Package 'spTimer'

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Title Spatio-Temporal Bayesian Modelling Using R			
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${\bf URL} \ {\tt http://www.southampton.ac.uk/~sks/research/papers/spTimeRpaper.pdf}$			
Depends R (>= 2.14.0), coda, lattice, forecast			
Description The package is able to fit, spatially predict and temporally forecast large amounts of space-time data using [1] Bayesian Gaussian Process (GP) Models, [2] Bayesian Auto-Regressive (AR) Models, and [3] Bayesian Gaussian Predictive Processes (GPP) based AR Models.			
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R topics documented:			
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Description

This package uses different hierarchical Bayesian spatio-temporal modelling strategies, namely the gaussian processes (GP) models, the autoregressive (AR) models, and models using Gaussian predictive processes (GPP) approximation to analyse space-time observations.

Details

Package: spTimer Type: Package

The back-end code of this package is built under c language.

Main functions used:

```
> spT.Gibbs
> predict.spT
Some other important functions:
> spT.priors
> spT.initials
> spT.decay
> spT.time
```

Data descriptions: > NYdata

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References

Bakar, K. S. and Sahu, S. K. (2013) spTimer: Spatio-Temporal Bayesian Modelling Using R. Technical Report, University of Southampton, UK.

Sahu, S. K. and Bakar, K. S. (2012) A comparison of Bayesian Models for Daily Ozone Concentration Levels Statistical Methodology, 9, 144-157.

Sahu, S. K. and Bakar, K. S. (2012) Hierarchical Bayesian auto-regressive models for large space time data with applications to ozone concentration modelling. Applied Stochastic Models in Business and Industry, 28, 395-415.

Sahu, S. K., Bakar, K. S. and Awang, N. (2013) Bayesian Forecasting Using Hierarchical Spatiotemporal Models with Applications to Ozone Levels in the Eastern United States. Technical Report, University of Southampton.

Sahu, S.K., Gelfand, A.E., & Holland, D.M. (2007). High-Resolution Space-Time Ozone Modelling for Assessing Trends. Journal of the American Statistical Association, 102, 1221-1234.

See Also

Packages 'forecast'; 'spBayes'; 'maps'; 'MBA'; 'coda'; website: http://www.r-project.org/.

as.forecast.object

Conversion of spT object into forecast object

Description

This function is used to convert the spT object from the temporal prediction output of "spTimer" into forecast object of the package "forecast".

Usage

```
as.forecast.object(object, site=1, level=c(80,95), ...)
```

Arguments

object Object of class inheriting from "spT".

site Selection of location/site for which the time-series will convert into forecast

object.

level Confidence level for temporal prediction intervals, see details in forecast.

... Other arguments, see details in forecast.

Value

An object class "forecast"

model Name of the fitted model.

method Name of the forecasting method.

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mean MCMC mean for the temporal predictions.

lower Lower limits for the temporal prediction intervals obtained from the MCMC

samples.

upper Upper limits for the temporal prediction intervals obtained from the MCMC

samples.

level Prediction interval limits.

The data used for model fitting in time-series format, see ts.

residuals Residuals of the fiited model.

fitted Fitted MCMC mean.

See Also

```
spT.Gibbs, predict.spT.
```

Examples

```
## Not run:
##

# 'out' is the output of spT class obtained from temporal prediction
fobj<-as.forecast.object(out)
class(fobj)
summary(fobj)
plot(fobj)

##

## End(Not run)</pre>
```

NYdata

Observations of ozone concentration levels, maximum temperature and wind speed.

Description

This data set contains values of daily 8-hour maximum average ozone concentrations (parts per billion (ppb)), maximum temperature (in degree Celsius), wind speed (knots), and relative humidity, obtained from 28 monitoring sites of New York, USA.

Total 4 seperate subset datasets are created from NYdata that contains same variables as NYdata.

- 1. DataFit: This data is used for model fitting, contains 20 monitoring locations with 62 days observations; total 1200 rows.
- 2. DataValPred: This data is used for model validation for spatial prediction, contains 8 monitoring locations with 60 days observations; total 480 rows.
- 3. DataValFore: This data is used for model validation for temporal prediction in the unobserved locations, contains 8 monitoring locations with 2 days observations; total 8 rows.

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4. DataFitFore: This data is used for model validation for temporal prediction in the observed locations, contains 20 monitoring locations with 2 days observations; total 40 rows.

We also include a dataset that contains gridded observations over NY state.

NYgrid: This dataset contains total 6200 rows for 62 days of observations for 10x10 = 100 grid points.

Usage

NYdata

Format

Columns for NYdata: each contains 1798 observations.

```
• 1st col = Site index (s.index),
```

- 2nd col = Longitude,
- 3rd col = Latitude,
- 4th col = Year,
- 5th col = Month,
- 6th col = Day,
- 7th col = Ozone (o8hrmax),
- 8th col = Maximum temperature (cMAXTMP),
- 9th col = Wind speed (WDSP).
- 10th col = Relative humidity (RH).

Source

US EPA

See Also

```
DataFit, DataFitFore, DataValFore, DataValPred, NYgrid.
```

```
## Not run:
##
  library("spTimer")
# NY data
  data(NYdata)
  head(NYdata)
# plots in NY map
  NYsite<-unique(cbind(NYdata[,1:3]))
  head(NYsite)
# map
  library(maps)
  map(database="state",regions="new york")
  points(NYsite[,2:3],pch=19)</pre>
```

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```
# DataFit
 data(DataFit)
 head(DataFit)
# DataValPred
 data(DataValPred)
 head(DataValPred)
# DataValFore
 data(DataValFore)
 head(DataValFore)
# DataFitFore
 data(DataFitFore)
 head(DataFitFore)
# Plot fitted and validation locations in map
 fit.coords<-unique(cbind(DataFit[,1:3]))</pre>
 val.coords<-unique(cbind(DataValPred[,1:3]))</pre>
 library(maps)
 map(database="state",regions="new york")
 points(fit.coords[,2:3],pch=19,col=2)
 points(val.coords[,2:3],pch=7,col=1)
 legend(x=-78,y=41.5,pch=c(19,7),col=c(2,1),bty="n",
 legend=c("Fitted locations", "Validation locations"))
# Grid data
 data(NYgrid)
 head(NYgrid)
 grid.coords<-unique(cbind(NYgrid[,8:9]))</pre>
 library(maps)
 plot(grid.coords,pch=19,col=1)
 map(database="state",regions="new york",add=TRUE)
##
## End(Not run)
```

plot.spT

Plots for spTimer output.

Description

This function is used to obtain MCMC summary and residual plots.

Usage

