

Package ‘bsts’

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Title Bayesian Structural Time Series

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add.ar	<i>AR(p) state component</i>
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Description

Add an AR(p) state component to the state specification.

Usage

```
AddAr(state.specification,
      y,
      lags = 1,
      sigma.prior,
      initial.state.prior = NULL,
      sdy)
```

Arguments

`state.specification` A list of state components. If omitted, an empty list is assumed.

`y` A numeric vector. The time series to be modeled.

`lags` The number of lags ("p") in the AR(p) process.

`sigma.prior` An object created by `SdPrior`. The prior for the standard deviation of the process increments.

`initial.state.prior` An object of class `MvnPrior` describing the values of the state at time 0. This argument can be `NULL`, in which case the stationary distribution of the AR(p) process will be used as the initial state distribution.

`sdy` The sample standard deviation of the time series to be modeled. Used to scale the prior distribution. This can be omitted if `y` is supplied.

Details

The model is

$$\alpha_t = \phi_1 \alpha_{i,t-1} + \dots + \phi_p \alpha_{i,t-p} + \epsilon_{t-1} \quad \epsilon_t \sim \mathcal{N}(0, \sigma^2)$$

The state consists of the last `p` lags of `alpha`. The state transition matrix has `phi` in its first row, ones along its first subdiagonal, and zeros elsewhere. The state variance matrix has `sigma^2` in its upper left corner and is zero elsewhere. The observation matrix has 1 in its first element and is zero otherwise.

Value

Returns `state.specification` with an AR(p) state component added to the end.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also[bsts.SdPrior](#)**Examples**

```

n <- 100
residual.sd <- .001

# Actual values of the AR coefficients
true.phi <- c(-.7, .3, .15)
ar <- arima.sim(model = list(ar = true.phi),
                n = n,
                sd = 3)

## Layer some noise on top of the AR process.
y <- ar + rnorm(n, 0, residual.sd)
ss <- AddAr(list(), lags = 3, sigma.prior = SdPrior(3.0, 1.0))

# Fit the model with knowledge with residual.sd essentially fixed at the
# true value.
model <- bsts(y, state.specification=ss, niter = 500, prior = SdPrior(residual.sd, 100000))

# Now compare the empirical ACF to the true ACF.
acf(y, lag.max = 30)
points(0:30, ARMAacf(ar = true.phi, lag.max = 30), pch = "+")
points(0:30, ARMAacf(ar = colMeans(model$AR3.coefficients), lag.max = 30))
legend("topright", leg = c("empirical", "truth", "MCMC"), pch = c(NA, "+", "o"))

```

add.dynamic.regression

Dynamic Regression State Component

Description

Add a dynamic regression component to the state specification of a bsts model.

Usage

```

AddDynamicRegression(
  state.specification,
  formula,
  data,
  sigma.mean.prior = NULL,
  shrinkage.parameter.prior = GammaPrior(a = 10, b = 1),
  contrasts = NULL,
  na.action = na.pass)

```

Arguments

<code>state.specification</code>	A list of state components that you wish to add to. If omitted, an empty list will be assumed.
<code>formula</code>	A formula describing the regression portion of the relationship between y and X. If no regressors are desired then the formula can be replaced by a numeric vector giving the time series to be modeled.
<code>data</code>	An optional data frame, list or environment (or object coercible by <code>as.data.frame</code> to a data frame) containing the variables in the model. If not found in data, the variables are taken from <code>'environment(formula)'</code> , typically the environment from which <code>AddDynamicRegression</code> is called.
<code>sigma.mean.prior</code>	An object created by <code>GammaPrior</code> describing the prior distribution of b/a (see details below).
<code>shrinkage.parameter.prior</code>	An object of class <code>GammaPrior</code> describing the shrinkage parameter, a (see details below).
<code>contrasts</code>	An optional list. See the <code>contrasts.arg</code> of <code>model.matrix.default</code> . This argument is only used if a model formula is specified. It can usually be ignored even then.
<code>na.action</code>	What to do about missing values. The default is to allow missing responses, but no missing predictors. Set this to <code>na.omit</code> or <code>na.exclude</code> if you want to omit missing responses altogether.

Details

The model is

$$\beta_{i,t+1} = \beta_{i,t} + \epsilon_t \quad \epsilon_t \sim \mathcal{N}(0, \sigma_i^2 / \text{variance}_{xi})$$

$$\frac{1}{\sigma_i^2} \sim Ga(a, b)$$

$$\sqrt{b/a} \sim \text{sigma.mean.prior}$$

$$a \sim \text{shrinkage.parameter.prior}$$

That is, each coefficient evolves independently, with its own variance term which is scaled by the variance of the *i*'th column of X. The parameters of the hyperprior are interpretable as: `sqrt(b/a)` typical amount that a coefficient might change in a single time period, and '`a`' is the 'sample size' or 'shrinkage parameter' measuring the degree of similarity in `sigma[i]` among the arms.

In most cases we hope `b/a` is small, so that `sigma[i]`'s will be small and the series will be forecastable. We also hope that '`a`' is large because it means that the `sigma[i]`'s will be similar to one another.

The default prior distribution is a pair of independent Gamma priors for `sqrt(b/a)` and `a`. The mean of `sigma[i]` is set to `.01 * sd(y)` with shape parameter equal to 1. The mean of the shrinkage parameter is set to 10, but with shape parameter equal to 1.

Value

Returns a list with the elements necessary to specify a dynamic regression model.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.SdPrior](#) [NormalPrior](#)

Examples

```
## Setting the seed to avoid small sample effects resulting from small
## number of iterations.
set.seed(8675309)
n <- 1000
x <- matrix(rnorm(n))

# beta follows a random walk with sd = .1 starting at -12.
beta <- cumsum(rnorm(n, 0, .1)) - 12

# level is a local level model with sd = 1 starting at 18.
level <- cumsum(rnorm(n)) + 18

# sigma.obs is .1
error <- rnorm(n, 0, .1)

y <- level + x * beta + error
par(mfrow = c(1, 3))
plot(y, main = "Raw Data")
plot(x, y - level, main = "True Regression Effect")
plot(y - x * beta, main = "Local Level Effect")

ss <- list()
ss <- AddLocalLevel(ss, y)
ss <- AddDynamicRegression(ss, y ~ x)
## In a real application you'd probably want more than 100
## iterations. See comment above about the random seed.
model <- bsts(y, state.specification = ss, niter = 100, seed = 8675309)
plot(model, "dynamic", burn = 10)
```

`add.holiday`*Holiday state models*

Description

Add a random-walk holiday model to a state specification.

This model allows each active day in a holiday window to move according to a random walk relative to the day's value the last time the holiday occurred.

Usage

```
AddFixedDateHoliday(state.specification = NULL,  
                    holiday.name,  
                    month,  
                    day,  
                    y,  
                    sigma.prior = NULL,  
                    initial.state.prior = NULL,  
                    sdy = sd(as.numeric(y), na.rm = TRUE),  
                    time0 = NULL,  
                    days.before = 1,  
                    days.after = 1)
```

```
AddNthWeekdayInMonthHoliday(state.specification = NULL,  
                             holiday.name,  
                             month,  
                             day.of.week,  
                             which.week,  
                             y,  
                             sigma.prior = NULL,  
                             initial.state.prior = NULL,  
                             sdy = sd(as.numeric(y), na.rm = TRUE),  
                             time0 = NULL,  
                             days.before = 1,  
                             days.after = 1)
```

```
AddLastWeekdayInMonthHoliday(state.specification = NULL,  
                              holiday.name,  
                              month,  
                              day.of.week,  
                              y,  
                              sigma.prior = NULL,  
                              initial.state.prior = NULL,  
                              sdy = sd(as.numeric(y), na.rm = TRUE),  
                              time0 = NULL,
```

```
days.before = 1,
days.after = 1)
```

```
NamedHolidays(except = NULL)
```

```
AddNamedHolidays(state.specification = NULL,
  named.holidays = NamedHolidays(),
  y,
  sigma.prior = NULL,
  initial.state.prior = NULL,
  sdy = sd(as.numeric(y), na.rm = TRUE),
  time0 = NULL,
  days.before = 1,
  days.after = 1)
```

Arguments

<code>state.specification</code>	A list of state components that you wish augment. If omitted, an empty list will be assumed.
<code>holiday.name</code>	A string that can be used to label the holiday in output.
<code>named.holidays</code>	A character vector containing one or more recognized holiday names.
<code>y</code>	The time series to be modeled, as a numeric vector convertible to <code>xts</code> . This state model assumes <code>y</code> contains daily data.
<code>sigma.prior</code>	An object created by <code>SdPrior</code> describing the prior distribution for the standard deviation of the random walk increments.
<code>initial.state.prior</code>	An object created using <code>NormalPrior</code> , describing the prior distribution of the the initial state vector (at time 1).
<code>sdy</code>	The standard deviation of the series to be modeled. This will be ignored if <code>y</code> is provided, or if all the required prior distributions are supplied directly.
<code>time0</code>	An object convertible to <code>POSIXt</code> containing the date of the initial observation in the training data. If omitted and <code>y</code> is a <code>zoo</code> or <code>xts</code> object, then <code>time0</code> will be obtained from the index of <code>y[1]</code> .
<code>days.before</code>	An integer giving the number of days the influence of the holiday extends prior to the actual holiday.
<code>days.after</code>	An integer giving the number of days the influence of the holiday extends after the actual holiday.
<code>month</code>	A string naming the month in which the holiday occurs. Unambiguous partial matches are acceptable. Capitalize the first letter.
<code>day</code>	An integer specifying the day of the month on which the <code>FixedDateHoliday</code> occurs.
<code>day.of.week</code>	A string giving the weekday on which the <code>NthWeekdayInMonthHoliday</code> occurs.
<code>which.week</code>	An integer specifying the week of the month on which the <code>NthWeekdayInMonthHoliday</code> occurs. If <code>which.week <= 0</code> then the holiday is assumed to occur on the last <code>day.of.week</code> in <code>month</code> .

except If NULL then all named holidays are returned. If except is a character vector containing partial matches to holiday names, the matched names will be omitted from the returned list.

Value

NamedHolidays returns a character vector with the names of the recognized holiday.

The other functions return state.specification, after adding the requested state components.

AddNthWeekdayInMonthHoliday, AddLastWeekdayInMonthHoliday, and AddFixedDateHoliday can each add one holiday at a time. AddNamedHolidays will add several holidays at once, if named.holidays is a vector.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.SdPrior](#) [NormalPrior](#)

Examples

```
## TODO(stevescott): add examples
```

add.local.level	<i>Local level trend state component</i>
-----------------	--

Description

Add a local level model to a state specification. The local level model assumes the trend is a random walk:

$$\alpha_{t+1} = \alpha_t + \epsilon_t \quad \epsilon_t \sim \mathcal{N}(0, \sigma).$$

The prior is on the σ parameter.

Usage

```
AddLocalLevel(  
  state.specification,  
  y,  
  sigma.prior,  
  initial.state.prior,  
  sdy,  
  initial.y)
```

Arguments

<code>state.specification</code>	A list of state components that you wish to add to. If omitted, an empty list will be assumed.
<code>y</code>	The time series to be modeled, as a numeric vector.
<code>sigma.prior</code>	An object created by SdPrior describing the prior distribution for the standard deviation of the random walk increments.
<code>initial.state.prior</code>	An object created using NormalPrior , describing the prior distribution of the the initial state vector (at time 1).
<code>sdy</code>	The standard deviation of the series to be modeled. This will be ignored if <code>y</code> is provided, or if all the required prior distributions are supplied directly.
<code>initial.y</code>	The initial value of the series being modeled. This will be ignored if <code>y</code> is provided, or if the priors for the initial state are all provided directly.

Value

Returns a list with the elements necessary to specify a local linear trend state model.

