

# Package ‘BayesSpec’

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**Type** Package

**Title** Bayesian Spectral Analysis Techniques

**Version** 0.5.3

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**Description** An implementation of methods for spectral analysis using the Bayesian framework. It includes functions for modelling spectrum as well as appropriate plotting and output estimates. There is segmentation capability with RJ MCMC (Reversible Jump Markov Chain Monte Carlo). The package takes these methods predominantly from the 2012 paper “AdaptSPEC: Adaptive Spectral Estimation for Nonstationary Time Series” <DOI:10.1080/01621459.2012.716340>.

**Imports** mvtnorm (>= 1.0-5), pscl (>= 1.4.9), trust (>= 0.1-7)

**License** GPL-3

**LazyLoad** TRUE

**LazyData** TRUE

**RoxygenNote** 6.0.1

**NeedsCompilation** no

**Repository** CRAN

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 adaptspec

*Adaptive Spectral Estimation for Non-stationary Time Series*


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### Description

Methodology for analyzing possibly non-stationary time series by adaptively dividing the time series into an unknown but finite number of segments and estimating the corresponding local spectra by smoothing splines.

### Usage

```
adaptspec(nloop, nwarmup, nexp_max, x,
          tmin, sigmasqalpha, tau_prior_a, tau_prior_b,
          tau_up_limit, prob_mm1, step_size_max,
          var_inflate, nbasis, nfreq_hat, plotting)
```

### Arguments

nloop	The total number of MCMC iterations
nwarmup	The number of burn-in iterations
nexp_max	The maximum number of segments allowed
x	The data, a univariate time series, not a time series object
tmin	The minimum number of observations per segment. An optional argument defaulted to $tmin = 40$ .
sigmasqalpha	An optional argument defaulted to $sigmasqalpha = 100$ .
tau_prior_a	An optional argument defaulted to $tau\_prior\_a = -1$ .
tau_prior_b	An optional argument defaulted to $tau\_prior\_b = 0$ .
tau_up_limit	An optional argument defaulted to $tau\_up\_limit = 10000$ .
prob_mm1	An optional argument defaulted to $prob\_mm1 = 0.8$ .
step_size_max	An optional argument defaulted to $step\_size\_max = 10$ .
var_inflate	An optional argument defaulted to $var\_inflate = 1$ .
nbasis	An optional argument defaulted to $nbasis = 7$ .
nfreq_hat	An optional argument defaulted to $nfreq\_hat = 50$ .
plotting	An optional argument for displaying output plots defaulted to FALSE. When set to TRUE, this displays the spectral and partition points.

### Value

xi The partition points  
 log\_spec\_hat Estimates of the log spectra for all segments  
 nexp\_curr The number of segments in each iteration.

**Author(s)**

Rosen, O., Wood, S. and Stoffer, D.

**References**

Rosen, O., Wood, S. and Stoffer, D. (2012). AdaptSPEc: Adaptive Spectral Estimation for Nonstationary Time Series. *J. of the American Statistical Association*, 107, 1575-1589

**Examples**

```
#Running adaptspec with the simulated_pieewise data.
data(simulated_pieewise)
model1 <- adaptspec(nloop = 80, nwarmup = 20,
  nexpt_max = 5, x = simulated_pieewise[1:100])
str(model1)
summary(model1$nexpt_curr)
plot(model1$nexpt_curr)
```

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intracranial\_eeg

*Intracranial Electroencephalograph (IEEG) Dataset*

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**Description**

A sample of IEEG data from a subject in an interictal state.

**Usage**

```
data(intracranial_eeg)
```

**Format**

A vector time series of 6,000 observations of intracranial electroencephalograph

**Source**

kaggle.com

**References**

<https://www.kaggle.com/c/melbourne-university-seizure-prediction>

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simulated\_piecewise     *Simulated Piecewise Time Series Dataset*

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**Description**

This dataset is simulated from a piecewise autoregressive process (model (11), p. 1581, in Rosen et al. (2012)), see Examples.

**Usage**

```
data(simulated_piecewise)
```

**Format**

A univariate numeric vector with 1,024 observations.

**Source**

Simulated

**References**

Rosen, O., Wood, S. and Stoffer, D. (2012). AdaptSPEC: Adaptive Spectral Estimation for Nonstationary Time Series. *J. of the American Statistical Association*, 107, 1575-1589

**Examples**

```
#Created using the following script:
set.seed(346)
phi_true <- matrix(list(),3,1)
phi_true[[1]] <- .9
phi_true[[2]] <- c(1.69, -.81)
phi_true[[3]] <- c(1.32, -.81)
sd_true <- rep(1,3)
x1 <- arima.sim(list(order=c(1,0,0), ar=phi_true[[1]]),512,sd=sd_true[1])
x2 <- arima.sim(list(order=c(2,0,0), ar=phi_true[[2]]),256,sd=sd_true[2])
x3 <- arima.sim(list(order=c(2,0,0), ar=phi_true[[3]]),256,sd=sd_true[3])
simulated_piecewise <- c(x1, x2, x3)
plot.ts(simulated_piecewise)
```

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