```
## S3 method for class 'spT'
## S3 method for class 'spT'
plot(x, residuals=FALSE, ...)
##
```

Arguments

x Object of class inheriting from "spT".

residuals If TRUE then plot residual vs. fitted and normal qqplot of the residuals. If

FALSE then plot MCMC samples of the parameters using coda package. De-

faults value is FALSE.

... Other arguments.

See Also

```
spT.Gibbs.
```

Examples

```
## Not run:
##

plot(out) # where out is the output from spT class
plot(out, residuals=TRUE) # where out is the output from spT class
##

## End(Not run)
```

predict.spT

Spatial and temporal predictions for the spatio-temporal models.

Description

This function is used to obtain spatial predictions in the unknown locations and also to get the temporal forecasts using MCMC samples.

Usage

Arguments

object Object of class inheriting from "spT".

newdata The data set for the covariate values for spatial prediction or temporal forecasts.

This data should have the same space-time structure as the original data frame.

newcoords The coordinates for the prediction or forecast sites. The locations are in similar

format to coords.

foreStep Number of K-step (time points) ahead forecast, K=1,2, ...; Only applicable if

type="temporal".

type If "spatial" the do spatial prediction and if "temporal" the do temporal prediction

or forecast. Default value is "spatial".

nBurn Number of burn-in. Initial MCMC samples to discard before making inference.

tol.dist Minimum tolerance distance limit between fitted and predicted locations.

predAR The prediction output, if forecasts are in the prediction locations. Only applica-

ble if type="forecast" and data fitted with the "AR" model.

... Other arguments.

Value

pred.samples or fore.samples

Prediction or forecast MCMC samples.

pred.coords or fore.coords

prediction or forecast coordinates.

Mean Average of the MCMC predictions
Median Median of the MCMC predictions

SD Standard deviation of the MCMC predictions

Low Lower limit for the 95 percent CI of the MCMC predictions

Up Upper limit for the 95 percent CI of the MCMC predictions

computation.time

The computation time.

model The model method used for prediction.

type "spatial" or "temporal".

... Other values "obsData", "fittedData" and "residuals" are provided only for tem-

poral prediction. This is to analyse the codespTimer forecast output using pack-

age forecast through function as.forecast.object.

References

Bakar, K. S. and Sahu, S. K. (2013) spTimer: Spatio-Temporal Bayesian Modelling Using R. Technical Report, University of Southampton, UK.

Sahu, S. K. and Bakar, K. S. (2012) A comparison of Bayesian Models for Daily Ozone Concentration Levels Statistical Methodology, 9, 144-157.

Sahu, S. K. and Bakar, K. S. (2012) Hierarchical Bayesian auto-regressive models for large space time data with applications to ozone concentration modelling. Applied Stochastic Models in Business and Industry, 28, 395-415.

Sahu, S. K., Bakar, K. S. and Awang, N. (2013) Bayesian Forecasting Using Hierarchical Spatiotemporal Models with Applications to Ozone Levels in the Eastern United States. Technical Report, University of Southampton.

See Also

spT.Gibbs, as.forecast.object.

```
## Not run:
## The GP models:
#####################################
## Spatial prediction/interpolation
##
# Read data
data(DataValPred)
# Define prediction coordinates
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
# Spatial prediction using spT.Gibbs output
set.seed(11)
pred.gp <- predict(post.gp, newdata=DataValPred, newcoords=pred.coords)</pre>
print(pred.gp)
names(pred.gp)
# validation criteria
spT.validation(DataValPred$o8hrmax,c(pred.gp$Mean))
## Temporal prediction/forecast
## 1. In the unobserved locations
##
# Read data
data(DataValFore);
# define forecast coordinates
fore.coords<-as.matrix(unique(cbind(DataValFore[,2:3])))</pre>
# Two-step ahead forecast, i.e., in day 61 and 62
# in the unobserved locations using output from spT.Gibbs
set.seed(11)
fore.gp <- predict(post.gp, newdata=DataValFore, newcoords=fore.coords,</pre>
           type="temporal", foreStep=2)
print(fore.gp)
names(fore.gp)
# Forecast validations
spT.validation(DataValFore$08hrmax,c(fore.gp$Mean))
# Use of "forecast" class
tmp<-as.forecast.object(fore.gp, site=1) # default for site 1</pre>
plot(tmp)
```

```
##
## Temporal prediction/forecast
## 2. In the observed/fitted locations
##
# Read data
data(DataFitFore)
# Define forecast coordinates
fore.coords<-as.matrix(unique(cbind(DataFitFore[,2:3])))</pre>
# Two-step ahead forecast, i.e., in day 61 and 62,
# in the fitted locations using output from spT.Gibbs
set.seed(11)
fore.gp <- predict(post.gp, newdata=DataFitFore, newcoords=fore.coords,</pre>
           type="temporal", foreStep=2)
print(fore.gp)
names(fore.gp)
# Forecast validations
spT.validation(DataFitFore$o8hrmax,c(fore.gp$Mean)) #
# Use of "forecast" class
tmp<-as.forecast.object(fore.gp, site=5) # for site 5</pre>
plot(tmp)
## Fit and spatially prediction simultaneously
##
# Read data
data(DataFit);
data(DataValPred)
# Define the coordinates
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
# MCMC via Gibbs will provide output in *.txt format
# from C routine to avoide large data problem in R
set.seed(11)
post.gp.fitpred <- spT.Gibbs(formula=o8hrmax ~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="GP", coords=coords,
         newcoords=pred.coords, newdata=DataValPred,
         scale.transform="SQRT")
print(post.gp.fitpred)
summary(post.gp.fitpred)
coef(post.gp.fitpred)
plot(post.gp.fitpred,residuals=TRUE)
{\tt names(post.gp.fitpred)}
# validation criteria
```