Author(s)

Steven L. Scott <stevescott@google.com>

References

- Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.
- Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.SdPrior](#) [NormalPrior](#)

 add.local.linear.trend

Local linear trend state component

Description

Add a local linear trend model to a state specification. The local linear trend model assumes that both the mean and the slope of the trend follow random walks. The equation for the mean is

$$\mu_{t+1} = \mu_t + \delta_t + \epsilon_t \quad \epsilon_t \sim \mathcal{N}(0, \sigma_\mu).$$

The equation for the slope is

$$\delta_{t+1} = \delta_t + \eta_t \quad \eta_t \sim \mathcal{N}(0, \sigma_\delta).$$

The prior distribution is on the level standard deviation σ_μ and the slope standard deviation σ_δ .

Usage

```
AddLocalLinearTrend(
  state.specification = NULL,
  y,
  level.sigma.prior = NULL,
  slope.sigma.prior = NULL,
  initial.level.prior = NULL,
  initial.slope.prior = NULL,
  sdy,
  initial.y)
```

Arguments

- state.specification
A list of state components that you wish to add to. If omitted, an empty list will be assumed.
- y
The time series to be modeled, as a numeric vector.
- level.sigma.prior
An object created by [SdPrior](#) describing the prior distribution for the standard deviation of the level component.
- slope.sigma.prior
An object created by [SdPrior](#) describing the prior distribution of the standard deviation of the slope component.
- initial.level.prior
An object created by [NormalPrior](#) describing the initial distribution of the level portion of the initial state vector.
- initial.slope.prior
An object created by [NormalPrior](#) describing the prior distribution for the slope portion of the initial state vector.

<code>sd</code>	The standard deviation of the series to be modeled. This will be ignored if <code>y</code> is provided, or if all the required prior distributions are supplied directly.
<code>initial.y</code>	The initial value of the series being modeled. This will be ignored if <code>y</code> is provided, or if the priors for the initial state are all provided directly.

Value

Returns a list with the elements necessary to specify a local linear trend state model.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.SdPrior](#) [NormalPrior](#)

Examples

```
data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bst(y, state.specification = ss, niter = 500)
pred <- predict(model, horizon = 12, burn = 100)
plot(pred)
```

`add.seasonal`

Seasonal state component

Description

Add a seasonal model to a state specification.

The seasonal model can be thought of as a regression on `nseasons` dummy variables with coefficients constrained to sum to 1 (in expectation). If there are S seasons then the state vector γ is of dimension $S-1$. The first element of the state vector obeys

$$\gamma_{t+1,1} = - \sum_{i=2}^S \gamma_{t,i} + \epsilon_t \quad \epsilon_t \sim \mathcal{N}(0, \sigma)$$

Usage

```
AddSeasonal(  
  state.specification,  
  y,  
  nseasons,  
  season.duration = 1,  
  sigma.prior,  
  initial.state.prior,  
  sdy)
```

Arguments

<code>state.specification</code>	A list of state components that you wish to add to. If omitted, an empty list will be assumed.
<code>y</code>	The time series to be modeled, as a numeric vector.
<code>nseasons</code>	The number of seasons to be modeled.
<code>season.duration</code>	The number of time periods in each season.
<code>sigma.prior</code>	An object created by SdPrior describing the prior distribution for the standard deviation of the random walk increments.
<code>initial.state.prior</code>	An object created using NormalPrior , describing the prior distribution of the the initial state vector (at time 1).
<code>sdy</code>	The standard deviation of the series to be modeled. This will be ignored if <code>y</code> is provided, or if all the required prior distributions are supplied directly.

Value

Returns a list with the elements necessary to specify a seasonal state model.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts](#). [SdPrior](#) [NormalPrior](#)

Examples

```

data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bst(y, state.specification = ss, niter = 500)
pred <- predict(model, horizon = 12, burn = 100)
plot(pred)

```

```
add.semilocal.linear.trend
```

Semilocal Linear Trend

Description

The semi-local linear trend model is similar to the local linear trend, but more useful for long-term forecasting. It assumes the level component moves according to a random walk, but the slope component moves according to an AR1 process centered on a potentially nonzero value D . The equation for the level is

$$\mu_{t+1} = \mu_t + \delta_t + \epsilon_t \quad \epsilon_t \sim \mathcal{N}(t, \sigma_\mu).$$

The equation for the slope is

$$\delta_{t+1} = D + \phi(\delta_t - D) + \eta_t \quad \eta_t \sim \mathcal{N}(t, \sigma_\delta).$$

This model differs from the local linear trend model in that the latter assumes the slope δ_t follows a random walk. A stationary AR(1) process is less variable than a random walk when making projections far into the future, so this model often gives more reasonable uncertainty estimates when making long term forecasts.

The prior distribution for the semi-local linear trend has four independent components. These are:

- an inverse gamma prior on the level standard deviation σ_μ ,
- an inverse gamma prior on the slope standard deviation σ_δ ,
- a Gaussian prior on the long run slope parameter D ,
- and a potentially truncated Gaussian prior on the AR1 coefficient ϕ . If the prior on ϕ is truncated to $(-1, 1)$, then the slope will exhibit short term stationary variation around the long run slope D .

Usage

```

AddSemilocalLinearTrend(
  state.specification = list(),
  y = NULL,
  level.sigma.prior = NULL,
  slope.mean.prior = NULL,

```

```
slope.ar1.prior = NULL,  
slope.sigma.prior = NULL,  
initial.level.prior = NULL,  
initial.slope.prior = NULL,  
sdy = NULL,  
initial.y = NULL)
```

Arguments

state.specification	A list of state components that you wish to add to. If omitted, an empty list will be assumed.
y	The time series to be modeled, as a numeric vector. This can be omitted if sdy and initial.y are supplied, or if all prior distributions are supplied directly.
level.sigma.prior	An object created by SdPrior describing the prior distribution for the standard deviation of the level component.
slope.mean.prior	An object created by NormalPrior giving the prior distribution for the mean parameter in the generalized local linear trend model (see below).
slope.ar1.prior	An object created by Ar1CoefficientPrior giving the prior distribution for the ar1 coefficient parameter in the generalized local linear trend model (see below).
slope.sigma.prior	An object created by SdPrior describing the prior distribution of the standard deviation of the slope component.
initial.level.prior	An object created by NormalPrior describing the initial distribution of the level portion of the initial state vector.
initial.slope.prior	An object created by NormalPrior describing the prior distribution for the slope portion of the initial state vector.
sdy	The standard deviation of the series to be modeled. This will be ignored if y is provided, or if all the required prior distributions are supplied directly.
initial.y	The initial value of the series being modeled. This will be ignored if y is provided, or if the priors for the initial state are all provided directly.

Value

Returns a list with the elements necessary to specify a generalized local linear trend state model.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.SdPrior NormalPrior](#)

```
add.student.local.linear.trend
      Robust local linear trend
```

Description

Add a local level model to a state specification. The local linear trend model assumes that both the mean and the slope of the trend follow random walks. The equation for the mean is

$$\mu_{t+1} = \mu_t + \delta_t + \epsilon_t \quad \epsilon_t \sim \mathcal{T}_{\nu_\mu}(0, \sigma_\mu).$$

The equation for the slope is

$$\delta_{t+1} = \delta_t + \eta_t \quad \eta_t \sim \mathcal{T}_{\nu_\delta}(0, \sigma_\delta).$$

Independent prior distributions are assumed on the level standard deviation, σ_μ the slope standard deviation σ_δ , the level tail thickness ν_μ , and the slope tail thickness ν_δ .

Usage

```
AddStudentLocalLinearTrend(
  state.specification = NULL,
  y,
  save.weights = FALSE,
  level.sigma.prior = NULL,
  level.nu.prior = NULL,
  slope.sigma.prior = NULL,
  slope.nu.prior = NULL,
  initial.level.prior = NULL,
  initial.slope.prior = NULL,
  sdy,
  initial.y)
```


Arguments

<code>state.specification</code>	A list of state components that you wish to add to. If omitted, an empty list will be assumed.
<code>y</code>	The time series to be modeled, as a numeric vector.
<code>save.weights</code>	A logical value indicating whether to save the draws of the weights from the normal mixture representation.
<code>level.sigma.prior</code>	An object created by <code>SdPrior</code> describing the prior distribution for the standard deviation of the level component.
<code>level.nu.prior</code>	An object inheriting from the class <code>DoubleModel</code> , representing the prior distribution on the nu tail thickness parameter of the T distribution for errors in the evolution equation for the level component.
<code>slope.sigma.prior</code>	An object created by <code>SdPrior</code> describing the prior distribution of the standard deviation of the slope component.
<code>slope.nu.prior</code>	An object inheriting from the class <code>DoubleModel</code> , representing the prior distribution on the nu tail thickness parameter of the T distribution for errors in the evolution equation for the slope component.
<code>initial.level.prior</code>	An object created by <code>NormalPrior</code> describing the initial distribution of the level portion of the initial state vector.
<code>initial.slope.prior</code>	An object created by <code>NormalPrior</code> describing the prior distribution for the slope portion of the initial state vector.
<code>sd</code>	The standard deviation of the series to be modeled. This will be ignored if <code>y</code> is provided, or if all the required prior distributions are supplied directly.
<code>initial.y</code>	The initial value of the series being modeled. This will be ignored if <code>y</code> is provided, or if the priors for the initial state are all provided directly.

Value

Returns a list with the elements necessary to specify a local linear trend state model.

Author(s)

Steven L. Scott <stevescott@google.com>

References

- Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.
- Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.SdPrior NormalPrior](#)

Examples

```
data(rsxfs)
ss <- AddStudentLocalLinearTrend(list(), rsxfs)
model <- bsts(rsxfs, state.specification = ss, niter = 500)
pred <- predict(model, horizon = 12, burn = 100)
plot(pred)
```

add.trig

Trigonometric seasonal state component

Description

Add a trigonometric seasonal model to a state specification.

The trig component adds a collection of sine and cosine terms with randomly varying coefficients to the state model. The coefficients are the states, while the sine and cosine values are part of the "observation matrix".

This state component adds the sum of its terms to the observation equation.

$$y_t = \sum_j \beta_{jt} \sin(f_j t) + \gamma_{jt} \cos(f_j t)$$

The evolution equation is that each of the sinusoid coefficients follows a random walk with standard deviation σ_j .

$$\beta_{jt} = \beta_{jt-1} + N(0, \sigma_{s_j}^2) \quad \gamma_{jt} = \gamma_{jt-1} + N(0, \sigma_{c_j}^2)$$

Usage

```
AddTrig(
  state.specification = NULL,
  y,
  period,
  frequencies,
  sigma.prior = NULL,
  initial.state.prior = NULL,
  sdy)
```

Arguments

state.specification	A list of state components that you wish to add to. If omitted, an empty list will be assumed.
y	The time series to be modeled, as a numeric vector.
period	A positive scalar giving the number of time steps required for the longest cycle to repeat.
frequencies	A vector of positive real numbers giving the number of times each cyclic component repeats in a period. One sine and one cosine term will be added for each frequency.
sigma.prior	An object created by SdPrior describing the prior distribution for the standard deviation of coefficients of the sinusoid terms.
initial.state.prior	An object created using NormalPrior , describing the prior distribution of the the initial state vector (at time 1).
sd	The standard deviation of the series to be modeled. This will be ignored if y is provided, or if all the required prior distributions are supplied directly.

Value

Returns a list with the elements necessary to specify a seasonal state model.

Author(s)

Steven L. Scott <stevescott@google.com>

References

- Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.
- Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts](#), [SdPrior](#) [MvnPrior](#)

Examples

```
data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddTrig(ss, y, period = 12, frequencies = 1)
model <- bsts(y, state.specification = ss, niter = 100)
plot(model)
```

aggregate.time.series *Aggregate a fine time series to a coarse summary*

Description

Aggregate measurements from a fine scaled time series into a coarse time series. This is similar to functions from the xts package, but it can handle aggregation from weeks to months.

