

Package ‘spatsurv’

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Type Package

Title Bayesian Spatial Survival Analysis with Parametric Proportional Hazards Models

Version 1.6

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Additional contributions
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Description Bayesian inference for parametric proportional hazards spatial survival models; flexible spatial survival models. See Benjamin M. Taylor, Barry S. Rowlingson (2017) <doi:10.18637/jss.v077.i04>.

License GPL-3

Imports survival, sp, spatstat.geom, spatstat.core, spatstat (>= 2.0-0), raster, iterators, RandomFields, fields, Matrix, stringr, rgeos, RColorBrewer, methods, lubridate

Suggests rgdal, rgl, OpenStreetMap

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spatsurv-package	<i>spatsurv</i>
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Description

An R package for spatially correlated parametric proportional hazards survival analysis.

Usage

spatsurv

Format

An object of class `logical` of length 1.

Details

Package: spatsurv
 Type: Package
 Title: Bayesian Spatial Survival Analysis with Parametric Proportional Hazards Models
 Version: 1.6
 Date: 2021-03-26
 Author: Benjamin M. Taylor and Barry S. Rowlingson Additional contributions Ziyu Zheng
 Maintainer: Benjamin M. Taylor <benjamin.taylor.software@gmail.com>
 Description: Bayesian inference for parametric proportional hazards spatial survival models; flexible spatial survival
 License: GPL-3
 Imports: survival, sp, spatstat.geom, spatstat.core, spatstat (>= 2.0-0), raster, iterators, RandomFields, fields, Mat
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 Encoding: UTF-8
 RoxygenNote: 7.1.1
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 Depends: R (>= 2.10)
 Repository: CRAN
 Date/Publication: 2020-04-26 07:00:02 UTC

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urlTemplate	urlTemplate function
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vcov.mlspatsurv	vcov.mlspatsurv function
weibullHaz	weibullHaz function

sectionDependencies The package spatsurv depends upon some other important contributions to CRAN in order to operate; their uses here are indicated:

survival, sp, spatstat, raster, iterators, RandomFields, fields, rgl, Matrix, stringr, RColorBrewer, geostatsp, rgeos.

sectionCitation To cite use of spatsurv, the user may refer to the following work:

Benjamin M. Taylor and Barry S. Rowlingson (2017).
 spatsurv: An R Package for Bayesian Inference with Spatial Survival Models.
 Journal of Statistical Software, 77(4), 1-32, doi:10.18637/jss.v077.i04.

references X

Author(s)

Benjamin Taylor, Health and Medicine, Lancaster University, Barry Rowlingson, Health and Medicine, Lancaster University

Description

A function to print a welcome message on loading package

Usage

```
.onAttach(libname, pkgname)
```

Arguments

libname	libname argument
pkgname	pkgname argument

Value

...

allocate	<i>allocate function</i>
----------	--------------------------

Description

A function to allocate coordinates to an observation whose spatial location is known to the regional level

Usage

```
allocate(poly, popden, survdat, pid, sid, n = 2, wid = 2000)
```

Arguments

poly	a SpatialPolygonsDataFrame, on which the survival data exist in aggregate form
popden	a sub-polygon raster image of population density
survdat	data.frame containing the survival data
pid	name of the variable in the survival data that gives the region identifier in poly
sid	the name of the variable in poly to match the region identifier in survdat to
n	the number of different allocations to make. e.g. if n is 2 (the default) two candidate sets of locations are available.
wid	The default is 2000, interpreted in metres ie 2Km. size of buffer to add to window for raster cropping purposes: this ensures that for each polygon, the cropped raster covers it completely.

Value

matrices x and y, both of size (number of observations in survdat x n) giving n potential candidate locations of points in the columns of x and y.

alpha	<i>alpha function</i>
-------	-----------------------

Description

A function used in calculating the coefficients of a B-spline curve

Usage

alpha(i, j, knots, knotidx)

Arguments

i	index i
j	index j
knots	knot vector
knotidx	knot index

Value

a vector

B	<i>B function</i>
---	-------------------

Description

A recursive function used in calculating the coefficients of a B-spline curve

Usage

B(x, i, j, knots)

Arguments

x	locations at which to evaluate the B-spline
i	index i
j	index j
knots	a knot vector

Value

a vector of polynomial coefficients

basehazard	<i>basehazard function</i>
------------	----------------------------

Description

Generic function for computing the baseline hazard

Usage

```
basehazard(obj, ...)
```

Arguments

obj	an object
...	additional arguments – currently there are none, but this is for extensibility

Value

method basehazard

See Also

[basehazard.basehazardspec](#), [exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

basehazard.basehazardspec	<i>basehazard.basehazardspec function</i>
---------------------------	---

Description

A function to retrieve the baseline hazard function

Usage

```
## S3 method for class 'basehazardspec'  
basehazard(obj, ...)
```

Arguments

obj	an object of class basehazardspec
...	additional arguments – currently there are none, but this is for extensibility

Value

a function returning the baseline hazard

See Also

[exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

baseHazST

baseHazST function

Description

A function to

Usage

```
baseHazST(  
  bh1 = NULL,  
  survobj,  
  t0,  
  nbreaks = 5,  
  breakmethod = "quantile",  
  MLimits = NULL  
)
```

Arguments

bh1	X
survobj	X
t0	X
nbreaks	X
breakmethod	X
MLimits	X

Value

...

baselinehazard	<i>baselinehazard function</i>
----------------	--------------------------------

Description

A function to compute quantiles of the posterior baseline hazard or cumulative baseline hazard.