```
spT.validation(DataValPred$o8hrmax,c(post.gp.fitpred$prediction[,1]))
## The AR models:
## Spatial prediction/interpolation
##
# Read data
data(DataValPred)
# Define prediction coordinates
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
# Spatial prediction using spT.Gibbs output
set.seed(11)
pred.ar <- predict(post.ar, newdata=DataValPred, newcoords=pred.coords)</pre>
print(pred.ar)
names(pred.ar)
# validation criteria
spT.validation(DataValPred$o8hrmax,c(pred.ar$Mean))
## Temporal prediction/forecast
## 1. In the unobserved locations
# Read data
data(DataValFore);
# define forecast coordinates
fore.coords<-as.matrix(unique(cbind(DataValFore[,2:3])))</pre>
# Two-step ahead forecast, i.e., in day 61 and 62
# in the unobserved locations using output from spT.Gibbs
set.seed(11)
fore.ar <- predict(post.ar, newdata=DataValFore, newcoords=fore.coords,</pre>
           type="temporal", foreStep=2, predAR=pred.ar)
print(fore.ar)
names(fore.ar)
# Forecast validations
spT.validation(DataValFore$08hrmax,c(fore.ar$Mean))
# Use of "forecast" class
tmp<-as.forecast.object(fore.ar, site=1) # default for site 1</pre>
plot(tmp)
```

```
## Temporal prediction/forecast
## 2. In the observed/fitted locations
# Read data
data(DataFitFore)
# Define forecast coordinates
fore.coords<-as.matrix(unique(cbind(DataFitFore[,2:3])))</pre>
# Two-step ahead forecast, i.e., in day 61 and 62,
\# in the fitted locations using output from spT.Gibbs
set.seed(11)
fore.ar <- predict(post.ar, newdata=DataFitFore, newcoords=fore.coords,</pre>
           type="temporal", foreStep=2)
print(fore.ar)
names(fore.ar)
# Forecast validations
spT.validation(DataFitFore$o8hrmax,c(fore.ar$Mean)) #
# Use of "forecast" class
tmp<-as.forecast.object(fore.ar, site=1) # default for site 1</pre>
plot(tmp)
## Fit and spatially prediction simultaneously
# Read data
data(DataFit);
data(DataValPred)
# Define the coordinates
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
# MCMC via Gibbs will provide output in *.txt format
# from C routine to avoide large data problem in R
set.seed(11)
post.ar.fitpred <- spT.Gibbs(formula=o8hrmax ~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="AR", coords=coords,
         newcoords=pred.coords, newdata=DataValPred,
         scale.transform="SQRT")
print(post.ar.fitpred)
summary(post.ar.fitpred)
coef(post.ar.fitpred)
names(post.ar.fitpred)
# validation criteria
spT.validation(DataValPred$o8hrmax,c(post.ar.fitpred$prediction[,1]))
```

```
## The GPP approximation models:
##
## Spatial prediction/interpolation
##
# Read data
data(DataValPred)
# Define prediction coordinates
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
# Spatial prediction using spT.Gibbs output
set.seed(11)
pred.gpp <- predict(post.gpp, newdata=DataValPred, newcoords=pred.coords)</pre>
print(pred.gpp)
names(pred.gpp)
# validation criteria
spT.validation(DataValPred$o8hrmax,c(pred.gpp$Mean))
##
## Temporal prediction/forecast
## 1. In the unobserved locations
##
# Read data
data(DataValFore);
# define forecast coordinates
fore.coords<-as.matrix(unique(cbind(DataValFore[,2:3])))</pre>
# Two-step ahead forecast, i.e., in day 61 and 62
# in the unobserved locations using output from spT.Gibbs
set.seed(11)
fore.gpp <- predict(post.gpp, newdata=DataValFore, newcoords=fore.coords,</pre>
           type="temporal", foreStep=2)
print(fore.gpp)
names(fore.gpp)
# Forecast validations
spT.validation(DataValFore$08hrmax,c(fore.gpp$Mean))
# Use of "forecast" class
tmp<-as.forecast.object(fore.gpp, site=1) # default for site 1</pre>
plot(tmp)
## Temporal prediction/forecast
## 2. In the observed/fitted locations
##
```

```
# Read data
data(DataFitFore)
# Define forecast coordinates
fore.coords<-as.matrix(unique(cbind(DataFitFore[,2:3])))</pre>
# Two-step ahead forecast, i.e., in day 61 and 62,
# in the fitted locations using output from spT.Gibbs
set.seed(11)
fore.gpp <- predict(post.gpp, newdata=DataFitFore, newcoords=fore.coords,</pre>
           type="temporal", foreStep=2)
print(fore.gpp)
names(fore.gpp)
# Forecast validations
spT.validation(DataFitFore$o8hrmax,c(fore.gpp$Mean)) #
# Use of "forecast" class
tmp<-as.forecast.object(fore.gpp, site=1) # default for site 1</pre>
plot(tmp)
## Fit and spatially prediction simultaneously
##
# Read data
data(DataFit);
data(DataValPred)
# Define the coordinates
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
knots.coords<-spT.grid.coords(Longitude=c(max(coords[,1]),</pre>
              min(coords[,1])),Latitude=c(max(coords[,2]),
              min(coords[,2])), by=c(4,4))
# MCMC via Gibbs will provide output in *.txt format
# from C routine to avoide large data problem in R
set.seed(11)
post.gpp.fitpred <- spT.Gibbs(formula=08hrmax ~cMAXTMP+WDSP+RH,</pre>
         {\tt data=DataFit,\ model="GP",\ coords=coords,\ knots.coords=knots.coords,}
         newcoords=pred.coords, newdata=DataValPred,
         scale.transform="SQRT")
print(post.gpp.fitpred)
summary(post.gpp.fitpred)
coef(post.gpp.fitpred)
plot(post.gpp.fitpred, residuals=TRUE)
names(post.gpp.fitpred)
# validation criteria
spT.validation(DataValPred$o8hrmax,c(post.gpp.fitpred$prediction[,1]))
##
```

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```
## End(Not run)
```

Description

This function is used to check the minimum distance between two locations.