Usage

```
AggregateTimeSeries(fine.series,  
                    contains.end,  
                    membership.fraction,  
                    trim.left = any(membership.fraction < 1),  
                    trim.right = NULL,  
                    byrow = TRUE)
```

Arguments

fine.series	A numeric vector or matrix giving the fine scale time series to be aggregated.
contains.end	A logical vector corresponding to fine.series indicating whether each fine time interval contains the end of a coarse time interval.
membership.fraction	A numeric vector corresponding to fine.series, giving the fraction of each time interval's observation attributable to the coarse interval containing the fine interval's first day. This will usually be a vector of 1's, unless fine.series is weekly.
trim.left	Logical indicating whether the first observation in the coarse aggregate should be removed.
trim.right	Logical indicating whether the final observation in the coarse aggregate should be removed.
byrow	Logical. If fine.series is a matrix, this argument indicates whether rows (TRUE) or columns (FALSE) correspond to time points.

Value

A matrix (if fine.series is a matrix) or vector (otherwise) containing the aggregated values of fine.series.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```

week.ending <- as.Date(c("2011-11-05",
                        "2011-11-12",
                        "2011-11-19",
                        "2011-11-26",
                        "2011-12-03",
                        "2011-12-10",
                        "2011-12-17",
                        "2011-12-24",
                        "2011-12-31"))
membership.fraction <- GetFractionOfDaysInInitialMonth(week.ending)
which.month <- MatchWeekToMonth(week.ending, as.Date("2011-11-01"))
contains.end <- WeekEndsMonth(week.ending)

weekly.values <- rnorm(length(week.ending))
monthly.values <- AggregateTimeSeries(weekly.values, contains.end, membership.fraction)

```

```
aggregate.weeks.to.months
```

Aggregate a weekly time series to monthly

Description

Aggregate measurements from a weekly time series into a monthly time series.

Usage

```

AggregateWeeksToMonths(weekly.series,
                        membership.fraction = NULL,
                        trim.left = TRUE,
                        trim.right = NULL)

```

Arguments

<code>weekly.series</code>	A numeric vector or matrix of class <code>zoo</code> giving the weekly time series to be aggregated. The index must be convertible to class <code>Date</code> .
<code>membership.fraction</code>	An optional numeric vector corresponding to <code>weekly.series</code> , giving the fraction of each week's observation attributable to the month containing the week's first day. If missing, then weeks will be split across months in proportion to the number of days in each month.
<code>trim.left</code>	Logical indicating whether the first observation in the monthly aggregate should be removed.
<code>trim.right</code>	Logical indicating whether the final observation in the monthly aggregate should be removed.

Value

A zoo-matrix (if `weekly.series` is a matrix) or vector (otherwise) containing the aggregated values of `weekly.series`.

Author(s)

Steven L. Scott <stevescott@google.com>

See Also

[AggregateTimeSeries](#)

Examples

```
week.ending <- as.Date(c("2011-11-05",
                        "2011-11-12",
                        "2011-11-19",
                        "2011-11-26",
                        "2011-12-03",
                        "2011-12-10",
                        "2011-12-17",
                        "2011-12-24",
                        "2011-12-31"))

weekly.values <- zoo(rnorm(length(week.ending)), week.ending)
monthly.values <- AggregateWeeksToMonths(weekly.values)
```

auto.ar

Sparse AR(p)

Description

Add a sparse AR(p) process to the state distribution. A sparse AR(p) is an AR(p) process with a spike and slab prior on the autoregression coefficients.

Usage

```
AddAutoAr(state.specification,
           y,
           lags = 1,
           prior = NULL,
           sdy = NULL,
           ...)
```

Arguments

state.specification	A list of state components. If omitted, an empty list is assumed.
y	A numeric vector. The time series to be modeled. This can be omitted if sdy is supplied.
lags	The maximum number of lags ("p") to be considered in the AR(p) process.
prior	An object inheriting from SpikeSlabArPrior , or NULL. If the latter, then a default SpikeSlabArPrior will be created.
sdy	The sample standard deviation of the time series to be modeled. Used to scale the prior distribution. This can be omitted if y is supplied.
...	Extra arguments passed to SpikeSlabArPrior .

Details

The model contributes $\alpha[t]$ to the expected value of $y[t]$, where the transition equation is

$$\alpha_t = \phi_1 \alpha_{i,t-1} + \dots + \phi_p \alpha_{t-p} + \epsilon_{t-1} \quad \epsilon_t \sim \mathcal{N}(0, \sigma^2)$$

The state consists of the last p lags of alpha. The state transition matrix has ϕ_i in its first row, ones along its first subdiagonal, and zeros elsewhere. The state variance matrix has σ^2 in its upper left corner and is zero elsewhere. The observation matrix has 1 in its first element and is zero otherwise.

This model differs from the one in [AddAr](#) only in that some of its coefficients may be set to zero.

Value

Returns state.specification with an AR(p) state component added to the end.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.SdPrior](#)

Examples

```

n <- 100
residual.sd <- .001

# Actual values of the AR coefficients
true.phi <- c(-.7, .3, .15)
ar <- arima.sim(model = list(ar = true.phi),
                n = n,
                sd = 3)

## Layer some noise on top of the AR process.
y <- ar + rnorm(n, 0, residual.sd)
ss <- AddAutoAr(list(), y, lags = 6)

# Fit the model with knowledge with residual.sd essentially fixed at the
# true value.
model <- bst(y, state.specification=ss, niter = 500, prior = SdPrior(residual.sd, 100000))

# Now compare the empirical ACF to the true ACF.
acf(y, lag.max = 30)
points(0:30, ARMAacf(ar = true.phi, lag.max = 30), pch = "+")
points(0:30, ARMAacf(ar = colMeans(model$AR6.coefficients), lag.max = 30))
legend("topright", leg = c("empirical", "truth", "MCMC"), pch = c(NA, "+", "o"))

```

bsts

*Bayesian structural time series***Description**

Uses MCMC to sample from the posterior distribution of a Bayesian structural time series model. This function can be used either with or without contemporaneous predictor variables (in a time series regression).

If predictor variables are present, the regression coefficients are fixed (as opposed to time varying, though time varying coefficients might be added as state component). The predictors and response in the formula are contemporaneous, so if you want lags and differences you need to put them in the predictor matrix yourself.

If no predictor variables are used, then the model is an ordinary state space time series model.

The model allows for several useful extensions beyond standard Bayesian dynamic linear models.

- A spike-and-slab prior is used for the (static) regression component of models that include predictor variables. This is especially useful with large numbers of regressor series.
- Both the spike-and-slab component (for static regressors) and the Kalman filter (for components of time series state) require observations and state variables to be Gaussian. The `bsts` package allows for non-Gaussian error families in the observation equation (as well as some state components) by using data augmentation to express these families as conditionally Gaussian.

- As of version 0.7.0, `bsts` supports having multiple observations at the same time point. In this case the basic model is taken to be

$$y_{t,j} = Z_t^T \alpha_t + \beta^T x_{t,j} + \epsilon_{t,j}.$$

This is a regression model where all observations with the same time point share a common time series effect.

Usage

```
bsts(formula,
      state.specification,
      family = c("gaussian", "logit", "poisson", "student"),
      data,
      prior,
      contrasts = NULL,
      na.action = na.pass,
      niter,
      ping = niter / 10,
      model.options = BstsOptions(),
      timestamps = NULL,
      seed = NULL,
      ### Deprecated arguments
      save.state.contributions,
      save.prediction.errors,
      bma.method,
      oda.options,
      ### End of deprecated arguments
      ...)
```

Arguments

<code>formula</code>	A formula describing the regression portion of the relationship between <code>y</code> and <code>X</code> . If no regressors are desired then the formula can be replaced by a numeric vector giving the time series to be modeled. Missing values are not allowed. If the response variable is of class <code>zoo</code> , <code>xts</code> , or <code>ts</code> , then the time series information it contains will be used in many of the plotting methods called from <code>plot.bsts</code> .
<code>state.specification</code>	A list with elements created by <code>AddLocalLinearTrend</code> , <code>AddSeasonal</code> , and similar functions for adding components of state. See the help page for <code>state.specification</code> .
<code>family</code>	The model family for the observation equation. Non-Gaussian model families use data augmentation to recover a conditionally Gaussian model.
<code>data</code>	An optional data frame, list or environment (or object coercible by <code>as.data.frame</code> to a data frame) containing the variables in the model. If not found in <code>data</code> , the variables are taken from <code>environment(formula)</code> , typically the environment from which <code>bsts</code> is called.

prior	If regressors are supplied in the model formula, then this is a prior distribution for the regression component of the model, as created by <code>SpikeSlabPrior</code> . The prior for the time series component of the model will be specified during the creation of <code>state.specification</code> . This argument is only used if a formula is specified. If the model contains no regressors, then this is simply the prior on the residual standard deviation, expressed as an object created by <code>SdPrior</code> .
contrasts	An optional list containing the names of contrast functions to use when converting factors numeric variables in a regression formula. This argument works exactly as it does in <code>lm</code> . The names of the list elements correspond to factor variables in your model formula. The list elements themselves are the names of contrast functions (see <code>help(contr.treatment)</code> and the <code>contrasts.arg</code> argument to <code>model.matrix.default</code>). This argument is only used if a model formula is specified, and even then the default is probably what you want.
na.action	What to do about missing values. The default is to allow missing responses, but no missing predictors. Set this to <code>na.omit</code> or <code>na.exclude</code> if you want to omit missing responses altogether.
niter	A positive integer giving the desired number of MCMC draws.
ping	A scalar giving the desired frequency of status messages. If <code>ping > 0</code> then the program will print a status message to the screen every <code>ping</code> MCMC iterations.
model.options	An object (list) returned by <code>BstsOptions</code> . See that function for details.
timestamps	The timestamp associated with each value of the response. This argument is primarily useful in cases where the response has missing gaps, or where there are multiple observations per time point. If the response is a "regular" time series with a single observation per time point then you can leave this argument as <code>NULL</code> . In that case, if either the response or the data argument is a type convertible to <code>zoo</code> then timestamps will be inferred.
seed	An integer to use as the random seed for the underlying C++ code. If <code>NULL</code> then the seed will be set using the clock.
save.state.contributions	***DEPRECATED*** Please use <code>BstsOptions</code> instead. Logical. If <code>TRUE</code> then a 3-way array named <code>state.contributions</code> will be stored in the returned object. The indices correspond to MCMC iteration, state model number, and time. Setting <code>save.state.contributions</code> to <code>FALSE</code> yields a smaller object, but <code>plot</code> will not be able to plot the the "state", "components", or "residuals" for the fitted model.
save.prediction.errors	***DEPRECATED*** Please use <code>BstsOptions</code> instead. Logical. If <code>TRUE</code> then a matrix named <code>one.step.prediction.errors</code> will be saved as part of the model object. The rows of the matrix represent MCMC iterations, and the columns represent time. The matrix entries are the one-step-ahead prediction errors from the Kalman filter.
bma.method	***DEPRECATED*** Please use <code>BstsOptions</code> instead. If the model contains a regression component, this argument specifies the method to use for Bayesian model averaging. "SSVS" is stochastic search variable selection, which is the classic approach from George and McCulloch (1997). "ODA" is orthogonal data augmentation, from Ghosh and Clyde (2011). It adds a set of latent

observations that make the $X^{\text{T}}X$ matrix diagonal, vastly simplifying complete data MCMC for model selection.

`oda.options` *****DEPRECATED*** Please use `BstsOptions` instead.** If `bma.method == "ODA"` then these are some options for fine tuning the ODA algorithm.

- `fallback.probability`: Each MCMC iteration will use SSVS instead of ODA with this probability. In cases where the latent data have high leverage, ODA mixing can suffer. Mixing in a few SSVS steps can help keep an errant algorithm on track.
- `eigenvalue.fudge.factor`: The latent X 's will be chosen so that the complete data $X'X$ matrix (after scaling) is a constant diagonal matrix equal to the largest eigenvalue of the observed (scaled) $X'X$ times $(1 + \text{eigenvalue.fudge.factor})$. This should be a small positive number.

`...` Extra arguments to be passed to [SpikeSlabPrior](#) (see the entry for the prior argument, above).

Details

If the model family is logit, then there are two ways one can format the response variable. If the response is 0/1, TRUE/FALSE, or 1/-1, then the response variable can be passed as with any other model family. If the response is a set of counts out of a specified number of trials then it can be passed as a two-column matrix, where the first column contains the counts of successes and the second contains the count of failures.

Likewise, if the model family is Poisson, the response can be passed as a single vector of counts, under the assumption that each observation has unit exposure. If the exposures differ across observations, then the response can be a two column matrix, with the first column containing the event counts and the second containing exposure times.

Value

An object of class `bsts` which is a list with the following components

<code>coefficients</code>	A <code>niter</code> by <code>ncol(X)</code> matrix of MCMC draws of the regression coefficients, where X is the design matrix implied by formula. This is only present if a model formula was supplied.
<code>sigma.obs</code>	A vector of length <code>niter</code> containing MCMC draws of the residual standard deviation.

The returned object will also contain named elements holding the MCMC draws of model parameters belonging to the state models. The names of each component are supplied by the entries in `state.specification`. If a model parameter is a scalar, then the list element is a vector with `niter` elements. If the parameter is a vector then the list element is a matrix with `niter` rows. If the parameter is a matrix then the list element is a 3-way array with first dimension `niter`.

Finally, if a model formula was supplied, then the returned object will contain the information necessary for the `predict` method to build the design matrix when a new prediction is made.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

Goerge and McCulloch (1997) "Approaches for Bayesian variable selection", *Statistica Sinica* pp 339–374.

Ghosh and Clyde (2011) "Rao-Blackwellization for Bayesian variable selection and model averaging in linear and binary regression: a novel data augmentation approach", *JASA* pp 1041 –1052.

See Also

[bsts](#), [AddLocalLevel](#), [AddLocalLinearTrend](#), [AddGeneralizedLocalLinearTrend](#), [AddSeasonal](#), [AddDynamicRegression](#), [SpikeSlabPrior](#), [SdPrior](#).

Examples