Usage

```
baselinehazard(
  x,
  t = NULL,
  n = 100,
  probs = c(0.025, 0.5, 0.975),
  cumulative = FALSE,
  plot = TRUE,
  bw = FALSE,
  ...
)
```

Arguments

<code>x</code>	an object inheriting class <code>mcmcspatsurv</code>
<code>t</code>	optional vector of times at which to compute the quantiles, Default is <code>NULL</code> , in which case a uniformly spaced vector of length <code>n</code> from 0 to the maximum time is used
<code>n</code>	the number of points at which to compute the quantiles if <code>t</code> is <code>NULL</code>
<code>probs</code>	vector of probabilities
<code>cumulative</code>	logical, whether to return the baseline hazard (default i.e. <code>FALSE</code>) or cumulative baseline hazard
<code>plot</code>	whether to plot the result
<code>bw</code>	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
<code>...</code>	additional arguments to be passed to <code>plot</code>

Value

the vector of times and quantiles of the baseline or cumulative baseline hazard at those times

See Also

[print.mcmcspatsurv](#), [quantile.mcmcspatsurv](#), [summary.mcmcspatsurv](#), [vcov.mcmcspatsurv](#), [frailty-lag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [predict.mcmcspatsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

baselinehazard_multiWay

baselinehazard_multiWay function

Description

A function to

Usage

```
baselinehazard_multiWay(  
  x,  
  probs = c(0.025, 0.5, 0.975),  
  cumulative = FALSE,  
  plot = TRUE,  
  joint = FALSE,  
  xlims = NULL,  
  ylims = NULL,  
  ...  
)
```

Arguments

x	X
probs	X
cumulative	X
plot	X
joint	X
xlims	X
ylims	X
...	X

Value

...

betapriorGauss	<i>betapriorGauss function</i>
----------------	--------------------------------

Description

A function to define Gaussian priors for beta. This function simply stores a vector of means and standard deviations to be passed to the main MCMC function, survspat.

Usage

```
betapriorGauss(mean, sd)
```

Arguments

mean	the prior mean, a vector of length 1 or more. 1 implies a common mean.
sd	the prior standard deviation, a vector of length 1 or more. 1 implies a common standard deviation.

Value

an object of class "betapriorGauss"

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

blockDiag	<i>A function to</i>
-----------	----------------------

Description

A function to

Usage

```
blockDiag(matlist)
```

Arguments

matlist	X
---------	---

Value

...

boxplotRisk	<i>boxplotRisk function</i>
-------------	-----------------------------

Description

A function to

Usage

```
boxplotRisk(g2r)
```

Arguments

g2r	X
-----	---

Value

...

Bspline.construct	<i>Bspline.construct function</i>
-------------------	-----------------------------------

Description

A function to construct a B-spline basis matrix for given data and basis coefficients. Used in evaluating the baseline hazard.

Usage

```
Bspline.construct(x, basis)
```

Arguments

x	a vector, the data
basis	an object created by the getBbasis function

Value

a basis matrix

BsplineHaz

*BsplineHaz function***Description**

A function to define a parametric proportional hazards model where the baseline hazard is modelled by a basis spline. This function returns an object inheriting class 'basehazardspec', list of functions 'distinfo', 'basehazard', 'gradbasehazard', 'hessbasehazard', 'cumbasehazard', 'gradcumbasehazard', 'hesscumbasehazard' and 'densityquantile'

Usage

```
BsplineHaz(times, knots = quantile(times), degree = 3, MLinits = NULL)
```

Arguments

times	vector of survival times (both censored and uncensored)
knots	vector of knots in ascending order, must include minimum and maximum values of 'times'
degree	degree of the spline basis, default is 3
MLinits	optional starting values for the non-spatial maximisation routine using optim. Note that we are working with the log of the parameters. Default is -10 for each parameter.

Details

The `distinfo` function is used to provide basic distribution specific information to other `spatsurv` functions. The user is required to provide the following information in the returned list: `npars`, the number of parameters in this distribution; `parnames`, the names of the parameters; `trans`, the transformation scale on which the priors will be provided; `itrans`, the inverse transformation function that will be applied to the parameters before the hazard, and other functions are evaluated; `jacobian`, the derivative of the inverse transformation function with respect to each of the parameters; and `hessian`, the second derivatives of the inverse transformation function with respect to each of the parameters – note that currently the package `spatsurv` only allows the use of functions where the parameters are transformed independently.