Usage

```
spT.check.locations(fit.locations, pred.locations,
   method="geodetic:km", tol = 5)
spT.check.sites.inside(coords, method)
```

Arguments

fit.locations The locations for the fitted observations.

pred.locations The locations for the predicted observations.

method The distance measurement. The available methods are "geodetic:km", "geode-

tic:mile", "euclidean", "maximum", "manhattan", "canberra", "binary" or "minkowski".

tol The tolerance limit for the distance.

coords The longitude and latitude positions in a matrix format.

Details

spT.check.locations is used to check minimum distance between two locations, e.g., fitted and prediction. spT.check.sites.inside is used to check distance within the location points. Here, the tol. limit is 0.01. If output shows nothing then we can say distances are alright. These functions are used to avoid of occuring non-positive definite correlation matrix.

See Also

```
spT.geodist, dist, spT.data.selection.
```

```
## Not run:
##

data(NYdata)
head(NYdata)
NYsite<-unique(NYdata[,1:3])
# Sample 4 sites randomly from the data NYdata.</pre>
```

spT.data.selection

```
r4<-spT.data.selection(data=NYsite, random=TRUE, num.rs=4)

# Choose purposively sites numbered as 2, 8, and 12, 15.

p4<-spT.data.selection(data=NYsite, random=FALSE, s=c(2,8,12,15))

# Check locations of datasets r4 and p4

spT.check.locations(fit.locations=r4, pred.locations=p4, method="geodetic:km", tol=5)

# spT.check.sites.inside(NYsite[,2:3],method="geodetic:km")

# if nothing appears then distances are alright

##

## End(Not run)</pre>
```

spT.data.selection

Selection of Spatial data from a big dataset.

Description

This command selects a part of the big spatial dataset using the site numbers.

Usage

Arguments

data	The dataset.
random	Logical value: if TRUE then the <code>num.rs</code> sites are randomly sampled, if FALSE then we need to provide the value for s .
num.rs	The number of sites to be selected, e.g., 3.
S	The site numbers to be selected, e.g., $c(2,8,12)$.
reverse	Logical value: if TRUE then num.rs will be discarded from the data.

See Also

NYdata.

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Examples

```
## Not run:
##

# Load ozone concentration data for New York.

data(NYdata)
NYdata

# Sample 4 sites randomly from the data NYdata.

r4<-spT.data.selection(data=NYdata, random=TRUE, num.rs=4)

# Choose purposively defined sites numbered as 2, 8, and 12.

p4<-spT.data.selection(data=NYdata, random=FALSE, s=c(2,8,12))

# Donot choose purposively defined sites numbered as 2, 8, and 12.

p4<-spT.data.selection(data=NYdata, random=FALSE, s=c(2,8,12), reverse=TRUE)

##
## End(Not run)</pre>
```

spT.decay

Choice for sampling spatial decay parameter ϕ *.*

Description

This function initialises the sampling method for the spatial decay parameter ϕ .

Usage

```
spT.decay(type="MH", tuning=NULL, limit=NULL, segments=NULL, value=NULL)
```

Arguments

type	The sampling method, currently available methods are, DISCRETE and MH. One can also used FIXED value for ϕ parameter.
tuning	If MH type is used then need to define the tuning parameter for ϕ .
limit	If DISCRETE type is used then need to define the lower and upper limits.
segments	If DISCRETE type is used then need to define the number of segments for the range of limits.
value	If FIXED type is used then need to define the value for ϕ .

spT.geodist

See Also

```
spT.Gibbs.
```

Examples

spT.geodist

Geodetic/geodesic Distance

Description

This geodetic distance provides the distance between the locations in Kilometers (k.m.) and Miles, using spherical law of Cosines.

Usage

```
spT.geodist(Lon, Lat, KM = TRUE)
spT.geo.dist(point1, point2)
spT.geo_dist(points)
```

Arguments

Lon	The longitude position.
Lat	The latitude position.
KM	A logical value, if 'TRUE' then output is in 'kilometers', otherwise in 'miles'.
point1	In the form of (longitude, latitude) position.
point2	In the form of (longitude, latitude) position.
points	In the form of points 1:(longitude, latitude) 2:(longitude, latitude) positions.

Details

spT.geodist is used to get geodetic distance in both miles and kilometers. spT.geo.dist is only used to get geodetic distance in kilometers with a different format. spT.geo_dist is only used to get geodetic distance in kilometers with a different format.

See Also

```
NYdata, spT.grid.coords.
```

Examples

```
## Not run:
##
# Load 28 ozone monitoring locations of New York.
data(NYdata)
head(NYdata)
NYsite<-unique(NYdata[,1:3])
# Find the geodetic distance in km
spT.geodist(Lon=NYsite$Longitude, Lat=NYsite$Latitude, KM=TRUE)
# Find the geodetic distance in miles
spT.geodist(Lon=NYsite$Longitude, Lat=NYsite$Latitude, KM=FALSE)
##
# using spT.geo.dist
point1<-c(-73.757,42.681)
point2<-c(-73.881,40.866)
spT.geo.dist(point1,point2)
# using spT.geo_dist
points<-c(point1,point2)</pre>
spT.geo_dist(points)
##
## End(Not run)
```

 ${\tt spT.Gibbs}$

MCMC sampling for the spatio-temporal models.

Description

This function is used to draw MCMC samples using the Gibbs sampler.

