Package 'spatialSPsurv'

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Type Package
Title Bayesian Spatial Split Population Survival Model
Version 0.1.0.9000
Description Contains functions to fit Bayesian spatial survival model for split population.
License MIT + file LICENSE
Encoding UTF-8
LazyData true
RoxygenNote 7.1.0
LinkingTo Rcpp, RcppArmadillo
Imports MCMCpack, FastGP, stats, Rcpp, RcppArmadillo, coda
R topics documented:
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Description

Markov Chain Monte Carlo (MCMC) to run Bayesian non-spatial frailty split population survival model

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Usage

```
frailtySPsurv(
  formula,
  duration,
  immune,
  Y0,
  LY,
  S,
  data = list(),
  N,
  burn,
  thin,
  w = c(1, 1, 1),
  m = 10,
  form,
  prop.var
)
```

Arguments

formula duration immune the elapsed time since inception until the beginning of time period (t-1) Υ0 LY last observation year S spatial information (e.g. district ID) for each observation that matches the spatial matrix row/column information data number of MCMC iterations Ν burn burn-in to be discarded thin thinning to prevent from autocorrelation size of the slice in the slice sampling for (betas, gammas, rho). Write it as a vector. E.g. c(1,1,1)limit on steps in the slice sampling. A vector of values for beta, gamma, rho. type of parametric model (Exponential or Weibull) form

Value

prop.var

chain of the variables of interest

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Description

Markov Chain Monte Carlo (MCMC) to run Bayesian spatial split population survival model

Usage

```
spatialSPsurv(
   duration,
   immune,
   Y0,
   LY,
   S,
   data = list(),
   A,
   N,
   burn,
   thin,
   w = c(1, 1, 1),
   m = 10,
   form,
   prop.var
)
```

Arguments

duration	
immune	
Y0	the elapsed time since inception until the beginning of time period (t-1)
LY	last observation year
S	spatial information (e.g. district ID) for each observation that matches the spatial matrix row/column information $\frac{1}{2}$
data	
A	Spatial Matrix (load separate spatial weights matrix file)
N	number of MCMC iterations
burn	burn-in to be discarded
thin	thinning to prevent from autocorrelation
W	size of the slice in the slice sampling for (betas, gammas, rho). Write it as a vector. E.g. $c(1,1,1)$
m	limit on steps in the slice sampling. A vector of values for beta, gamma, rho.
form	type of parametric model (Exponential or Weibull)
prop.var	proposal variance for Metropolis-Hastings

Value

chain of the variables of interest

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Description

Markov Chain Monte Carlo (MCMC) to run Bayesian split population survival model with no frailties

Usage

```
SPsurv(
   duration,
   immune,
   Y0,
   LY,
   data = list(),
   N,
   burn,
   thin,
   w = c(1, 1, 1),
   m = 10,
   form
)
```

Arguments

```
duration
immune
                   ...
Υ0
                   the elapsed time since inception until the beginning of time period (t-1)
LY
                   last observation year
data
                   number of MCMC iterations
Ν
                   burn-in to be discarded
burn
thin
                   thinning to prevent from autocorrelation
                   size of the slice in the slice sampling for (betas, gammas, rho). Write it as a
                   vector. E.g. c(1,1,1)
                   limit on steps in the slice sampling. A vector of values for beta, gamma, rho.
m
                   type of parametric model (Exponential or Weibull)
form
```

Value

chain of the variables of interest

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