The `basehazard` function is used to evaluate the baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradbasehazard` function is used to evaluate the gradient of the baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hessbasehazard` function is used to evaluate the Hessian of the baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `cumbasehazard` function is used to evaluate the cumulative baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradcumbasehazard` function is used to evaluate the gradient of the cumulative baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hesscumbasehazard` function is used to evaluate the Hessian of the cumulative baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `densityquantile` function is used to return quantiles of the density function. This is NOT REQUIRED for running the MCMC, merely for us in post-processing with the `predict` function where `type` is `'densityquantile'`. In the case of the Weibull model for the baseline hazard, it can be shown that the `q`-th quantile is:

Value

an object inheriting class `'basehazardspec'`

See Also

[exponentialHaz](#), [gompertzHaz](#), [makehamHaz](#), [weibullHaz](#)

checkSurvivalData	<i>checkSurvivalData function</i>
-------------------	-----------------------------------

Description

A function to check whether the survival data to be passed to `survspat` is in the correct format

Usage

```
checkSurvivalData(s)
```

Arguments

`s` an object of class `Surv`, from the `survival` package

Value

if there are any issues with data format, these are returned with the data an error message explaining any issues with the data

circulant	<i>circulant function</i>
-----------	---------------------------

Description

generic function for constructing circulant matrices

Usage

```
circulant(x, ...)
```

Arguments

x	an object
...	additional arguments

Value

method circulant

circulant.matrix	<i>circulant.matrix function</i>
------------------	----------------------------------

Description

If x is a matrix whose columns are the bases of the sub-blocks of a block circulant matrix, then this function returns the block circulant matrix of interest.

Usage

```
## S3 method for class 'matrix'
circulant(x, ...)
```

Arguments

x	a matrix object
...	additional arguments

Value

If x is a matrix whose columns are the bases of the sub-blocks of a block circulant matrix, then this function returns the block circulant matrix of interest.

circulant.numeric *circulant.numeric function*

Description

returns a circulant matrix with base x

Usage

```
## S3 method for class 'numeric'
circulant(x, ...)
```

Arguments

x an numeric object
 ... additional arguments

Value

a circulant matrix with base x

circulantij *circulantij function*

Description

A function to return the "idx" i.e. c(i,j) element of a circulant matrix with base "base".

Usage

```
circulantij(idx, base)
```

Arguments

idx vector of length 2 th (i,j) (row,column) index to return
 base the base matrix of a circulant matrix

Value

the ij element of the full circulant

covmodel	<i>covmodel function</i>
----------	--------------------------

Description

A function to define the spatial covariance model, see also ?CovarianceFct. Note that the parameters defined by the 'pars' argument are fixed, i.e. not estimated by the MCMC algorithm. To have spatsurv estimate these parameters, the user must construct a new covariance function to do so, see the spatsurv vignette.

Usage

```
covmodel(model, pars)
```

Arguments

model	correlation type, a string see ?CovarianceFct
pars	vector of additional parameters for certain classes of covariance function (eg Matern), these must be supplied in the order given in ?CovarianceFct and are not estimated

Value

an object of class covmodel

See Also

CovarianceFct

CSplot	<i>CSplot function</i>
--------	------------------------

Description

A function to produce a diagnostic plot for model fit using the Cox-Snell residuals.

Usage

```
CSplot(mod, plot = TRUE, bw = FALSE, ...)
```

Arguments

mod	an object produced by the function survspat
plot	whether to plot the result, default is TRUE
bw	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
...	other arguments to pass to plot

Value

the x and y values used in the plot

cumbasehazard	<i>cumbasehazard function</i>
---------------	-------------------------------

Description

Generic function for computing the cumulative baseline hazard

Usage

```
cumbasehazard(obj, ...)
```

Arguments

obj	an object
...	additional arguments – currently there are none, but this is for extensibility

Value

method cumbasehazard

See Also

[cumbasehazard.basehazardspec](#), [exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpow-Haz](#)

cumbasehazard.basehazardspec	<i>cumbasehazard.basehazardspec function</i>
------------------------------	--

Description

A function to retrieve the cumulative baseline hazard function

Usage

```
## S3 method for class 'basehazardspec'
cumbasehazard(obj, ...)
```

Arguments

obj	an object of class basehazardspec
...	additional arguments – currently there are none, but this is for extensibility

Value

a function returning the cumulative baseline hazard

See Also

[exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

cumulativeBspline.construct

cumulativeBspline.construct function

Description

A function to construct the integral of a B-spline curve given data and basis coefficients. Used in evaluating the cumulative baseline hazard.

Usage

```
cumulativeBspline.construct(x, basis)
```

Arguments

x	a vector, the data
basis	an object created by the getBbasis function

Value

an object that allows the integral of a given B-spline curve to be computed

densityquantile

densityquantile function

Description

Generic function for computing quantiles of the density function for a given baseline hazard. This may not be analytically tractable.

Usage

```
densityquantile(obj, ...)
```

Arguments

obj	an object
...	additional arguments – currently there are none, but this is for extensibility

Value

method densityquantile

See Also

[densityquantile.basehazardspec](#), [exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

densityquantile.basehazardspec

densityquantile.basehazardspec function

Description

A function to retrieve the quantiles of the density function

Usage

```
## S3 method for class 'basehazardspec'  
densityquantile(obj, ...)
```

Arguments

obj	an object of class basehazardspec
...	additional arguments – currently there are none, but this is for extensibility

Value

a function returning the density quantiles

See Also

[exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

densityquantile_PP *densityquantile_PP function*

Description

A function to compute quantiles of the density function

Usage

```
densityquantile_PP(inputs)
```

Arguments

inputs	inputs for the function including the model matrix, frailties, fixed effects and the parameters of the baseline hazard derived from this model
--------	--

Value

quantiles of the density function for the individual

density_PP *density_PP function*

Description

A function to compute an individual's density function

Usage

```
density_PP(inputs)
```

Arguments

inputs	inputs for the function including the model matrix, frailties, fixed effects and the parameters of the baseline hazard derived from this model
--------	--