Usage

```
spT.Gibbs(formula, data=parent.frame(), model, time.data,
coords, knots.coords, newcoords, newdata, priors, initials,
nItr, nBurn, report, tol.dist, distance.method, cov.fnc,
scale.transform, spatial.decay, annual.aggrn)
```

Arguments

formula

The symnbolic description of the model equation of the regression part of the space-time model. data An optional data frame containing the variables in the model. If omitted, the variables are taken from environment(formula), typically the environment from which spT.Gibbs is called. The data should be ordered first by the time and then by the sites specified by the coords below. One can also supply coordinates through this argument, where coordinate names should be "Latitude"/"Longitude"

or "xcoords"/"ycoords".

The spatio-temporal models to be fitted, current input: "GP", "AR", and "GPP". model

Defining the segments of the time-series set up using the function spT.time. time.data

coords The n by 2 matrix defining the locations (e.g., longitude/easting, latitude/northing) of the fitting sites, where n is the number of fitting sites. One can also supply coordinates through the argument data, where coordinate names should be "Lat-

itude"/"Longitude" or "xcoords"/"ycoords".

The locations of the knots in similar format to coords above, only required if knots.coords

model="GPP".

The locations of the prediction sites in similar format to coords above, only newcoords

required if fit and predictions are done simultaneously.

newdata The covariate values at the prediction sites specified by pred.coords. This

should have same space-time structure as the original data frame.

The prior distributions for the parameters. Default distributions are specified if priors

these are not specified. If priors=NULL a flat prior distribution will be used with

large variance. See details in spT.priors.

initials The preferred initial values for the parameters. If omitted, default values are

provided automatically. Further details are provided in spT.initials.

nItr Number of MCMC iterations. Default value is 13000.

nBurn Number of burn-in samples. This number of samples will be discarded before

making any inference. Default value is 3000.

Number of reports to display while running the Gibbs sampler. Defaults to numreport

ber of iterations.

distance.method

The preferred method to calculate the distance between any two locations. The available options are "geodetic:km", "geodetic:mile", "euclidean", "maximum", "manhattan", and "canberra". See details in dist.

tol.dist Minimum separation distance between any two locations out of those specified

by coords, knots.coords and pred.coords. The default is 0.005. The programme will exit if the minimum distance is less than the non-zero specified value. This

will ensure non-singularity of the covariance matrices.

cov.fnc Covariance function for the spatial effects. The available options are "exponen-

tial", "gaussian", "spherical" and "matern". If "matern" is used then by default the smooth parameter (ν) is estimated from (0,1) uniform distribution using dis-

crete samples.

scale.transform

The transformation method for the response variable. Currently implemented

options are: "NONE", "SQRT", and "LOG" with "NONE" as the deault.

spatial.decay Provides the options for sampling the spatial decay parameter ϕ . Currently im-

plemented options are "DISCRETE", "MH" or "FIXED" and further options for

each of these are specified by spT. decay. The default is "MH".

annual aggrn This provides the options for calculating annual summary statistics by aggregat-

ing different time segments (e.g., annual mean). Currently implemented options are: "NONE", "ave" and "an4th", where "ave" = annual average, "an4th"= annual 4th highest. Only applicable if spT.time inputs more than one segment

and when fit and predict are done simultaneously.

Value

accept The acceptance rate for the ϕ parameter if the "MH" method of sampling is

chosen.

phip MCMC samples for the parameter ϕ .

nup MCMC samples for the parameter ν . Only available if "matern" covariance

function is used.

sig2eps MCMC samples for the parameter σ_{ϵ}^2 . sig2etap MCMC samples for the parameter σ_{η}^2 . betap MCMC samples for the parameter β . op MCMC samples for the true observations.

fitted MCMC summary (mean and sd) for the fitted values.

tol.dist Minimum tolerance distance limit between the locations.

distance.method

Name of the distance calculation method.

cov. fnc Name of the covariance function used in model fitting.

scale.transform

Name of the scale.transformation method.

sampling.sp.decay

The method of sampling for the spatial decay parameter ϕ .

covariate.names

Name of the covariates used in the model.

Distance.matrix

The distance matrix.

coords The coordinate values.

n Total number of sites.

Total number of segments in time, e.g., years.Total points of time, e.g., days within each year.

p Total number of model coefficients, i.e., β 's including the intercept.

initials The initial values used in the model.

priors The prior distributions used in the model.

PMCC The predictive model choice criteria obtained by minimising the expected value

of a loss function, see Gelfand and Ghosh (1998). Results for both goodness of

fit and penalty are given.

iterations The number of samples for the MCMC chain, without burn-in.

nBurn The number of burn-in period for the MCMC chain.

computation.time

The computation time required for the fitted model.

model The spatio-temporal model used for analyse the data.

Text Output This option is only applicable when fit and predictions are done simultaneously.

For GP models:

OutGP_Values_Parameter.txt: (nItr x parameters matrix) has the MCMC samples for the parameters, ordered as: beta's, sig2eps, sig2eta, and phi.

OutGP_Stats_FittedValue.txt: (N x 2) matrix of fitted summary, with 1st column as mean and 2nd column as standard deviations, where N=nrT.

OutGP_Stats_PredValue.txt: ((predsites*r*T) x 2) matrix of prediction summary, with 1st column as mean and 2nd column as standard deviations.

OutGP_Values_Prediction.txt: (nItr x (predsites*r*T)) matrix of MCMC predicted values in the predicted sites.

If annual.aggregation="ave" then we get text output as:

OutGP_Annual_Average_Prediction.txt: (nItr x (predsites*r)) matrix.

If annual.aggregation="an4th" then we get text output as:

OutGP_Annual_4th_Highest_Prediction.txt: (nItr x (predsites*r)) matrix.

For AR models:

OutAR_Values_Parameter.txt: (nItr x parameters matrix) has the MCMC samples for the parameters, ordered as: beta's, rho, sig2eps, sig2eta, mu_l's, sig2l's and phi.

OutAR_Stats_TrueValue.txt: (N x 2) matrix of true summary values, with 1st column as mean and 2nd column as standard deviations.

OutAR_Stats_FittedValue.txt: (N x 2) matrix of fitted summary, with 1st column as mean and 2nd column as standard deviations.

OutAR_Stats_PredValue.txt: ((predsites*r*T) x 2) matrix of prediction summary, with 1st column as mean and 2nd column as standard deviations.