```
## Example 1: Time series (ts) data
data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bsts(y, state.specification = ss, niter = 500)
pred <- predict(model, horizon = 12, burn = 100)
par(mfrow = c(1,2))
plot(model)
plot(pred)

## Not run:

MakePlots <- function(model, ask = TRUE) {
  ## Make all the plots callable by plot.bsts.
  opar <- par(ask = ask)
  on.exit(par(opar))
  plot.types <- c("state", "components", "residuals",
                 "prediction.errors", "forecast.distribution")
  for (plot.type in plot.types) {
    plot(model, plot.type)
  }
  if (model$has.regression) {
    regression.plot.types <- c("coefficients", "predictors", "size")
    for (plot.type in regression.plot.types) {
      plot(model, plot.type)
    }
  }
}

## Example 2: G00G is the Google stock price, an xts series of daily
## data.
data(goog)
```

```

ss <- AddGeneralizedLocalLinearTrend(list(), goog)
model <- bsts(goog, state.specification = ss, niter = 500)
MakePlots(model)

## Example 3: Change GOOG to be zoo, and not xts.
goog <- zoo(goog, index(goog))
ss <- AddGeneralizedLocalLinearTrend(list(), goog)
model <- bsts(goog, state.specification = ss, niter = 500)
MakePlots(model)

## Example 4: Naked numeric data works too
y <- rnorm(100)
ss <- AddLocalLinearTrend(list(), y)
model <- bsts(y, state.specification = ss, niter = 500)
MakePlots(model)

## Example 5: zoo data with intra-day measurements
y <- zoo(rnorm(100),
         seq(from = as.POSIXct("2012-01-01 7:00 EST"), len = 100, by = 100))
ss <- AddLocalLinearTrend(list(), y)
model <- bsts(y, state.specification = ss, niter = 500)
MakePlots(model)

\dontrun{
## Example 6: Including regressors
data(iclaims)
ss <- AddLocalLinearTrend(list(), initial.claims$iclaimsNSA)
ss <- AddSeasonal(ss, initial.claims$iclaimsNSA, nseasons = 52)
model <- bsts(iclaimsNSA ~ ., state.specification = ss, data =
              initial.claims, niter = 1000)

plot(model)
plot(model, "components")
plot(model, "coefficients")
plot(model, "predictors")
}

## End(Not run)

## Not run:
## Example 7: Regressors with multiple time stamps.
number.of.time.points <- 50
sample.size.per.time.point <- 10
total.sample.size <- number.of.time.points * sample.size.per.time.point
sigma.level <- .1
sigma.obs <- 1

## Simulate some fake data with a local level state component.
trend <- cumsum(rnorm(number.of.time.points, 0, sigma.level))
predictors <- matrix(rnorm(total.sample.size * 2), ncol = 2)
colnames(predictors) <- c("X1", "X2")
coefficients <- c(-10, 10)
regression <- as.numeric(predictors)
y.hat <- rep(trend, sample.size.per.time.point) + regression

```

```

y <- rnorm(length(y.hat), y.hat, sigma.obs)

## Create some time stamps, with multiple observations per time stamp.
first <- as.POSIXct("2013-03-24")
dates <- seq(from = first, length = number.of.time.points, by = "month")
timestamps <- rep(dates, sample.size.per.time.point)

## Run the model with a local level trend, and an unnecessary seasonal component.
ss <- AddLocalLevel(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 7)
model <- bst(y ~ predictors, ss, niter = 250, timestamps = timestamps,
             seed = 8675309)
plot(model)
plot(model, "components")

## End(Not run)

```

bsts.options.Rd

Bsts Model Options

Description

Rarely used modeling options for bsts models.

Usage

```

BstsOptions(save.state.contributions = TRUE,
            save.prediction.errors = TRUE,
            bma.method = c("SSVS", "ODA"),
            oda.options = list(
              fallback.probability = 0.0,
              eigenvalue.fudge.factor = 0.01),
            timeout.seconds = Inf)

```

Arguments

`save.state.contributions`

Logical. If TRUE then a 3-way array named `state.contributions` will be stored in the returned object. The indices correspond to MCMC iteration, state model number, and time. Setting `save.state.contributions` to FALSE yields a smaller object, but plot will not be able to plot the the "state", "components", or "residuals" for the fitted model.

`save.prediction.errors`

Logical. If TRUE then a matrix named `one.step.prediction.errors` will be saved as part of the model object. The rows of the matrix represent MCMC iterations, and the columns represent time. The matrix entries are the one-step-ahead prediction errors from the Kalman filter.

bma.method	If the model contains a regression component, this argument specifies the method to use for Bayesian model averaging. "SSVS" is stochastic search variable selection, which is the classic approach from George and McCulloch (1997). "ODA" is orthogonal data augmentation, from Ghosh and Clyde (2011). It adds a set of latent observations that make the $X^T X$ matrix diagonal, vastly simplifying complete data MCMC for model selection.
oda.options	If bma.method == "ODA" then these are some options for fine tuning the ODA algorithm. <ul style="list-style-type: none"> • fallback.probability: Each MCMC iteration will use SSVS instead of ODA with this probability. In cases where the latent data have high leverage, ODA mixing can suffer. Mixing in a few SSVS steps can help keep an errant algorithm on track. • eigenvalue.fudge.factor: The latent X's will be chosen so that the complete data $X^T X$ matrix (after scaling) is a constant diagonal matrix equal to the largest eigenvalue of the observed (scaled) $X^T X$ times (1 + eigenvalue.fudge.factor). This should be a small positive number.
timeout.seconds	The number of seconds that sampler will be allowed to run. If the timeout is exceeded the returned object will be truncated to the final draw that took place before the timeout occurred, as if that had been the requested number of iterations.

Value

The arguments are checked to make sure they have legal types and values, then a list is returned containing the arguments.

Author(s)

Steven L. Scott <stevescott@google.com>

compare.bsts.models *Compare bsts models*

Description

Produce a set of line plots showing the cumulative absolute one step ahead prediction errors for different models. This plot not only shows which model is doing the best job predicting the data, it highlights regions of the data where the predictions are particularly good or bad.

Usage

```
CompareBstsModels(model.list,
                  burn = SuggestBurn(.1, model.list[[1]]),
                  filename = "",
                  colors = NULL,
```

```

lwd = 2,
xlab = "Time",
main = "",
grid = TRUE)

```

Arguments

model.list	A list of <code>bsts</code> models.
burn	The number of initial MCMC iterations to remove from each model as burn-in.
filename	A string. If non-empty string then a pdf of the plot will be saved in the specified file.
colors	A vector of colors to use for the different lines in the plot. If NULL then the <code>rainbow</code> palette will be used.
lwd	The width of the lines to be drawn.
xlab	Label for the horizontal axis.
main	Main title for the plot.
grid	Logical. Should gridlines be drawn in the background?

Value

Invisibly returns the matrix of cumulative one-step ahead prediction errors (the lines in the top panel of the plot). Each row in the matrix corresponds to a model in `model.list`.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```

data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
trend.only <- bsts(y, ss, niter = 500)

ss <- AddSeasonal(ss, y, nseasons = 12)
trend.and.seasonal <- bsts(y, ss, niter = 500)

CompareBstsModels(list(trend = trend.only,
                       "trend and seasonal" = trend.and.seasonal))

```

estimate.time.scale *Intervals between dates*

Description

Estimate the time scale used in time series data.

Usage

```
EstimateTimeScale(dates)
```

Arguments

dates A sorted vector of class `Date`.

Value

A character string. Either "daily", "weekly", "yearly", "monthly", "quarterly", or "other". The value is determined based on counting the number of days between successive observations in dates.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```
weekly.data <- as.Date(c("2011-10-01",
                        "2011-10-08",
                        "2011-10-15",
                        "2011-10-22",
                        "2011-10-29",
                        "2011-11-05"))

EstimateTimeScale(weekly.data) # "weekly"

almost.weekly.data <- as.Date(c("2011-10-01",
                                "2011-10-08",
                                "2011-10-15",
                                "2011-10-22",
                                "2011-10-29",
                                "2011-11-06")) # last day is one later

EstimateTimeScale(weekly.data) # "other"
```

extend.time	<i>Extends a vector of dates to a given length</i>
-------------	--

Description

Pads a vector of dates to a specified length.

Usage

```
ExtendTime(dates, number.of.periods, dt = NULL)
```

Arguments

dates	An ordered vector of class Date .
number.of.periods	The desired length of the output.
dt	A character string describing the frequency of the dates in dates. Possible values are "daily", "weekly", "monthly", "quarterly", "yearly", or "other". An attempt to deduce dt will be made if it is missing.

Value

If `number.of.periods` is longer than `length(dates)`, then dates will be padded to the desired length. Extra dates are added at time intervals matching the average interval in dates. Thus they may not be

Author(s)

Steven L. Scott <stevescott@google.com>

See Also

[bsts.mixed](#).

Examples

```
origin.month <- as.Date("2011-09-01")
week.ending <- as.Date(c("2011-10-01", ## 1
                        "2011-10-08", ## 2
                        "2011-12-03", ## 3
                        "2011-12-31")) ## 4
MatchWeekToMonth(week.ending, origin.month) == 1:4
```

Description

Tools for checking if a series of timestamps is 'regular' meaning that it has no duplicates, and no gaps. Checking for regularity can be tricky. For example, if you have monthly observations with [Date](#) or [POSIXt](#) timestamps then gaps between timestamps can be 28, 29, 30, or 31 days, but the series is still "regular".

Usage

```
NoDuplicates(timestamps)
NoGaps(timestamps)
IsRegular(timestamps)

HasDuplicateTimestamps(bsts.object)
```

Arguments

`timestamps` A set of (possibly irregular or non-unique) timestamps. This could be a set of integers (like 1, 2, , 3...), a set of numeric like (1945, 1945.083, 1945.167, ...) indicating years and fractions of years, a [Date](#) object, or a [POSIXt](#) object.

`bsts.object` A `bsts` model object.

Value

All four functions return scalar logical values. `NoDuplicates` returns `TRUE` if all elements of `timestamps` are unique.

`NoGaps` examines the smallest nonzero gap between time points. As long as no gaps between time points are more than twice as wide as the smallest gap, it returns `TRUE`, indicating that there are no missing timestamps. Otherwise it returns `FALSE`.

`IsRegular` returns `TRUE` if `NoDuplicates` and `NoGaps` both return `TRUE`.

`HasDuplicateTimestamps` returns `FALSE` if the data used to fit `bsts.model` either has `NULL` timestamps, or if the timestamps contain no duplicate values.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```
first <- as.POSIXct("2015-04-19 08:00:04")
monthly <- seq(from = first, length.out = 24, by = "month")
IsRegular(monthly) ## TRUE

skip.one <- monthly[-8]
```

```
IsRegular(skip.one) ## FALSE

has.duplicates <- monthly
has.duplicates[1] <- has.duplicates[2]
IsRegular(has.duplicates) ## FALSE
```

geometric.sequence *Create a Geometric Sequence*

Description

Create a geometric sequence.