Value

the density function for the individual

derivindepGaussianprior

derivindepGaussianprior function

Description

A function for evaluating the first and second derivatives of the log of an independent Gaussian prior

Usage

```
derivindepGaussianprior(beta = NULL, omega = NULL, eta = NULL, priors)
```

Arguments

beta	a vector, the parameter beta
omega	a vector, the parameter omega
eta	a vector, the parameter eta
priors	an object of class 'mcmcPrior', see ?mcmcPrior

Value

returns the first and second derivatives of the prior

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

derivindepGaussianpriorST

derivindepGaussianpriorST function

Description

A function to

Usage

```
derivindepGaussianpriorST(beta = NULL, omega = NULL, eta = NULL, priors)
```

Arguments

beta	X
omega	X
eta	X
priors	X

Value

...

derivpsplineprior *derivpsplineprior function*

Description

A function for evaluating the first and second derivatives of the log of an independent Gaussian prior

Usage

```
derivpsplineprior(beta = NULL, omega = NULL, eta = NULL, priors)
```

Arguments

beta	a vector, the parameter beta
omega	a vector, the parameter omega
eta	a vector, the parameter eta
priors	an object of class 'mcmcPrior', see ?mcmcPrior

Value

returns the first and second derivatives of the prior

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

distinfo	<i>distinfo function</i>
----------	--------------------------

Description

Generic function for returning information about the class of baseline hazard functions employed.

Usage

```
distinfo(obj, ...)
```

Arguments

obj	an object
...	additional argument – currently there are none, but this is for extensibility

Value

method distinfo

See Also

[distinfo.basehazardspec](#), [exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

distinfo.basehazardspec	<i>distinfo.basehazardspec function</i>
-------------------------	---

Description

A function to retrieve information on the baseline hazard distribution of choice

Usage

```
## S3 method for class 'basehazardspec'  
distinfo(obj, ...)
```

Arguments

obj	an object of class basehazardspec
...	additional arguments – currently there are none, but this is for extensibility

Value

a function returning information on the baseline hazard distribution of choice

See Also

[exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

estimateY	<i>estimateY function</i>
-----------	---------------------------

Description

A function to get an initial estimate of Y, to be used in calibrating the MCMC. Not for general use

Usage

```
estimateY(X, betahat, omegahat, surv, control)
```

Arguments

X	the design matrix containing covariate information
betahat	an estimate of beta
omegahat	an estimate of omega
surv	an object of class Surv
control	a list containing various control parameters for the MCMC and post-processing routines

Value

an estimate of Y, to be used in calibrating the MCMC

etapriorGauss	<i>etapriorGauss function</i>
---------------	-------------------------------

Description

A function to define Gaussian priors for eta. This function simply stores a vector of means and standard deviations to be passed to the main MCMC function, survspat.

Usage

```
etapriorGauss(mean, sd)
```

Arguments

mean	the prior mean, a vector of length 1 or more. 1 implies a common mean.
sd	the prior standard deviation, a vector of length 1 or more. 1 implies a common standard deviation.

Value

an object of class "etapriorGauss"

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

Et_PP

Et_PP function

Description

A function to compute an individual's approximate expected survival time using numerical integration. Note this appears to be unstable; the function is based on R's integrate function. Not intended for general use (yet!).

Usage

Et_PP(inputs)

Arguments

inputs inputs for the function including the model matrix, frailties, fixed effects and the parameters of the baseline hazard derived from this model

Value

the expected survival time for the individual, obtained by numerical integration of the density function.

EvalCov

EvalCov function

Description

This function is used to evaluate the covariance function within the MCMC run. Not intended for general use.

Usage

EvalCov(cov.model, u, parameters)

Arguments

cov.model	an object of class covmodel
u	vector of distances
parameters	vector of parameters

Value

method EvalCov

ExponentialCovFct *ExponentialCovFct function*

Description

A function to declare and also evaluate an exponential covariance function.

Usage

```
ExponentialCovFct()
```

Value

the exponential covariance function

See Also

[SpikedExponentialCovFct](#), [covmodel](#), [CovarianceFct](#)

exponentialHaz *exponentialHaz function*

Description

A function to define a parametric proportional hazards model where the baseline hazard is taken from the exponential model. This function returns an object inheriting class 'basehazardspec', list of functions 'distinfo', 'basehazard', 'gradbasehazard', 'hessbasehazard', 'cumbasehazard', 'gradcumbasehazard', 'hesscumbasehazard' and 'densityquantile'

Usage

```
exponentialHaz()
```

Details

The `distinfo` function is used to provide basic distribution specific information to other `spatsurv` functions. The user is required to provide the following information in the returned list: `npars`, the number of parameters in this distribution; `parnames`, the names of the parameters; `trans`, the transformation scale on which the priors will be provided; `itrans`, the inverse transformation function that will be applied to the parameters before the hazard, and other functions are evaluated; `jacobian`, the derivative of the inverse transformation function with respect to each of the parameters; and `hessian`, the second derivatives of the inverse transformation function with respect to each of the parameters – note that currently the package `spatsurv` only allows the use of functions where the parameters are transformed independently.