OutAR_Values_Prediction.txt: (nItr x (predsites*r*T)) matrix of MCMC predicted values in the predicted sites.

If annual.aggregation="ave" then we get text output as:

OutAR_Annual_Average_Prediction.txt: (nItr x (predsites*r)) matrix.

If annual.aggregation="an4th" then we get text output as:
OutAR_Annual_4th_Highest_Prediction.txt: (nItr x (predsites*r)) matrix.

For models using GPP approximations:

OutGPP_Values_Parameter.txt: (nItr x parameters matrix) has the MCMC samples for the parameters, ordered as: beta's, rho, sig2eps, sig2eta, and phi.

OutGPP_Stats_FittedValue.txt: (N x 2) matrix of fitted summary, with 1st column as mean and 2nd column as standard deviations.

OutGPP_Stats_PredValue.txt: ((predsites*r*T) x 2) matrix of prediction summary, with 1st column as mean and 2nd column as standard deviations.

OutGPP_Values_Prediction.txt: (nItr x (predsites*r*T)) matrix of MCMC predicted values in the predicted sites.

If annual.aggregation="ave" then we get text output as:

OutGPP_Annual_Average_Prediction.txt: (nItr x (predsites*r)) matrix.

If annual.aggregation="an4th" then we get text output as:

OutGPP_Annual_4th_Highest_Prediction.txt: (nItr x (predsites*r)) matrix.

References

Bakar, K. S. and Sahu, S. K. (2013) spTimer: Spatio-Temporal Bayesian Modelling Using R. Technical Report, University of Southampton, UK.

Sahu, S. K. and Bakar, K. S. (2012) A comparison of Bayesian Models for Daily Ozone Concentration Levels Statistical Methodology, 9, 144-157.

Sahu, S. K. and Bakar, K. S. (2012) Hierarchical Bayesian auto-regressive models for large space time data with applications to ozone concentration modelling. Applied Stochastic Models in Business and Industry, 28, 395-415.

Sahu, S. K., Bakar, K. S. and Awang, N. (2013) Bayesian Forecasting Using Hierarchical Spatiotemporal Models with Applications to Ozone Levels in the Eastern United States. Technical Report, University of Southampton.