Usage

```
GeometricSequence(length, initial.value = 1, discount.factor = .5)
```

Arguments

`length` A positive integer giving the length of the desired sequence.
`initial.value` The first term in the sequence. Cannot be zero.
`discount.factor` The ratio between a sequence term and the preceding term. Cannot be zero.

Value

A numeric vector containing the desired sequence.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```
GeometricSequence(4, .8, .6)
# [1] 0.8000 0.4800 0.2880 0.1728

GeometricSequence(5, 2, 3)
# [1] 2 6 18 54 162

## Not run:
GeometricSequence(0, -1, -2)
# Error: length > 0 is not TRUE

## End(Not run)
```

get.fraction	<i>Compute membership fractions</i>
--------------	-------------------------------------

Description

Returns the fraction of days in a week that occur in the ear

Usage

```
GetFractionOfDaysInInitialMonth(week.ending)
GetFractionOfDaysInInitialQuarter(week.ending)
```

Arguments

week.ending A vector of class [Date](#). Each entry contains the date of the last day in a week.

Value

Returns a numeric vector of the same length as `week.ending`. Each entry gives the fraction of days in the week that occur in the coarse time interval (month or quarter) containing the start of the week (i.e the date 6 days before).

Author(s)

Steven L. Scott <stevescott@google.com>

See Also

[bsts.mixed](#).

Examples

```
dates <- as.Date(c("2003-03-31",
                  "2003-04-01",
                  "2003-04-02",
                  "2003-04-03",
                  "2003-04-04",
                  "2003-04-05",
                  "2003-04-06",
                  "2003-04-07"))
fraction <- GetFractionOfDaysInInitialMonth(dates)
fraction == c(1, 6/7, 5/7, 4/7, 3/7, 2/7, 1/7, 1)
```

goog	<i>Google stock price</i>
------	---------------------------

Description

Daily closing price of Google stock.

Usage

```
data(goog)
```

Format

xts time series

Source

The Internets

HarveyCumulator	<i>HarveyCumulator</i>
-----------------	------------------------

Description

Given a state space model on a fine scale, the Harvey cumulator aggregates the model to a coarser scale (e.g. from days to weeks, or weeks to months).

Usage

```
HarveyCumulator(fine.series,
                 contains.end,
                 membership.fraction)
```

Arguments

<code>fine.series</code>	The fine-scale time series to be aggregated.
<code>contains.end</code>	A logical vector, with length matching <code>fine.series</code> indicating whether each fine scale time interval contains the end of a coarse time interval. For example, months don't contain a fixed number of weeks, so when cumulating a weekly time series into a monthly series, you need to know which weeks contain the end of a month.
<code>membership.fraction</code>	The fraction of each fine-scale time observation belonging to the coarse scale time observation at the beginning of the time interval. For example, if week <code>i</code> started in March and ended in April, <code>membership.fraction[i]</code> is the fraction of <code>fine.series[i]</code> that should be attributed to March. This should be 1 for most observations.

Value

Returns a vector containing the course scale partial aggregates of `fine.series`.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.mixed](#),

Examples

```
data(goog)
days <- factor(weekdays(index(goog)),
               levels = c("Monday", "Tuesday", "Wednesday",
                          "Thursday", "Friday"),
               ordered = TRUE)

## Because of holidays, etc the days do not always go in sequence.
## (Sorry, Rebecca Black! https://www.youtube.com/watch?v=kfVsf0SbJY0)
## diff.days[i] is the number of days between days[i-1] and days[i].
## We know that days[i] is the end of a week if diff.days[i] < 0.
diff.days <- tail(as.numeric(days), -1) - head(as.numeric(days), -1)
contains.end <- c(FALSE, diff.days < 0)

goog.weekly <- HarveyCumulator(goog, contains.end, 1)
```

iclaims

Initial Claims Data

Description

A weekly time series of US initial claims for unemployment. The first column contains the initial claims numbers from FRED. The others contain a measure of the relative popularity of various search queries identified by Google Correlate.

Usage

```
data(iclaims)
```

Format

zoo time series

Source

FRED. <http://research.stlouisfed.org/fred2/series/ICNSA>,
Google correlate. <http://www.google.com/trends/correlate>

See Also

[bsts](#)

Examples

```
data(iclaims)  
plot(initial.claims)
```

<code>last.day.in.month</code>	<i>Find the last day in a month</i>
--------------------------------	-------------------------------------

Description

Finds the last day in the month containing a specified date.

Usage

```
LastDayInMonth(dates)
```

Arguments

`dates` A vector of class [Date](#).

Value

A vector of class [Date](#) where each entry is the last day in the month containing the corresponding entry in `dates`.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```

inputs <- as.Date(c("2007-01-01",
                  "2007-01-31",
                  "2008-02-01",
                  "2008-02-29",
                  "2008-03-14",
                  "2008-12-01",
                  "2008-12-31"))
expected.outputs <- as.Date(c("2007-01-31",
                             "2007-01-31",
                             "2008-02-29",
                             "2008-02-29",
                             "2008-03-31",
                             "2008-12-31",
                             "2008-12-31"))
LastDayInMonth(inputs) == expected.outputs

```

MATCH.NumericTimestamps

Match Numeric Timestamps

Description

S3 generic method for MATCH function supplied in the zoo package.

Usage

```

## S3 method for class 'NumericTimestamps'
MATCH(x, table, nomatch = NA, ...)

```

Arguments

x	A numeric set of timestamps.
table	A set of regular numeric timestamps to match against.
nomatch	The value to be returned in the case when no match is found. Note that it is coerced to integer.
...	Additional arguments passed to match .

Details

Numeric timestamps match if they agree to 8 significant digits.

Value

Returns the index of the entry in table matched by each argument in x. If an entry has no match then nomatch is returned at that position.

See Also[MATCH](#)

match.week.to.month *Find the month containing a week*

Description

Returns the index of a month, in a sequence of months, that contains a given week.

Usage

```
MatchWeekToMonth(week.ending, origin.month)
```

Arguments

`week.ending` A vector of class [Date](#). Each entry contains the date of the last day in a week.
`origin.month` A [Date](#), giving any day in the month to use as the origin of the sequence (month 1).

Value

The index of the month matching the month containing the first day in `week.ending`. The origin is month 1. It is the caller's responsibility to ensure that these indices correspond to legal values in a particular vector of months.

Author(s)

Steven L. Scott <stevescott@google.com>

See Also

[bsts.mixed](#).

Examples

```
origin.month <- as.Date("2011-09-01")
week.ending <- as.Date(c("2011-10-01", ## 1
                        "2011-10-08", ## 2
                        "2011-12-03", ## 3
                        "2011-12-31")) ## 4
MatchWeekToMonth(week.ending, origin.month) == 1:4
```

mixed.frequency *Models for mixed frequency time series*

Description

Fit a structured time series to mixed frequency data.

Usage

```
bsts.mixed(target.series,
           predictors,
           which.coarse.interval,
           membership.fraction,
           contains.end,
           state.specification,
           regression.prior,
           niter,
           ping = niter / 10,
           seed = NULL,
           truth = NULL,
           ...)
```

Arguments

- `target.series` A vector object of class `zoo` indexed by calendar dates. The date associated with each element is the LAST DAY in the time interval measured by the corresponding value. The value is what Harvey (1989) calls a 'flow' variable. It is a number that can be viewed as an accumulation over the measured time interval.
- `predictors` A matrix of class `zoo` indexed by calendar dates. The date associated with each row is the LAST DAY in the time interval encompassing the measurement. The dates are expected to be at a finer scale than the dates in `target.series`. Any predictors should be at sufficient lags to be able to predict the rest of the cycle.
- `which.coarse.interval` A numeric vector of length `nrow(predictors)` giving the index of the coarse interval corresponding to the end of each fine interval.
- `membership.fraction` A numeric vector of length `nrow(predictors)` giving the fraction of activity attributed to the coarse interval corresponding to the beginning of each fine interval. This is always positive, and will be 1 except when a fine interval spans the boundary between two coarse intervals.
- `contains.end` A logical vector of length `nrow(predictors)` indicating whether each fine interval contains the end of a coarse interval.
- `state.specification` A state specification like that required by `bsts`.

regression.prior	A prior distribution created by <code>SpikeSlabPrior</code> . A default prior will be generated if none is specified.
niter	The desired number of MCMC iterations.
ping	An integer indicating the frequency with which progress reports get printed. E.g. setting <code>ping = 100</code> will print a status message with a time and iteration stamp every 100 iterations. If you don't want these messages set <code>ping < 0</code> .
seed	An integer to use as the random seed for the underlying C++ code. If NULL then the seed will be set using the clock.
truth	For debugging purposes only. A list containing one or more of the following elements. If any are present then corresponding values will be held fixed in the MCMC algorithm. <ul style="list-style-type: none"> • A matrix named <code>state</code> containing the state of the coarse model from a fake-data simulation. • A vector named <code>beta</code> of regression coefficients. • A scalar named <code>sigma.obs</code>.
...	Extra arguments passed to <code>SpikeSlabPrior</code>

Value

An object of class `bsts.mixed`, which is a list with the following elements. Many of these are arrays, in which case the first index of the array corresponds to the MCMC iteration number.

<code>coefficients</code>	A matrix containing the MCMC draws of the regression coefficients. Rows correspond to MCMC draws, and columns correspond to variables.
<code>sigma.obs</code>	The standard deviation of the weekly latent observations.
<code>state.contributions</code>	A three-dimensional array containing the MCMC draws of each state model's contributions to the state of the weekly model. The three dimensions are MCMC iteration, state model, and week number.
<code>weekly</code>	A matrix of MCMC draws of the weekly latent observations. Rows are MCMC iterations, and columns are weekly time points.
<code>cumulator</code>	A matrix of MCMC draws of the cumulator variable.

The returned object also contains MCMC draws for the parameters of the state models supplied as part of `state.specification`, relevant information passed to the function call, and other supplemental information.

Author(s)

Steven L. Scott <stevescott@google.com>

References

- Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.
- Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts](#), [AddLocalLevel](#), [AddLocalLinearTrend](#), [AddGeneralizedLocalLinearTrend](#), [SpikeSlabPrior](#), [SdPrior](#).

Examples

```
data <- SimulateFakeMixedFrequencyData(nweeks = 104, xdim = 20)

## Setting an upper limit on the standard deviations can help keep the
## MCMC from flying off to infinity.
sd.limit <- sd(data$coarse.target)
state.specification <-
  AddLocalLinearTrend(list(),
    data$coarse.target,
    level.sigma.prior = SdPrior(1.0, 5, upper.limit = sd.limit),
    slope.sigma.prior = SdPrior(.5, 5, upper.limit = sd.limit))
weeks <- index(data$predictor)
months <- index(data$coarse.target)
which.month <- MatchWeekToMonth(weeks, months[1])
membership.fraction <- GetFractionOfDaysInInitialMonth(weeks)
contains.end <- WeekEndsMonth(weeks)

model <- bsts.mixed(target.series = data$coarse.target,
  predictors = data$predictors,
  membership.fraction = membership.fraction,
  contains.end = contains.end,
  which.coarse = which.month,
  state.specification = state.specification,
  niter = 500,
  expected.r2 = .999,
  prior.df = 1)

plot(model, "state")
plot(model, "components")
```

month.distance	<i>Elapsed time in months</i>
----------------	-------------------------------

Description

The (integer) number of months between dates.