The `basehazard` function is used to evaluate the baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradbasehazard` function is used to evaluate the gradient of the baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hessbasehazard` function is used to evaluate the Hessian of the baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `cumbasehazard` function is used to evaluate the cumulative baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradcumbasehazard` function is used to evaluate the gradient of the cumulative baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hesscumbasehazard` function is used to evaluate the Hessian of the cumulative baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `densityquantile` function is used to return quantiles of the density function. This is NOT REQUIRED for running the MCMC, merely for us in post-processing with the `predict` function where `type` is `'densityquantile'`. In the case of the Weibull model for the baseline hazard, it can be shown that the `q`-th quantile is:

Value

an object inheriting class `'basehazardspec'`

See Also

[tpowHaz](#), [gompertzHaz](#), [makehamHaz](#), [weibullHaz](#)

FFTgrid	<i>FFTgrid function</i>
---------	-------------------------

Description

A function to generate an FFT grid and associated quantities including cell dimensions, size of extended grid, centroids,

Usage

```
FFTgrid(spatialdata, cellwidth, ext, boundingbox = NULL)
```

Arguments

spatialdata	a SpatialPixelsDataFrame object
cellwidth	width of computational cells
ext	multiplying constant: the size of the extended grid: ext*M by ext*N
boundingbox	optional bounding box over which to construct computational grid, supplied as an object on which the function 'bbox' returns the bounding box

Value

a list

fixedpars	<i>fixedpars function</i>
-----------	---------------------------

Description

A function to return the mcmc chains for the covariate effects

Usage

```
fixedpars(x)
```

Arguments

x	an object of class mcmcspsurv
---	-------------------------------

Value

the beta mcmc chains

See Also

[print.mcmcspatsurv](#), [quantile.mcmcspatsurv](#), [summary.mcmcspatsurv](#), [vcov.mcmcspatsurv](#), [frailty-lag1](#), [spatialpars](#), [hazardpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspatsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

fixmatrix	<i>fixmatrix function</i>
-----------	---------------------------

Description

!! THIS FUNCTION IS NOT INTENDED FOR GENERAL USE !!

Usage

```
fixmatrix(mat)
```

Arguments

mat	a matrix
-----	----------

Details

A function to fix up an estimated covariance matrix using a VERY ad-hoc method.

Value

the fixed matrix

fixParHaz	<i>fixParHaz function</i>
-----------	---------------------------

Description

A function to

Usage

```
fixParHaz(bh, idx, fixval)
```

Arguments

bh	X
idx	X
fixval	X

Value

...

frailtylag1	<i>frailtylag1 function</i>
-------------	-----------------------------

Description

A function to produce a plot of, and return, the lag 1 (or higher, see argument 'lag') autocorrelation for each of the spatially correlated frailty chains

Usage

```
frailtylag1(object, plot = TRUE, lag = 1, ...)
```

Arguments

object	an object inheriting class <code>mcmcspsurv</code>
plot	logical whether to plot the result, default is TRUE
lag	the lag to plot, the default is 1
...	other arguments to be passed to the plot function

Value

the lag 1 autocorrelation for each of the spatially correlated frailty chains

See Also

[print.mcmcspsurv](#), [quantile.mcmcspsurv](#), [summary.mcmcspsurv](#), [vcov.mcmcspsurv](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

fs	<i>London Fire Brigade property</i>
----	-------------------------------------

Description

London Fire Brigade property

Usage

```
data(fs)
```

Format

```
data.frame
```

Source

<https://data.london.gov.uk/>

References

<https://data.london.gov.uk/>, <http://www.nationalarchives.gov.uk/doc/open-government-licence/version/3/>

Examples

```
fire <- data(fs)
```

fstimes

London Fire Brigade response times to dwelling fires, 2009

Description

London Fire Brigade response times to dwelling fires, 2009

Usage

```
data(fstimes)
```

Format

data.frame

Source

<https://data.london.gov.uk/>

References

<https://data.london.gov.uk/>, <http://www.nationalarchives.gov.uk/doc/open-government-licence/version/3/>

Examples

```
firetimes <- data(fstimes)
```

gamma2risk	<i>gamma2risk function</i>
------------	----------------------------

Description

A function to

Usage

```
gamma2risk(mod)
```

Arguments

mod	X
-----	---

Value

...

GammafromY	<i>GammafromY function</i>
------------	----------------------------

Description

A function to change Ys (spatially correlated noise) into Gammas (white noise). Used in the MALA algorithm.

Usage

```
GammafromY(Y, rootQeigs, mu)
```

Arguments

Y	Y matrix
rootQeigs	square root of the eigenvectors of the precision matrix
mu	parameter of the latent Gaussian field

Value

Gamma

GammaFromY_SPDE *GammaFromY_SPDE function*

Description

A function to go from Y to Gamma

Usage

```
GammaFromY_SPDE(Y, U, mu)
```

Arguments

Y	Y
U	upper Cholesky matrix
mu	the mean

Value

the value of Gamma for the given Y

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.
2. Finn Lindgren, Havard Rue, Johan Lindstrom. An explicit link between Gaussian fields and Gaussian Markov random fields: the stochastic partial differential equation approach. *Journal of the Royal Statistical Society: Series B* 73(4)

gencens *gencens function*

Description

A function to generate observed times given a vector of true survival times and a vector of censoring times. Used in the simulation of survival data.