See Also

```
spT.priors, spT.initials, spT.geodist, dist, summary.spT, plot.spT, predict.spT.
```

```
## Model fitting
##
# Read data
data(DataFit);
# Define the coordinates
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
# MCMC via Gibbs using default choices
set.seed(11)
post.gp <- spT.Gibbs(formula=o8hrmax ~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="GP", coords=coords,
         scale.transform="SQRT")
print(post.gp)
# MCMC via Gibbs not using default choices
# define the time-series
time.data<-spT.time(t.series=60,segment=1)</pre>
# hyper-parameters for the prior distributions
priors<-spT.priors(model="GP",var.prior=Gam(2,1),</pre>
        beta.prior=Nor(0,10^4))
# initial values for the model parameters
initials<-spT.initials(model="GP", sig2eps=0.01,</pre>
            sig2eta=0.5, beta=NULL, phi=0.001)
# input for spatial decay, any one approach from below
#spatial.decay<-spT.decay(type="FIXED", value=0.01)</pre>
spatial.decay<-spT.decay(type="MH", tuning=0.08)</pre>
#spatial.decay<-spT.decay(type="DISCRETE",limit=c(0.01,0.02),segments=5)</pre>
# Iterations for the MCMC algorithms
nItr<-5000
# MCMC via Gibbs
set.seed(11)
post.gp <- spT.Gibbs(formula=08hrmax ~ cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="GP", time.data=time.data,
         coords=coords, priors=priors, initials=initials,
         nItr=nItr, nBurn=0, report=nItr,
         tol.dist=2, distance.method="geodetic:km",
         cov.fnc="exponential", scale.transform="SQRT",
         spatial.decay=spatial.decay)
print(post.gp)
# Summary and plots
summary(post.gp)
summary(post.gp,pack="coda")
plot(post.gp)
plot(post.gp,residuals=TRUE)
```

```
coef(post.gp)
terms(post.gp)
formula(post.gp)
model.frame(post.gp)
model.matrix(post.gp)
# Model selection criteria
post.gp$PMCC
## Fit and spatially prediction simultaneously
##
# Read data
data(DataFit);
data(DataValPred)
# Define the coordinates
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
# MCMC via Gibbs will provide output in *.txt format
# from C routine to avoide large data problem in R
set.seed(11)
post.gp.fitpred <- spT.Gibbs(formula=o8hrmax ~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="GP", coords=coords,
         newcoords=pred.coords, newdata=DataValPred,
         scale.transform="SQRT")
print(post.gp.fitpred)
summary(post.gp.fitpred)
coef(post.gp.fitpred)
plot(post.gp.fitpred)
names(post.gp.fitpred)
# validation criteria
spT.validation(DataValPred$o8hrmax,c(post.gp.fitpred$prediction[,1]))
#####################################
## The AR models:
#####################################
## Model fitting
##
# Read data
data(DataFit);
# Define the coordinates
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
# MCMC via Gibbs using default choices
```

```
set.seed(11)
post.ar <- spT.Gibbs(formula=o8hrmax ~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="AR", coords=coords,
         scale.transform="SQRT")
print(post.ar)
# MCMC via Gibbs not using default choices
# define the time-series
time.data<-spT.time(t.series=60,segment=1)</pre>
# hyper-parameters for the prior distributions
priors<-spT.priors(model="AR",var.prior=Gam(2,1),</pre>
        beta.prior=Nor(0,10<sup>4</sup>))
# initial values for the model parameters
initials<-spT.initials(model="AR", sig2eps=0.01,</pre>
            sig2eta=0.5, beta=NULL, phi=0.001)
# Input for spatial decay
#spatial.decay<-spT.decay(type="FIXED", value=0.01)</pre>
spatial.decay<-spT.decay(type="MH", tuning=0.08)</pre>
#spatial.decay<-spT.decay(type="DISCRETE",limit=c(0.01,0.02),segments=5)</pre>
# Iterations for the MCMC algorithms
nItr<-5000
# MCMC via Gibbs
set.seed(11)
post.ar <- spT.Gibbs(formula=08hrmax~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="AR", time.data=time.data,
         coords=coords, priors=priors, initials=initials,
         nItr=nItr, nBurn=0, report=nItr,
         tol.dist=2, distance.method="geodetic:km",
         cov.fnc="exponential", scale.transform="SQRT",
         spatial.decay=spatial.decay)
print(post.ar)
# Summary and plots
summary(post.ar)
plot(post.ar)
# Model selection criteria
post.ar$PMCC
## Fit and spatially prediction simultaneously
##
# Read data
data(DataFit);
data(DataValPred)
# Define the coordinates
```

```
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
# MCMC via Gibbs will provide output in *.txt format
# from C routine to avoide large data problem in R
set.seed(11)
post.ar.fitpred <- spT.Gibbs(formula=o8hrmax ~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="AR", coords=coords,
         newcoords=pred.coords, newdata=DataValPred,
         scale.transform="SQRT")
print(post.ar.fitpred)
summary(post.ar.fitpred)
names(post.ar.fitpred)
# validation criteria
spT.validation(DataValPred$o8hrmax,c(post.ar.fitpred$prediction[,1]))
## The GPP approximation models:
######################################
## Model fitting
##
# Read data
data(DataFit);
# Define the coordinates
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
# Define knots
knots<-spT.grid.coords(Longitude=c(max(coords[,1]),</pre>
              min(coords[,1])),Latitude=c(max(coords[,2]),
              min(coords[,2])), by=c(4,4))
# MCMC via Gibbs using default choices
set.seed(11)
post.gpp <- spT.Gibbs(formula=08hrmax ~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="GPP", coords=coords,
         knots.coords=knots, scale.transform="SQRT")
print(post.gpp)
# MCMC via Gibbs not using default choices
# define the time-series
time.data<-spT.time(t.series=60, segment=1)</pre>
# hyper-parameters for the prior distributions
priors<-spT.priors(model="GPP",var.prior=Gam(2,1),</pre>
        beta.prior=Nor(0,10<sup>4</sup>))
# initial values for the model parameters
initials<-spT.initials(model="GPP", sig2eps=0.01,</pre>
            sig2eta=0.5, beta=NULL, phi=0.001)
```

```
# input for spatial decay
#spatial.decay<-spT.decay(type="FIXED", value=0.001)</pre>
spatial.decay<-spT.decay(type="MH", tuning=0.05) #</pre>
#spatial.decay<-spT.decay(type="DISCRETE",limit=c(0.001,0.009),segments=10)</pre>
# Iterations for the MCMC algorithms
nItr<-5000
# MCMC via Gibbs
set.seed(11)
post.gpp <- spT.Gibbs(formula=08hrmax~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="GPP", time.data=time.data,
         coords=coords, knots.coords=knots,
         priors=priors, initials=initials,
         nItr=nItr, nBurn=0, report=nItr,
         tol.dist=2, \ distance.method="geodetic:km",
         cov.fnc="exponential", scale.transform="SQRT",
         spatial.decay=spatial.decay)
print(post.gpp)
# Summary and plots
summary(post.gpp)
plot(post.gpp)
# Model selection criteria
post.gpp$PMCC
## Fit and spatially prediction simultaneously
##
# Read data
data(DataFit);
data(DataValPred)
# Define the coordinates
coords<-as.matrix(unique(cbind(DataFit[,2:3])))</pre>
pred.coords<-as.matrix(unique(cbind(DataValPred[,2:3])))</pre>
knots<-spT.grid.coords(Longitude=c(max(coords[,1]),</pre>
              min(coords[,1])),Latitude=c(max(coords[,2]),
              min(coords[,2])), by=c(4,4))
# MCMC via Gibbs will provide output in *.txt format
# from C routine to avoide large data problem in R
set.seed(11)
post.gpp.fitpred <- spT.Gibbs(formula=08hrmax ~cMAXTMP+WDSP+RH,</pre>
         data=DataFit, model="GP", coords=coords, knots.coords=knots,
         newcoords=pred.coords, newdata=DataValPred,
         scale.transform="SQRT")
print(post.gpp.fitpred)
summary(post.gpp.fitpred)
plot(post.gpp.fitpred)
```

spT.grid.coords 29

```
names(post.gpp.fitpred)

# validation criteria
spT.validation(DataValPred$o8hrmax,c(post.gpp.fitpred$prediction[,1]))
##

## End(Not run)
```

spT.grid.coords

Grid Coordinates

Description

This function is used to obtain Longitude/x and Latitude/y coordinates in a grid set.

Usage

```
spT.grid.coords(Longitude = c(max, min),
    Latitude = c(max, min), by = c(NA,NA))
```

Arguments

Longitude The maximum and minimum longitude position.

Latitude The maximum and minimum latitude position.

by The number of x and y points in each axis.

See Also

```
spT.geodist.
```

spT.initials

```
## ## End(Not run)
```

spT.initials

Initial values for the spatio-temporal models.

Description

This command is useful to assign the initial values of the hyper-parameters of the prior distributions.

Usage

```
spT.initials(model, sig2eps=0.01, sig2eta=NULL,
    rho=NULL, beta=NULL, phi=NULL)
```

Arguments

model	The spatio-temporal models, current options are: "GP", "AR", and "GPP".
sig2eps	Initial value for the parameter $\sigma^2 - \epsilon$.
sig2eta	Initial value for the parameter $\sigma^2\eta$.
rho	Initial value for the parameter ρ .
beta	Initial value for the parameter β .
phi	Initial value for the parameter ϕ .

Note

Initial values are automatically given if the user does not provide these.

See Also

```
spT.Gibbs, spT.priors.
```

spT.keep.morethan.dist 31

```
spT.keep.morethan.dist
```

Present one coordinate in a defined area for presentation

Description

This function is used to present one coordinate in a defined area to avoid cutter.

