dirichlet process prior is used to find the weights for the B-spline densities used in the finite mixture and a seperate and independent Dirichlet process prior used to place knots. The algorithm therefore allows for a data-driven choice of the number of knots/mixture components and their locations.

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References

Edwards, M. C., Meyer, R., and Christensen, N. (2018), Bayesian nonparametric spectral density estimation using B-spline priors, *Statistics and Computing*, https://doi.org/10.1007/s11222-017-9796-9.

Choudhuri, N., Ghosal, S., and Roy, A. (2004), Bayesian estimation of the spectral density of a time series, *Journal of the American Statistical Association*, 99(468):1050–1059.

dbspline

Generate a B-spline density basis of any degree

Description

This function generates a B-spline density basis of any degree.

Usage

```
dbspline(x, knots, degree = 3)
```

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Arguments

x numeric vector for which the B-spline densities are to be generated

knots used to generate the B-spline densities

degree positive integer specifying the degree of the B-spline densities (default is 3 for

cubic B-splines)

Details

splineDesign is used to generate a B-spline basis of any degree. Each B-spline is then normalised to become a B-spline density using analytical integration. Note that the two boundary knots (0 and 1) are each coincident degree + 1 times.

Value

matrix of the B-spline density basis

See Also

splineDesign

Examples

```
## Not run:

# Generate basis functions
set.seed(1)
x = seq(0, 1, length = 256)
knots = sort(c(0, runif(10), 1))
basis = dbspline(x, knots)

# Plot basis functions
plot(x, basis[1, ], type = "1", ylim = c(min(basis), max(basis)),
    ylab = expression(b[3](x)), main = "Cubic B-spline Density Basis Functions")
for (i in 2:nrow(basis)) lines(x, basis[i, ], col = i)

## End(Not run)
```

gibbs_bspline

Metropolis-within-Gibbs sampler for spectral inference of a stationary time series using a B-spline prior

Description

This function updates the B-spline prior using the Whittle likelihood and obtains samples from the pseudo-posterior to infer the spectral density of a stationary time series.

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Usage

```
gibbs_bspline(data, Ntotal, burnin, thin = 1, k.theta = 0.01, MG = 1,
G0.alpha = 1, G0.beta = 1, LG = 20, MH = 1, H0.alpha = 1,
H0.beta = 1, LH = 20, tau.alpha = 0.001, tau.beta = 0.001,
kmax = 100, k1 = 20, degree = 3)
```

Arguments

data numeric vector

Ntotal total number of iterations to run the Markov chain

burnin number of initial iterations to be discarded

thin thinning number (post-processing)

k. theta prior parameter for number of B-spline densities k (proportional to exp(-k.theta*k^2))

in mixture

MG Dirichlet process base measure constant for weights of B-spline densities in mix-

ture (>0)

G0.alpha, G0.beta

parameters of Beta base measure of Dirichlet process for weights of B-spline

densities in mixture (default is Uniform[0, 1])

LG truncation parameter of Dirichlet process in stick breaking representation for

weights of B-spline densities

MH Dirichlet process base measure constant for knot placements of B-spline densi-

ties (>0)

H0.alpha, H0.beta

parameters of Beta base measure of Dirichlet process for knot placements of

B-spline densities (default is Uniform[0, 1])

LH truncation parameter of Dirichlet process in stick breaking representation for

knot placements of B-spline densities

tau.alpha, tau.beta

prior parameters for tau (Inverse-Gamma)

kmax upper bound for number of B-spline densities in mixture

k1 starting value for k. If k1 = NA then a random starting value between degree +

2 and kmax is selected

degree positive integer specifying the degree of the B-spline densities (default is 3)

Details

The function gibbs_bspline is an implementation of the (serial version of the) MCMC algorithm presented in Edwards et al. (2018). This algorithm uses a nonparametric B-spline prior to estimate the spectral density of a stationary time series and can be considered a generalisation of the algorithm of Choudhuri et al. (2004), which used the Bernstein polynomial prior. A Dirichlet process prior is used to find the weights for the B-spline densities used in the finite mixture and a seperate and independent Dirichlet process prior used to place knots. The algorithm therefore allows for a data-driven choice of the number of knots/mixtures and their locations.

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Value

```
A list with S3 class 'psd' containing the following components:
psd.median,psd.mean
                  psd estimates: (pointwise) posterior median and mean
psd.p05,psd.p95
                  90% pointwise credibility interval
psd.u05,psd.u95
                  90% uniform credibility interval
k,tau,V,Z,U,X
                  posterior traces of model parameters
knots.trace
                  trace of knot placements
                  trace of log likelihood
ll.trace
pdgrm
                  periodogram
                  integer length of input time series
```

References

Edwards, M. C., Meyer, R., and Christensen, N. (2018), Bayesian nonparametric spectral density estimation using B-spline priors, *Statistics and Computing*, https://doi.org/10.1007/s11222-017-9796-9.

Choudhuri, N., Ghosal, S., and Roy, A. (2004), Bayesian estimation of the spectral density of a time series, *Journal of the American Statistical Association*, 99(468):1050–1059.

See Also

plot.psd

Examples

```
## Not run:
set.seed(123456)

# Generate AR(1) data with rho = 0.9
n = 128
data = arima.sim(n, model = list(ar = 0.9))
data = data - mean(data)

# Run MCMC (may take some time)
mcmc = gibbs_bspline(data, 10000, 5000)

require(beyondWhittle)  # For psd_arma() function
freq = 2 * pi / n * (1:(n / 2 + 1) - 1)[-c(1, n / 2 + 1)]  # Remove first and last frequency
psd.true = psd_arma(freq, ar = 0.9, ma = numeric(0), sigma2 = 1)  # True PSD
plot(mcmc)  # Plot log PSD (see documentation of plot.psd)
lines(freq, log(psd.true), col = 2, lty = 3, lwd = 2)  # Overlay true PSD

## End(Not run)
```

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plot.psd

Plot method for psd class

Description

This function plots the log periodogram, log posterior median PSD, and log 90% credible region PSD. The x-axis uses angular frequency and the y-axis is plotted on the log scale. The PSD at the zero frequency is removed from the plot. If the time series is even length, the PSD at the last frequency is also removed from the plot.

Usage

```
## S3 method for class 'psd'
plot(x, legend.loc = "topright", ylog = TRUE, ...)
```

Arguments

x	an object of class psd
legend.loc	location of legend out of "topright" (default), "topleft", "bottomright", "bottom-left". If set to NA then no legend will be produced
ylog	logical value (default is TRUE) to determine if PSD (y-axis) should be on natural log scale $$
	other graphical parameters from the plot.default function

Value

plot of the estimate of the (log) PSD

See Also

```
gibbs_bspline
```

Examples

```
## Not run:
set.seed(12345)

# Simulate AR(4) data
n = 2 ^ 7
ar.ex = c(0.9, -0.9, 0.9, -0.9)
data = arima.sim(n, model = list(ar = ar.ex))
data = data - mean(data)

# Run MCMC with linear B-spline prior (may take some time)
mcmc = gibbs_bspline(data, 10000, 5000, degree = 1)

# Plot result
```

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```
plot(mcmc)

# Plot result on original scale with title
plot(mcmc, ylog = FALSE, main = "Estimate of PSD using the linear B-spline prior")

## End(Not run)
```

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