Usage

```
gencens(survtimes, censtimes, type = "right")
```


Arguments

<code>survtimes</code>	a vector of survival times
<code>censtimes</code>	a vector of censoring times for left or right censored data, 2-column matrix of censoring times for interval censoring (number of rows equal to the number of observations).
<code>type</code>	the type of censoring to generate can be 'right' (default), 'left' or 'interval'

Value

an object of class 'Surv', the censoring indicator is equal to 1 if the event is uncensored and 0 otherwise for right/left censored data, or for interval censored data, the indicator is 0 uncensored, 1 right censored, 2 left censored, or 3 interval censored.

<code>getBackground</code>	<i>getBackground function</i>
----------------------------	-------------------------------

Description

A function to

Usage

```
getBackground(poly, type = "stamen-toner")
```

Arguments

<code>poly</code>	a spatial object that can be transformed and the extent obtained using the <code>bbox</code> function.
<code>type</code>	see <code>?openmap</code>

Value

...

getbb	<i>getbb function</i>
-------	-----------------------

Description

A function to get the bounding box of a Spatial object

Usage

```
getbb(obj)
```

Arguments

obj	a spatial object e.g. a SpatialPolygonsDataFrame, SpatialPolygons, etc ... anything with a bounding box that can be computed with bbox(obj)
-----	---

Value

a SpatialPolygons object: the bounding box

getBbasis	<i>getBbasis function</i>
-----------	---------------------------

Description

A function returning the piecewise polynomial coefficients for a B-spline basis function i.e. the basis functions.

Usage

```
getBbasis(x, knots, degree, force = FALSE)
```

Arguments

x	a vector of data
knots	a vector of knots in ascending order. The first and last knots must be respectively the minimum and maximum of x.
degree	the degree of the spline
force	logical: skip check on knots? (not recommended!)

Value

the knots and the piecewise polynomial coefficients for a B-spline basis function i.e. the basis functions.

getcov	<i>getcov function</i>
--------	------------------------

Description

A function to return the covariance from a model based on the randomFields covariance functions. Not intended for general use.

Usage

```
getcov(u, sigma, phi, model, pars)
```

Arguments

u	distance
sigma	variance parameter
phi	scale parameter
model	correlation type, see ?CovarianceFct
pars	vector of additional parameters for certain classes of covariance function (eg Matern), these must be supplied in the order given in ?CovarianceFct and are not estimated

Value

this is just a wrapper for CovarianceFct

getgrd	<i>getgrd function</i>
--------	------------------------

Description

A function to create a regular grid over an observation window in order to model the spatial random effects as a Gaussian Markov random field.

Usage

```
getgrd(shape, cellwidth)
```

Arguments

shape	an object of class SpatialPolygons or SpatialPolygonsDataFrame
cellwidth	a scalar, the width of the grid cells

Value

a SpatialPolygons object: the grid on which prediction of the spatial effects will occur

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.
2. Finn Lindgren, Havard Rue, Johan Lindstrom. An explicit link between Gaussian fields and Gaussian Markov random fields: the stochastic partial differential equation approach. *Journal of the Royal Statistical Society: Series B* 73(4)

getGrid	<i>getGrid function</i>
---------	-------------------------

Description

A function to extract and return the computational grid from a gridded analysis.

Usage

```
getGrid(mod, returnclass = "SpatialPolygonsDataFrame")
```

Arguments

mod	an object of class mcmcspatsurv, returned by the function survspat
returnclass	the class of object to return, default is a 'SpatialPolygonsDataFrame'. Other options are 'raster', which returns a raster brick; or 'SpatialPixelsDataFrame'

Value

a SpatialPolygonsDataFrame in which Monte Carlo expectations can be stored and later plotted.

getleneta	<i>getleneta function</i>
-----------	---------------------------

Description

A function to compute the length of eta

Usage

```
getleneta(cov.model)
```

Arguments

cov.model	a covariance model
-----------	--------------------

Value

the length of eta

getOptCellwidth	<i>getOptCellwidth function</i>
-----------------	---------------------------------

Description

A function to compute an optimal cellwidth close to an initial suggestion. This maximises the efficiency of the MCMC algorithm when in the control argument of the function `survspat`, the option `gridded` is set to `TRUE`

Usage

```
getOptCellwidth(dat, cellwidth, ext = 2, plot = TRUE, boundingbox = NULL)
```

Arguments

<code>dat</code>	any spatial data object whose bounding box can be computed using the function <code>bbox</code> .
<code>cellwidth</code>	an initial suggested cellwidth
<code>ext</code>	the extension parameter for the FFT transform, set to 2 by default
<code>plot</code>	whether to plot the grid and data to illustrate the optimal grid
<code>boundingbox</code>	optional bounding box over which to construct computational grid, supplied as an object on which the function <code>'bbox'</code> returns the bounding box

Value

the optimum cell width

getparranges	<i>getparranges function</i>
--------------	------------------------------

Description

A function to extract parameter ranges for creating a grid on which to evaluate the log-posterior, used in calibrating the MCMC. This function is not intended for general use.