Usage

```
spT.keep.morethan.dist(coords, tol.dist=100)
```

Arguments

coords X and Y axes/ longitude and latitude values.

tol.dist The tolerance limit for the distance.

See Also

```
spT.geodist, dist, spT.data.selection.
```

```
## Not run:
##

data(NYdata)
head(NYdata)
NYsite<-unique(NYdata[,2:3])
head(NYsite)
spT.keep.morethan.dist(NYsite,tol.dist=100)

# Including values
dat<-cbind(NYsite,value=rnorm(dim(NYsite)[[1]]))
head(dat)
spT.keep.morethan.dist(dat,tol.dist=100)
##

## End(Not run)</pre>
```

32 spT.pCOVER

Description

This function is used to obtain nominal coverage.

Usage

```
spT.pCOVER(z=NULL,zup=NULL,zlow=NULL,zsample=NULL,level=95)
```

Arguments

z The original values (matrix or vector).
 zup The predicted values for upper interval (matrix or vector).
 zlow The predicted values for lower interval (matrix or vector).
 zsample Predicted MCMC samples.

level Level of coverages.

See Also

```
spT.validation.
```

```
## Not run:
##

# Create 'x': the true values.
# Create 'yup': the upper interval.
# Create 'ylow': the lower interval.

x <- rnorm(1000,5,0.1)
yup <- rnorm(1000,7,2)
ylow <- rnorm(1000,3,2)

# The pCOVER is:

spT.pCOVER(z=x, zup=yup, zlow=ylow)

# create predicted MCMC samples
y <- matrix(rnorm(1000*5000,5,1),1000,5000)
# The pCOVER is:

spT.pCOVER(z=x, zsamples=y)
spT.pCOVER(z=x, zsamples=y, level=50)</pre>
```

spT.priors 33

```
## ## End(Not run)
```

spT.priors

Priors for the spatio-temporal models.

Description

This command is useful to assign the hyper-parameters of the prior distributions.

Usage

```
spT.priors(model,var.prior=Gam(a=2,b=1),
beta.prior=Nor(0,10^10), rho.prior=Nor(0,10^10),
phi.prior=Gam(a=2,b=1))
```

Arguments

model	The spatio-temporal models, current input: "GP", "AR", and "GPP".
var.prior	The hyper-parameter for the Gamma prior distribution (with mean = a/b) of the variance model parameters (e.g., $\sigma 2_{\epsilon}$, $\sigma 2_{\eta}$).
beta.prior	The hyper-parameter for the Normal prior distribution of the β model parameters.
rho.prior	The hyper-parameter for the Normal prior distribution of the ρ model parameter.
phi.prior	The hyper-parameter for the Gamma prior distribution (with mean = a/b) of the spatial decay (ϕ) parameter.

Note

If no prior information are given (assigned as NULL), then it use flat prior values of the corresponding distributions.

Gam and Nor refers to Gamma and Normal distributions respectively.

See Also

```
spT.Gibbs, predict.spT, spT.initials.
```

34 spT.segment.plot

```
## ## End(Not run)
```

spT.segment.plot

Utility plot for prediction/forecast

Description

This function is used to obtain scatter plots with 95 percent CI for predictions/forecasts.

Usage

```
spT.segment.plot(obs, est, up, low, limit = NULL)
```

Arguments

obs	Observed values.
est	Estimated values.
up	Upper limit of the estimated values.
low	Lower limit of the estimated values.
limit	x-axis and y-axis limits.

See Also

```
summary.spT, plot.spT.
```

```
## Not run:
##

obs<-rnorm(10,15,1)
est<-rnorm(10,15,1.5)
up<-rnorm(10,25,0.5)
low<-rnorm(10,5,0.5)
spT.segment.plot(obs,est,up,low,limit=c(0,30))
##

## End(Not run)</pre>
```

spT.time 35

spT.time

Timer series information.

Description

This function defines the time series in the spatio-temporal data.

Usage

```
spT.time(t.series, segment=1)
```

Arguments

t.series

Number of times within each segment in each series. This should be a constant. Currently it is not possible to make it a variable? Like its 30 for April, and 31

for May etc.

segment

Number of segments in each time series. This should be a constant.

See Also

```
spT.Gibbs.
```

Examples

```
## Not run:
##

time.data<-spT.time(t.series=31,segment=2)
##

## End(Not run)</pre>
```

spT.validation

Validation Commands

Description

The following function is used to validate the predicted observations with the actual values.

Usage

```
spT.validation(z, zhat)
```

36 summary.spT

Arguments

z The original values (matrix or vector). zhat The predicted values (matrix or vector).

Value

MSE Mean Squared Error.

RMSE Root Mean Squared Error.

MAE Mean Absolute Error.

MAPE Mean Absolute Percentage Error.

BIAS Bias.

rBIAS Relative Bias.

rMSEP Relative Mean Separation.

See Also

```
spT.pCOVER.
```

Examples

```
## Not run:
##

# Create 'x', which is the true values.
# Create 'y', which is the predicted values.

x <- rnorm(10,5,0.1)
y <- rnorm(10,5,1)
spT.validation(x, y)
##

## End(Not run)</pre>
```

summary.spT

Summary statistics of the parameters.

Description

This function is used to obtain MCMC summary statistics.

Usage

```
## S3 method for class 'spT'
## S3 method for class 'spT'
summary(object, digits=4, package="spTimer", ...)
##
```

summary.spT 37

Arguments

object Object of class inheriting from "spT".

digits Rounds the specified number of decimal places (default 4).

package If "coda" then summary statistics are given using coda package. Defaults value

is "spTimer".

.. Other arguments.

Value

sig2eps Summary statistics for σ_{ϵ}^2 . sig2eta Summary statistics for σ_{η}^2 .

phi Summary statistics for spatial decay parameter ϕ , if estimated using spT. decay.

. . . Summary statistics for other parameters used in the models.

See Also

spT.Gibbs.

```
## Not run:
##
summary(out) # where out is the output from spT class
summary(out, digits=2) # where out is the output from spT class
summary(out, pack="coda") # where out is the output from spT class
##
## End(Not run)
```

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