Usage

```
MonthDistance(dates, origin)
```

Arguments

dates A vector of class `Date` to be measured.
origin A scalar of class `Date`.

Value

Returns a numeric vector giving the integer number of months that have elapsed between `origin` and each element in `dates`. The daily component of each date is ignored, so two dates that are in the same month will have the same measured distance. Distances are signed, so months that occur before `origin` will have negative values.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```
dates <- as.Date(c("2008-04-17",
                  "2008-05-01",
                  "2008-05-31",
                  "2008-06-01"))
origin <- as.Date("2008-05-15")
MonthDistance(dates, origin) == c(-1, 0, 0, 1)
```

new.home.sales

New home sales and Google trends

Description

The first column, HSN1FNSA is a time series of new home sales in the US, obtained from the FRED online data base. The series has been manually deseasonalized. The remaining columns contain search terms from Google trends (obtained from <http://trends.google.com/correlate>). These show the relative popularity of each search term among all search terms typed into Google. All series in this data set have been standardized by subtracting off their mean and dividing by their standard deviation.

Usage

```
data(new.home.sales)
```

Format

zoo time series

Source

FRED and trends.google.com

`one.step.prediction.errors`*Prediction Errors*

Description

Computes the one-step-ahead prediction errors for a `bsts` model.

Usage

```
bsts.prediction.errors(bsts.object,  
                       cutpoints = NULL,  
                       burn = SuggestBurn(.1, bsts.object))
```

Arguments

<code>bsts.object</code>	An object of class <code>bsts</code> .
<code>cutpoints</code>	An increasing sequence of integers between 1 and the number of time points in the training data for <code>bsts.object</code> , or <code>NULL</code> . If <code>NULL</code> then the in-sample one-step prediction errors from the <code>bsts</code> object will be extracted and returned. Otherwise the model will be re-fit with a separate MCMC run for each entry in 'cutpoints'. Data up to each cutpoint will be included in the fit, and one-step prediction errors for data after the cutpoint will be computed.
<code>burn</code>	An integer giving the number of MCMC iterations to discard as burn-in. If <code>burn <= 0</code> then no burn-in sample will be discarded.

Details

Returns the posterior distribution of the one-step-ahead prediction errors from the `bsts.object`. The errors are computed using the Kalman filter, and are of two types.

Purely in-sample errors are computed as a by-product of the Kalman filter as a result of fitting the model. These are stored in the `bsts.object` assuming the `save.prediction.errors` option is `TRUE`, which is the default (See [BstsOptions](#)). The in-sample errors are 'in-sample' in the sense that the parameter values used to run the Kalman filter are drawn from their posterior distribution given complete data. Conditional on the parameters in that MCMC iteration, each 'error' is the difference between the observed $y[t]$ and its expectation given data to $t-1$.

Purely out-of-sample errors can be computed by specifying the 'cutpoints' argument. If cutpoints are supplied then a separate MCMC is run using just data up to the cutpoint. The Kalman filter is then run on the remaining data, again finding the difference between $y[t]$ and its expectation given data to $t-1$, but conditional on parameters estimated using data up to the cutpoint.

Value

A matrix of draws of the one-step-ahead prediction errors. Rows of the matrix correspond to MCMC draws. Columns correspond to time.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts](#), [AddLocalLevel](#), [AddLocalLinearTrend](#), [AddGeneralizedLocalLinearTrend](#), [SpikeSlabPrior](#), [SdPrior](#).

Examples

```
data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bsts(y, state.specification = ss, niter = 500)
errors <- bsts.prediction.errors(model, burn = 100)
PlotDynamicDistribution(errors$in.sample)

## Compute out of sample prediction errors beyond times 80 and 120.
errors <- bsts.prediction.errors(model, cutpoints = c(80, 120))
plot(model, "prediction.errors", cutpoints = c(80, 120))
str(errors)      ## three matrices, with 400 (= 500 - 100) rows
                 ## and length(y) columns
```

plot.bsts

Plotting functions for Bayesian structural time series

Description

Functions to plot the results of a model fit using [bsts](#).

Usage

```
## S3 method for class 'bsts'
plot(x, y = c("state", "components", "residuals",
             "coefficients", "prediction.errors",
             "forecast.distribution",
             "predictors", "size", "dynamic", "seasonal", "help"),
     ...)

PlotBstsCoefficients(bsts.object, burn = SuggestBurn(.1, bsts.object),
```



```

        inclusion.threshold = 0, number.of.variables = NULL, ...)
PlotBstsComponents(bsts.object, burn = SuggestBurn(.1, bsts.object),
  time, same.scale = TRUE,
  layout = c("square", "horizontal", "vertical"),
  style = c("dynamic", "boxplot"),
  ylim = NULL, ...)
PlotDynamicRegression(bsts.object, burn = SuggestBurn(.1, bsts.object),
  time = NULL, style = c("dynamic", "boxplot"),
  layout = c("square", "horizontal", "vertical"),
  ...)
PlotBstsState(bsts.object, burn = SuggestBurn(.1, bsts.object),
  time, show.actuals = TRUE,
  style = c("dynamic", "boxplot"),
  scale = c("linear", "mean"),
  ylim = NULL,
  ...)
PlotBstsResiduals(bsts.object, burn = SuggestBurn(.1, bsts.object),
  time, style = c("dynamic", "boxplot"), ...)

PlotBstsPredictionErrors(bsts.object, cutpoints = NULL,
  burn = SuggestBurn(.1, bsts.object),
  style = c("dynamic", "boxplot"),
  xlab = "Time", ylab = "", main = "",
  ...)

PlotBstsForecastDistribution(bsts.object, cutpoints = NULL,
  burn = SuggestBurn(.1, bsts.object),
  style = c("dynamic", "boxplot"),
  xlab = "Time",
  ylab = "",
  main = "",
  show.actuals = TRUE,
  col.actuals = "blue",
  ...)

PlotBstsSize(bsts.object, burn = SuggestBurn(.1, bsts.object), style =
  c("histogram", "ts"), ...)
PlotSeasonalEffect(bsts.object, nseasons = 7, season.duration = 1,
  same.scale = TRUE, ylim = NULL, get.season.name = NULL,
  burn = SuggestBurn(.1, bsts.object), ...)

```

Arguments

x	An object of class <code>bsts</code> .
bsts.object	An object of class <code>bsts</code> .
y	A character string indicating the aspect of the model that should be plotted.
burn	The number of MCMC iterations to discard as burn-in.
col.actuals	The color to use for the actual data when comparing actuals vs forecasts.

cutpoints	A numeric vector of integers, or NULL. For diagnostic plots of prediction errors or forecast distributions, the model will be re-fit with a separate MCMC run for each entry in 'cutpoints'. Data up to each cutpoint will be included in the fit, and one-step prediction errors for data after the cutpoint will be computed.
get.season.name	A function that can be used to infer the title of each seasonal plot. It should take a single <code>POSIXt</code> , <code>Date</code> , or similar object as an argument, and return a single string that can be used as a panel title. If <code>get.season.name</code> is NULL and <code>nseasons</code> is specified or inferred to be one of the following values, then the following functions will be used. <ul style="list-style-type: none"> • 4: <code>quarters</code> • 7: <code>weekdays</code> • 12: <code>months</code>
inclusion.threshold	An inclusion probability that individual coefficients must exceed in order to be displayed when what <code>==</code> "coefficients". See the help file for <code>plot.lm.spike</code> .
layout	For controlling the layout of functions that generate multiple plots.
main	Main title for the plot.
nseasons	If there is only one seasonal component in the model, this argument is ignored. If there are multiple seasonal components then <code>nseasons</code> and <code>season.duration</code> are used to select the desired one.
number.of.variables	If non-NULL this specifies the number of coefficients to plot, taking precedence over <code>inclusion.threshold</code> . See <code>plot.lm.spike</code> .
same.scale	Logical. If TRUE then all the state components will be plotted with the same scale on the vertical axis. If FALSE then each component will get its own scale for the vertical axis.
scale	The scale on which to plot the state. If the error family is "logit" or "poisson" then the state can either be plotted on the scale of the linear predictor (e.g. trend + seasonal + regression) or the linear predictor can be passed through the link function so as to plot the distribution of the conditional mean.
season.duration	If there is only one seasonal component in the model, this argument is ignored. If there are multiple seasonal components then <code>nseasons</code> and <code>season.duration</code> are used to select the desired one.
show.actuals	Logical. If TRUE then actual values from the fitted series will be shown on the plot.
style	The desired plot style. Partial matching is allowed, so "dyn" would match "dynamic", for example.
time	An optional vector of values to plot against. If missing, the default is to diagnose the time scale of the original time series.
xlab	Label for the horizontal axis.
ylab	Label for the vertical axis.

ylim	Limits for the vertical axis. If NULL these will be inferred from the state components and the same .scale argument. Otherwise all plots will be created with the same ylim values.
...	Additional arguments to be passed to PlotDynamicDistribution , or TimeSeriesBoxplot .

Details

[PlotBstsState](#), [PlotBstsComponents](#), and [PlotBstsResiduals](#) all produce dynamic distribution plots. [PlotBstsState](#) plots the aggregate state contribution (including regression effects) to the mean, while [PlotBstsComponents](#) plots the contribution of each state component. [PlotBstsResiduals](#) plots the posterior distribution of the residuals given complete data (i.e. looking forward and backward in time). [PlotBstsPredictionErrors](#) plots filtering errors (i.e. the one-step-ahead prediction errors given data up to the previous time point). [PlotBstsForecastDistribution](#) plots the one-step-ahead forecasts instead of the prediction errors.

[PlotBstsCoefficients](#) creates a significance plot for the predictors used in the state space regression model. It is obviously not useful for models with no regressors.

[PlotBstsSize](#) plots the distribution of the number of predictors included in the model.

[PlotSeasonalEffect](#) generates an array of plots showing how the distribution of the seasonal effect changes, for each season.

Value

These functions are called for their side effect, which is to produce a plot on the current graphics device.

[PlotBstsState](#) invisibly returns the state object being plotted.

See Also

[bsts](#) [PlotDynamicDistribution](#) [plot.lm.spike](#)

Examples

```
data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bst(y, state.specification = ss, niter = 500)
plot(model, burn = 100)
plot(model, "residuals", burn = 100)
plot(model, "components", burn = 100)
plot(model, "forecast.distribution", burn = 100)
```

plot.bsts.mixed *Plotting functions for mixed frequency Bayesian structural time series*

Description

Functions for plotting the output of a mixed frequency time series regression.

Usage

```
## S3 method for class 'bsts.mixed'
plot(x,
      y = c("state", "components",
            "coefficients", "predictors", "size"),
      ...)

PlotBstsMixedState(bsts.mixed.object,
                   burn = SuggestBurn(.1, bsts.mixed.object),
                   time = NULL,
                   fine.scale = FALSE,
                   style = c("dynamic", "boxplot"),
                   trim.left = NULL,
                   trim.right = NULL,
                   ...)