Usage

```
getparranges(priors, leneta, mult = 1.96)
```

Arguments

priors	an object of class mcmcPriors
leneta	the length of eta passed to the function
mult	defaults to 1.96 so the grid formed will be mean plus/minus 1.96 times the standard deviation

Value

an appropriate range used to calibrate the MCMC: the mean of the prior for eta plus/minus 1.96 times the standard deviation

getsurvdata	<i>getsurvdata function</i>
-------------	-----------------------------

Description

A function to return the survival data from an object of class mcmcsurv. This function is not intended for general use.

Usage

```
getsurvdata(x)
```

Arguments

x	an object of class mcmcsurv
---	-----------------------------

Value

the survival data from an object of class mcmcsurv

gompertzHaz	<i>gompertzHaz function</i>
-------------	-----------------------------

Description

A function to define a parametric proportional hazards model where the baseline hazard is taken from a Gompertz model. This function returns an object inheriting class 'basehazardspec', list of functions 'distinfo', 'basehazard', 'gradbasehazard', 'hessbasehazard', 'cumbasehazard', 'gradcumbasehazard', 'hesscumbasehazard' and 'densityquantile'

Usage

```
gompertzHaz()
```

Details

The `distinfo` function is used to provide basic distribution specific information to other `spatsurv` functions. The user is required to provide the following information in the returned list: `npars`, the number of parameters in this distribution; `parnames`, the names of the parameters; `trans`, the transformation scale on which the priors will be provided; `itrans`, the inverse transformation function that will be applied to the parameters before the hazard, and other functions are evaluated; `jacobian`, the derivative of the inverse transformation function with respect to each of the parameters; and `hessian`, the second derivatives of the inverse transformation function with respect to each of the parameters – note that currently the package `spatsurv` only allows the use of functions where the parameters are transformed independently.

The `basehazard` function is used to evaluate the baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradbasehazard` function is used to evaluate the gradient of the baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hessbasehazard` function is used to evaluate the Hessian of the baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `cumbasehazard` function is used to evaluate the cumulative baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradcumbasehazard` function is used to evaluate the gradient of the cumulative baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hesscumbasehazard` function is used to evaluate the Hessian of the cumulative baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `densityquantile` function is used to return quantiles of the density function. This is NOT REQUIRED for running the MCMC, merely for us in post-processing with the `predict` function where `type` is `'densityquantile'`. In the case of the Weibull model for the baseline hazard, it can be shown that the `q`-th quantile is:

Value

an object inheriting class `'basehazardspec'`

See Also

[tpowHaz](#), [exponentialHaz](#), [makehamHaz](#), [weibullHaz](#)

gradbasehazard *gradbasehazard function*

Description

Generic function for computing the gradient of the baseline hazard

Usage

```
gradbasehazard(obj, ...)
```

Arguments

obj an object
... additional arguments – currently there are none, but this is for extensibility

Value

method gradbasehazard

See Also

[gradbasehazard.basehazardspec](#), [exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpow-Haz](#)

gradbasehazard.basehazardspec
gradbasehazard.basehazardspec function

Description

A function to retrieve the gradient of the baseline hazard function

Usage

```
## S3 method for class 'basehazardspec'  
gradbasehazard(obj, ...)
```

Arguments

obj an object of class basehazardspec
... additional arguments – currently there are none, but this is for extensibility

Value

a function returning the gradient of the baseline hazard

See Also

[exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

gradcumbasehazard *gradcumbasehazard function*

Description

Generic function for computing the gradient of the cumulative baseline hazard

Usage

```
gradcumbasehazard(obj, ...)
```

Arguments

obj an object
 ... additional arguments – currently there are none, but this is for extensibility

Value

method gradcumbasehazard

See Also

[gradcumbasehazard.basehazardspec](#), [exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

gradcumbasehazard.basehazardspec
 gradcumbasehazard.basehazardspec function

Description

A function to retrieve the gradient of the cumulative baseline hazard function

Usage

```
## S3 method for class 'basehazardspec'  
gradcumbasehazard(obj, ...)
```

Arguments

obj an object of class basehazardspec
 ... additional arguments – currently there are none, but this is for extensibility

Value

a function returning the gradient of the cumulative baseline hazard

See Also

[exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

grid2spdf *grid2spdf function*

Description

A function to convert a regular (x,y) grid of centroids into a SpatialPoints object

Usage

```
grid2spdf(xgrid, ygrid, proj4string = CRS(as.character(NA)))
```

Arguments

xgrid vector of x centroids (equally spaced)
ygrid vector of x centroids (equally spaced)
proj4string an optional proj4string, projection string for the grid, set using the function CRS

Value

a SpatialPolygonsDataFrame

grid2spix *grid2spix function*

Description

A function to convert a regular (x,y) grid of centroids into a SpatialPixels object

Usage

```
grid2spix(xgrid, ygrid, proj4string = CRS(as.character(NA)))
```

Arguments

xgrid vector of x centroids (equally spaced)
ygrid vector of x centroids (equally spaced)
proj4string an optional proj4string, projection string for the grid, set using the function CRS

Value

a SpatialPixels object

grid2spts	<i>grid2spts function</i>
-----------	---------------------------

Description

A function to convert a regular (x,y) grid of centroids into a SpatialPoints object

Usage

```
grid2spts(xgrid, ygrid, proj4string = CRS(as.character(NA)))
```

Arguments

xgrid	vector of x centroids (equally spaced)
ygrid	vector of x centroids (equally spaced)
proj4string	an optional proj4string, projection string for the grid, set using the function CRS