PlotBstsMixedComponents(bsts.mixed.object,
                        burn = SuggestBurn(.1, bsts.mixed.object),
                        time = NULL,
                        same.scale = TRUE,
                        fine.scale = FALSE,
                        style = c("dynamic", "boxplot"),
                        layout = c("square", "horizontal", "vertical"),
                        ylim = NULL,
                        trim.left = NULL,
                        trim.right = NULL,
                        ...)
```

Arguments

x	An object of class <code>bsts.mixed</code> .
bsts.mixed.object	An object of class <code>bsts.mixed</code> .
y	A character string indicating the aspect of the model that should be plotted.
burn	The number of MCMC iterations to discard as burn-in.
time	An optional vector of values to plot against. If missing, the default is to obtain the time scale from the original time series.

fine.scale	Logical. If TRUE then the plots will be at the weekly level of granularity. If FALSE they will be at the monthly level.
same.scale	Logical. If TRUE then all the state components will be plotted with the same scale on the vertical axis. If FALSE then each component will get its own scale for the vertical axis.
style	character. If "dynamic" then a dynamic distribution plot will be shown. If "box" then boxplots will be shown.
layout	A character string indicating whether the plots showing components of state should be laid out in a square, horizontally, or vertically.
trim.left	A logical indicating whether the first (presumably partial) observation in the aggregated state time series should be removed.
trim.right	A logical indicating whether the last (presumably partial) observation in the aggregated state time series should be removed.
ylim	Limits for the vertical axis. Optional.
...	Additional arguments to be passed to PlotDynamicDistribution or TimeSeriesBoxplot

Details

[PlotBstsMixedState](#) plots the aggregate state contribution (including regression effects) to the mean, while [PlotBstsComponents](#) plots the contribution of each state component separately. [PlotBstsCoefficients](#) creates a significance plot for the predictors used in the state space regression model.

Value

These functions are called for their side effect, which is to produce a plot on the current graphics device.

See Also

[bsts.mixed](#) [PlotDynamicDistribution](#) [plot.lm.spike](#) [PlotBstsSize](#)

Examples

```
## Not run:
## This example is flaky and needs to be fixed
data <- SimulateFakeMixedFrequencyData(nweeks = 104, xdim = 20)
state.specification <- AddLocalLinearTrend(list(), data$coarse.target)
weeks <- index(data$predictor)
months <- index(data$coarse.target)
which.month <- MatchWeekToMonth(weeks, months[1])
membership.fraction <- GetFractionOfDaysInInitialMonth(weeks)
contains.end <- WeekEndsMonth(weeks)

model <- bsts.mixed(target.series = data$coarse.target,
  predictors = data$predictors,
  membership.fraction = membership.fraction,
  contains.end = contains.end,
  which.coarse = which.month,
```

```

state.specification = state.specification,
niter = 500)

plot(model, "state")
plot(model, "components")

## End(Not run)

```

plot.bsts.prediction *Plot predictions from Bayesian structural time series*

Description

Plot the posterior predictive distribution from a `bsts` prediction object.

Usage

```

## S3 method for class 'bsts.prediction'
plot(x,
      y = NULL,
      burn = 0,
      plot.original = TRUE,
      median.color = "blue",
      median.type = 1,
      median.width = 3,
      interval.quantiles = c(.025, .975),
      interval.color = "green",
      interval.type = 2,
      interval.width = 2,
      style = c("dynamic", "boxplot"),
      ylim = NULL,
      ...)

```

Arguments

<code>x</code>	An object of class <code>bsts.prediction</code> created by calling <code>predict</code> on a <code>bsts</code> object.
<code>y</code>	A dummy argument necessary to match the signature of the <code>plot</code> generic function. This argument is unused.
<code>plot.original</code>	Logical or numeric. If <code>TRUE</code> then the prediction is plotted after a time series plot of the original series. If <code>FALSE</code> , the prediction fills the entire plot. If numeric, then it specifies the number of trailing observations of the original time series to plot in addition to the predictions.
<code>burn</code>	The number of observations you wish to discard as burn-in from the posterior predictive distribution. This is in addition to the burn-in discarded using <code>predict.bsts</code> .
<code>median.color</code>	The color to use for the posterior median of the prediction.

median.type	The type of line (lty) to use for the posterior median of the prediction.
median.width	The width of line (lwd) to use for the posterior median of the prediction.
interval.quantiles	The lower and upper limits of the credible interval to be plotted.
interval.color	The color to use for the upper and lower limits of the 95% credible interval for the prediction.
interval.type	The type of line (lty) to use for the upper and lower limits of the 95% credible interval for of the prediction.
interval.width	The width of line (lwd) to use for the upper and lower limits of the 95% credible interval for of the prediction.
style	Either "dynamic", for dynamic distribution plots, or "boxplot", for box plots. Partial matching is allowed, so "dyn" or "box" would work, for example.
ylim	Limits on the vertical axis.
...	Extra arguments to be passed to PlotDynamicDistribution and lines .

Details

Plots the posterior predictive distribution described by x using a dynamic distribution plot generated by [PlotDynamicDistribution](#). Overlays the posterior median and 95% prediction limits for the predictive distribution.

Value

Returns NULL.

See Also

[bsts](#) [PlotDynamicDistribution](#) [plot.lm.spike](#)

Examples

```
data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bst(y, state.specification = ss, niter = 500)
pred <- predict(model, horizon = 12, burn = 100)
plot(pred)
```

plot.bsts.predictors *Plot the most likely predictors*

Description

Creates a time series plot showing the most likely predictors of a time series used to fit a [bsts](#) object.

Usage

```
PlotBstsPredictors(bsts.object,  
                   burn = SuggestBurn(.1, bsts.object),  
                   inclusion.threshold = .1,  
                   ylim = NULL,  
                   flip.signs = TRUE,  
                   show.legend = TRUE,  
                   grayscale = TRUE,  
                   short.names = TRUE,  
                   ...)
```

Arguments

bsts.object	An object of class bsts .
burn	The number of observations you wish to discard as burn-in.
inclusion.threshold	Plot predictors with marginal inclusion probabilities above this threshold.
ylim	Scale for the vertical axis.
flip.signs	If true then a predictor with a negative sign will be flipped before being plotted, to better align visually with the target series.
show.legend	Should a legend be shown indicating which predictors are plotted?
grayscale	Logical. If TRUE then lines for different predictors grow progressively lighter as their inclusion probability decreases. If FALSE then lines are drawn in black.
short.names	Logical. If TRUE then a common prefix or suffix shared by all the variables will be discarded.
...	Extra arguments to be passed to plot .

See Also

[bsts](#) [PlotDynamicDistribution](#) [plot.lm.spike](#)

Examples

```

data(AirPassengers)
y <- log(AirPassengers)
lag.y <- c(NA, head(y, -1))
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
## Call bsts with na.action = na.omit to omit the leading NA in lag.y
model <- bst(y ~ lag.y, state.specification = ss, niter = 500,
             na.action = na.omit)
plot(model, "predictors")

```

plot.holidays

Plot bsts holidays

Description

Makes a series of side-by-side boxplots for all the holiday state components in model.

Usage

```
PlotHolidays(model, ylim = NULL, same.scale = TRUE, ...)
```

Arguments

model	A model fit by bsts containing one or more holiday state components.
ylim	Limits on the vertical axis of the plots. If <code>ylim</code> is specified, all plots will have the same vertical axis.
same.scale	If <code>ylim</code> is <code>NULL</code> , this flag determines whether all plots share the same scale for their vertical axis (<code>same.scale == TRUE</code>), or each plot is independently scaled (<code>same.scale == FALSE</code>).
...	Extra arguments passed to boxplot .

Value

Returns `invisible{NULL}`.

See Also

[bsts](#) [AddNamedHolidays](#) [AddFixedDateHoliday](#) [AddNthWeekdayInMonthHoliday](#)

Examples

```
## TODO(stevescott): add examples
```

predict.bsts *Prediction for bayesian structural time series*

Description

Generated draws from the posterior predictive distribution of a `bsts` object.

Usage

```
## S3 method for class 'bsts'
predict(object,
        newdata = NULL,
        horizon = 1,
        burn = SuggestBurn(.1, object),
        na.action = na.exclude,
        olddata = NULL,
        trials.or.exposure = 1,
        quantiles = c(.025, .975),
        ...)
```

Arguments

<code>object</code>	An object of class <code>bsts</code> created by a call to the function <code>bsts</code> .
<code>newdata</code>	a vector, matrix, or data frame containing the predictor variables to use in making the prediction. This is only required if <code>object</code> contains a regression component. If a data frame, it must include variables with the same names as the data used to fit <code>object</code> . The first observation in <code>newdata</code> is assumed to be one time unit after the end of the last observation used in fitting <code>object</code> , and the subsequent observations are sequential time points. If the regression part of <code>object</code> contains only a single predictor then <code>newdata</code> can be a vector. If <code>newdata</code> is passed as a matrix it is the caller's responsibility to ensure that it contains the correct number of columns and that the columns correspond to those in <code>object\$coefficients</code> .
<code>horizon</code>	An integer specifying the number of periods into the future you wish to predict. If <code>object</code> contains a regression component then the forecast horizon is <code>nrow(X)</code> , and this argument is not used.
<code>burn</code>	An integer describing the number of MCMC iterations in <code>object</code> to be discarded as burn-in. If <code>burn <= 0</code> then no burn-in period will be discarded.
<code>na.action</code>	A function determining what should be done with missing values in <code>newdata</code> .
<code>olddata</code>	This is an optional component allowing predictions to be made conditional on data other than the data used to fit the model. If omitted, then it is assumed that forecasts are to be made relative to the final observation in the training data. If <code>olddata</code> is supplied then it will be filtered to get the distribution of the next state before a prediction is made, and it is assumed that the first entry in <code>newdata</code> comes immediately after the last entry in <code>olddata</code> .

The value for `olddata` depends on whether or not `object` contains a regression component.

- If a regression component is present, then `olddata` is a `data.frame` including variables with the same names as the data used to fit `object`, including the response.
- If no regression component is present, then `olddata` is a vector containing historical values of a time series.

`trials.or.exposure`

For logit or Poisson models, the number of binomial trials (or the exposure time) to assume at each time point in the forecast period. This can either be a scalar (if the number of trials is to be the same for each time period), or it can be a vector with length equal to `horizon` (if the model contains no regression term) or `nrow(newdata)` if the model contains a regression term.

`quantiles`

A numeric vector of length 2 giving the lower and upper quantiles to use for the forecast interval estimate.

...

This is a dummy argument included to match the signature of the generic `predict` function. It is not used.

Details

Samples from the posterior distribution of a Bayesian structural time series model. This function can be used either with or without contemporaneous predictor variables (in a time series regression).

If predictor variables are present, the regression coefficients are fixed (as opposed to time varying, though time varying coefficients might be added as state component). The predictors and response in the formula are contemporaneous, so if you want lags and differences you need to put them in the predictor matrix yourself.

If no predictor variables are used, then the model is an ordinary state space time series model.

Value

Returns an object of class `bsts.prediction`, which is a list with the following components.

<code>mean</code>	A vector giving the posterior mean of the prediction.
<code>interval</code>	A two (column/row?) matrix giving the upper and lower bounds of the 95 percent credible interval for the prediction.
<code>distribution</code>	A matrix of draws from the posterior predictive distribution. Each row in the matrix is one MCMC draw. Columns represent time.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts](#). [AddLocalLevel](#). [AddLocalLinearTrend](#). [AddGeneralizedLocalLinearTrend](#).

Examples

```
data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bst(y, state.specification = ss, niter = 500)
pred <- predict(model, horizon = 12, burn = 100)
plot(pred)
```

quarter

Find the quarter in which a date occurs

Description

Returns the quarter and year in which a date occurs.

Usage

```
Quarter(date)
```

Arguments

date A vector convertible to [POSIXlt](#). A [Date](#) or character is fine.

Value

A numeric vector identifying the quarter that each element of date corresponds to, expressed as a number of years since 1900. Thus Q1-2000 is 100.00, and Q3-2007 is 107.50.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```
Quarter(c("2008-02-29", "2008-04-29"))
# [1] 108.00 108.25
```

regularize.timestamps *Produce a Regular Series of Time Stamps*

Description

Given an set of timestamps that might contain duplicates and gaps, produce a set of timestamps that has no duplicates and no gaps.

Usage

```
RegularizeTimestamps(timestamps)

## Default S3 method:
RegularizeTimestamps(timestamps)

## S3 method for class 'numeric'
RegularizeTimestamps(timestamps)

## S3 method for class 'Date'
RegularizeTimestamps(timestamps)

## S3 method for class 'POSIXt'
RegularizeTimestamps(timestamps)
```

Arguments

timestamps A set of (possibly irregular or non-unique) timestamps. This could be a set of integers (like 1, 2, , 3...), a set of numeric like (1945, 1945.083, 1945.167, ...) indicating years and fractions of years, a [Date](#) object, or a [POSIXt](#) object. If the argument is NULL a NULL will be returned.

Value

A set of regularly spaced timestamps of the same class as the argument (which might be NULL).

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```
first <- as.POSIXct("2015-04-19 08:00:04")
monthly <- seq(from = first, length.out = 24, by = "month")
skip.one <- monthly[-8]
has.duplicates <- monthly
has.duplicates[2] <- has.duplicates[3]

reg1 <- RegularizeTimestamps(skip.one)
```

```

all.equal(reg1, monthly) ## TRUE

reg2 <- RegularizeTimestamps(has.duplicates)
all.equal(reg2, monthly) ## TRUE

```

residuals.bsts *Residuals from a bsts Object*

Description

Residuals (or posterior distribution of residuals) from a bsts object.

Usage

```

## S3 method for class 'bsts'
residuals(object,
  burn = SuggestBurn(.1, object),
  mean.only = FALSE,
  ...)

```

Arguments

object	An object of class bsts created by the function of the same name.
burn	The number of MCMC iterations to discard as burn-in.
mean.only	Logical. If TRUE then the mean residual for each time period is returned. If FALSE then the full posterior distribution is returned.
...	Not used. This argument is here to comply with the signature of the generic residuals function.

Value

If `mean.only` is TRUE then this function returns a vector of residuals with the same "time stamp" as the original series. If `mean.only` is FALSE then the posterior distribution of the residuals is returned instead, as a matrix of draws. Each row of the matrix is an MCMC draw, and each column is a time point. The colnames of the returned matrix will be the timestamps of the original series, as text.

See Also

[bsts](#), [plot.bsts](#).

rsxfs	<i>Retail sales, excluding food services</i>
-------	--

Description

A monthly time series of retail sales in the US, excluding food services. In millions of dollars. Seasonally adjusted.

Usage

```
data(rsxfs)
```

Format

zoo time series

Source

FRED. See <http://research.stlouisfed.org/fred2/series/RSXFS>

Examples

```
data(rsxfs)
plot(rsxfs)
```

shorten	<i>Shorten long names</i>
---------	---------------------------

Description

Removes common prefixes and suffixes from character vectors.

Usage

```
Shorten(words)
```

Arguments

words A character vector to be shortened.

Value

The argument words is returned, after common prefixes and suffixes have been removed. If all arguments are identical then no shortening is done.

Author(s)

Steven L. Scott <stevescott@google.com>

See Also

[bsts.mixed.](#)

Examples

```
Shorten(c("/usr/common/foo.tex", "/usr/common/barbarian.tex"))  
# returns c("foo", "barbarian")
```

```
Shorten(c("hello", "hellobye"))  
# returns c("", "bye")
```

```
Shorten(c("hello", "hello"))  
# returns c("hello", "hello")
```

```
Shorten(c("", "x", "xx"))  
# returns c("", "x", "xx")
```

```
Shorten("abcde")  
# returns "abcde"
```

simulate.fake.mixed.frequency.data

Simulate fake mixed frequency data

Description

Simulate a fake data set that can be used to test mixed frequency code.

Usage

```
SimulateFakeMixedFrequencyData(nweeks,  
                                xdim,  
                                number.nonzero = xdim,  
                                start.date = as.Date("2009-01-03"),  
                                sigma.obs = 1.0,  
                                sigma.slope = .5,  
                                sigma.level = .5,  
                                beta.sd = 10)
```


Arguments

nweeks	The number of weeks of data to simulate.
xdim	The dimension of the predictor variables to be simulated.
number.nonzero	The number nonzero coefficients. Must be less than or equal to xdim.
start.date	The date of the first simulated week.
sigma.obs	The residual standard deviation for the fine time scale model.
sigma.slope	The standard deviation of the slope component of the local linear trend model for the fine time scale data.
sigma.level	The standard deviation of the level component fo the local linear trend model for the fine time scale data.
beta.sd	The standard deviation of the regression coefficients to be simulated.

Details

The simulation begins by simulating a local linear trend model for nweeks to get the trend component.

Next a nweeks by xdim matrix of predictor variables is simulated as IID normal(0, 1) deviates, and a xdim-vector of regression coefficients is simulated as IID normal(0, beta.sd). The product of the predictor matrix and regression coefficients is added to the output of the local linear trend model to get fine.target.

Finally, fine.target is aggregated to the month level to get coarse.target.

Value

Returns a list with the following components

coarse.target	A zoo time series containing the monthly values to be modeled.
fine.target	A zoo time series containing the weekly observations that aggregate to coarse.target.
predictors	A zoo matrix corresponding to fine.target containing the set of predictors variables to use in <code>bsts.mixed</code> prediction.
true.beta	The vector of "true" regression coefficients used to simulate fine.target.
true.sigma.obs	The residual standard deviation that was used to simulate fine.target.
true.sigma.slope	The value of sigma.slope used to simulate fine.target.
true.sigma.level	The value of sigma.level use to simulate fine.target.
true.trend	The combined contribution of the simulated latent state on fine.target, including regression effects.
true.state	A matrix containin the fine-scale state of the model being simulated. Columns represent time (weeks). Rows correspond to regression (a constant 1), the local linear trend level, the local linear trend slope, the values of fine.target, and the weekly partial aggregates of coarse.target.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts.mixed](#), [AddLocalLinearTrend](#),

Examples

```
fake.data <- SimulateFakeMixedFrequencyData(nweeks = 100, xdim = 10)
plot(fake.data$coarse.target)
```

spike.slab.ar.prior *Spike and Slab Priors for AR Processes*

Description

Returns a spike and slab prior for the parameters of an AR(p) process.

Usage

```
SpikeSlabArPrior(
  lags,
  prior.inclusion.probabilities =
    GeometricSequence(lags, initial.value = .8, discount.factor = .8),
  prior.mean = rep(0, lags),
  prior.sd =
    GeometricSequence(lags, initial.value = .5, discount.factor = .8),
  sdy,
  prior.df = 1,
  expected.r2 = .5,
  sigma.upper.limit = Inf,
  truncate = TRUE)
```

Arguments

lags	A positive integer giving the maximum number of lags to consider.
prior.inclusion.probabilities	A vector of length lags giving the prior probability that the corresponding AR coefficient is nonzero.
prior.mean	A vector of length lags giving the prior mean of the AR coefficients. This should almost surely stay set at zero.
prior.sd	A vector of length lags giving the prior standard deviations of the AR coefficients, which are modeled as a-priori independent of one another.
sd	The sample standard deviation of the series being modeled.
expected.r2	The expected fraction of variation in the response explained by this AR proces.
prior.df	A positive number indicating the number of observations (time points) worth of weight to assign to the guess at expected.r2.
sigma.upper.limit	A positive number less than infinity truncates the support of the prior distribution to regions where the residual standard deviation is less than the specified limit. Any other value indicates support over the entire positive real line.
truncate	If TRUE then the support of the distribution is truncated to the region where the AR coefficients imply a stationary process. If FALSE the coefficients are unconstrained.

Value

A list of class SpikeSlabArPrior containing the information needed for the underlying C++ code to instantiate this prior.

Author(s)

Steven L. Scott <stevescott@google.com>

state.sizes *Compute state dimensions*

Description

Returns a vector containing the size of each state component (i.e. the state dimension) in the state vector.

Usage

```
StateSizes(state.specification)
```

Arguments

state.specification

A list containing state specification components, such as would be passed to [bsts](#).

Value

A numeric vector giving the dimension of each state component.

Author(s)

Steven L. Scott <stevescott@google.com>

Examples

```
y <- rnorm(1000)
state.specification <- AddLocalLinearTrend(list(), y)
state.specification <- AddSeasonal(state.specification, y, 7)
StateSizes(state.specification)
```

StateSpecification *Add a state component to a Bayesian structural time series model*

Description

Add a state component to the state.specification argument in a [bsts](#) model.

Author(s)

Steven L. Scott <stevescott@google.com>

References

Harvey (1990), "Forecasting, structural time series, and the Kalman filter", Cambridge University Press.

Durbin and Koopman (2001), "Time series analysis by state space methods", Oxford University Press.

See Also

[bsts](#). [SdPrior](#) [NormalPrior](#) [Ar1CoefficientPrior](#)

Examples

```

data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bst(y, state.specification = ss, niter = 500)
pred <- predict(model, horizon = 12, burn = 100)
plot(pred)

```

SuggestBurn

Suggested burn-in size

Description

Suggest the size of an MCMC burn in sample as a proportion of the total run.

Usage

```
SuggestBurn(proportion, bst.object)
```

Arguments

proportion The proportion of the MCMC run to discard as burn in.
bst.object An object of class [bsts](#).

Value

An integer number of iterations to discard.

See Also

[bsts](#)

summary.bsts

Summarize a Bayesian structural time series object

Description

Print a summary of a [bsts](#) object.

Usage

```

## S3 method for class 'bsts'
summary(object, burn = SuggestBurn(.1, object), ...)

```

Arguments

object	An object of class <code>bsts</code> created by the function of the same name.
burn	The number of MCMC iterations to discard as burn-in.
...	Additional arguments passed to <code>summary.lm.spike</code> if object has a regression component.

Value

Returns a list with the following elements.

residual.sd	The posterior mean of the residual standard deviation parameter.
prediction.sd	The standard deviation of the one-step-ahead prediction errors for the training data.
rsquare	Proportion by which the residual variance is less than the variance of the original observations.
relative.gof	Harvey's goodness of fit statistic. Let ν denote the one step ahead prediction errors, n denote the length of the series, and y denote the original series. The goodness of fit statistic is

$$1 - \frac{\sum_{i=1}^n \nu_i^2}{\sum_{i=2}^n n(\Delta y_i - \Delta \bar{y})^2}$$

This statistic is analogous to R^2 in a regression model, but the baseline model is a random walk with drift, instead of the mean of the data. Unlike a traditional R-square statistic, this can be negative.

size	Distribution of the number of nonzero coefficients appearing in the model
coefficients	If object contains a regression component then the output contains matrix with rows corresponding to coefficients, and columns corresponding to: <ul style="list-style-type: none"> • The posterior probability the variable is included. • The posterior probability that the variable is positive. • The conditional expectation of the coefficient, given inclusion. • The conditional standard deviation of the coefficient, given inclusion.

References

Harvey's goodness of fit statistic is from Harvey (1989) *Forecasting, structural time series models, and the Kalman filter*. Page 268.

See Also

[bsts](#), [plot.bsts](#), [summary.lm.spike](#)

Examples

```
data(AirPassengers)
y <- log(AirPassengers)
ss <- AddLocalLinearTrend(list(), y)
ss <- AddSeasonal(ss, y, nseasons = 12)
model <- bsts(y, state.specification = ss, niter = 100)
summary(model, burn = 20)
```

week.ends

Check to see if a week contains the end of a month or quarter

Description

Returns a logical vector indicating whether the given week contains the end of a month or quarter.

Usage

```
WeekEndsMonth(week.ending)
WeekEndsQuarter(week.ending)
```

Arguments

week.ending A vector of class [Date](#). Each entry contains the date of the last day in a week.

Value

A logical vector indicating whether the given week contains the end of a month or a quarter.

Author(s)

Steven L. Scott <stevescott@google.com>

See Also

[bsts.mixed](#).

Examples

```
week.ending <- as.Date(c("2011-10-01",
                        "2011-10-08",
                        "2011-12-03",
                        "2011-12-31"))
WeekEndsMonth(week.ending) == c(TRUE, FALSE, TRUE, TRUE)
WeekEndsQuarter(week.ending) == c(TRUE, FALSE, FALSE, TRUE)
```

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