Value

a SpatialPoints object

gridY	<i>gridY function</i>
-------	-----------------------

Description

A function to put estimated individual Y's onto a grid

Usage

```
gridY(Y, control)
```

Arguments

Y	estimate of Y
control	control parameters

Value

...

gridY_polygonal	<i>gridY_polygonal function</i>
-----------------	---------------------------------

Description

A function to put estimated individual Y's onto a grid

Usage

```
gridY_polygonal(Y, control)
```

Arguments

Y	estimate of Y
control	control parameters

Value

...

guess_t	<i>guess_t function</i>
---------	-------------------------

Description

A function to get an initial guess of the failure time t, to be used in calibrating the MCMC. Not for general use

Usage

```
guess_t(surv)
```

Arguments

surv	an object of class Surv
------	-------------------------

Value

a guess at the failure times

hasNext	<i>generic hasNext method</i>
---------	-------------------------------

Description

test if an iterator has any more values to go

Usage

```
hasNext(obj)
```

Arguments

obj	an iterator
-----	-------------

hasNext.iter	<i>hasNext.iter function</i>
--------------	------------------------------

Description

method for iter objects test if an iterator has any more values to go

Usage

```
## S3 method for class 'iter'
hasNext(obj)
```

Arguments

obj	an iterator
-----	-------------

hazardexceedance	<i>hazardexceedance function</i>
------------------	----------------------------------

Description

A function to compute exceedance probabilities for the spatially correlated frailties.

Usage

```
hazardexceedance(threshold, direction = "upper")
```

Arguments

threshold	vector of thresholds
direction	default is "upper" which will calculate $P(Y > \text{threshold})$, alternative is "lower", which will calculate $P(Y < \text{threshold})$

Value

a function that can be passed to the function MCE in order to compute the exceedance probabilities

See Also

[print.mcmcpatsurv](#), [quantile.mcmcpatsurv](#), [summary.mcmcpatsurv](#), [vcov.mcmcpatsurv](#), [frailty-lag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcpatsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#),

hazardpars

hazardpars function

Description

A function to return the mcmc chains for the hazard function parameters

Usage

```
hazardpars(x)
```

Arguments

x an object of class mcmcpatsurv

Value

the omega mcmc chains

See Also

[print.mcmcpatsurv](#), [quantile.mcmcpatsurv](#), [summary.mcmcpatsurv](#), [vcov.mcmcpatsurv](#), [frailty-lag1](#), [spatialpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcpatsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

hazard_PP	<i>hazard_PP function</i>
-----------	---------------------------

Description

A function to compute an individual's hazard function.

Usage

```
hazard_PP(inputs)
```

Arguments

inputs	inputs for the function including the model matrix, frailties, fixed effects and the parameters of the baseline hazard derived from this model
--------	--

Value

the hazard function for the individual

hessbasehazard	<i>hessbasehazard function</i>
----------------	--------------------------------

Description

Generic function for computing the hessian of the baseline hazard

Usage

```
hessbasehazard(obj, ...)
```

Arguments

obj	an object
...	additional arguments – currently there are none, but this is for extensibility

Value

method hessbasehazard

See Also

[hessbasehazard.basehazardspec](#), [exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpow-Haz](#)

hessbasehazard.basehazardspec
hessbasehazard.basehazardspec function

Description

A function to retrieve the Hessian of the baseline hazard function

Usage

```
## S3 method for class 'basehazardspec'  
hessbasehazard(obj, ...)
```

Arguments

obj an object of class basehazardspec
... additional arguments – currently there are none, but this is for extensibility

Value

a function returning the Hessian of the baseline hazard

See Also

[exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

hesscumbasehazard *hesscumbasehazard function*

Description

Generic function for computing the Hessian of the cumulative baseline hazard

Usage

```
hesscumbasehazard(obj, ...)
```

Arguments

obj an object
... additional arguments – currently there are none, but this is for extensibility

Value

method hesscumbasehazard

See Also

[hesscumbasehazard.basehazardspec](#), [exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

hesscumbasehazard.basehazardspec
hesscumbasehazard.basehazardspec function

Description

A function to retrieve the hessian of the cumulative baseline hazard function

Usage

```
## S3 method for class 'basehazardspec'
hesscumbasehazard(obj, ...)
```

Arguments

obj an object of class basehazardspec
 ... additional arguments – currently there are none, but this is for extensibility

Value

a function returning the hessian of the cumulative baseline hazard

See Also

[exponentialHaz](#), [weibullHaz](#), [gompertzHaz](#), [makehamHaz](#), [tpowHaz](#)

imputationModel *imputationModel function*

Description

A function to

Usage

```
imputationModel(formula, offset, covariateData, priors)
```

Arguments

formula	X
offset	X
covariateData	X
priors	X

Value

...

Independent	<i>Independent function</i>
-------------	-----------------------------

Description

A function to declare and also evaluate an exponential covariance function.

Usage

Independent()

Value

the exponential covariance function

See Also

[SpikedExponentialCovFct](#), [covmodel](#), [CovarianceFct](#)

indepGaussianprior	<i>indepGaussianprior function</i>
--------------------	------------------------------------

Description

A function for evaluating the log of an independent Gaussian prior for a given set of parameter values.

Usage

indepGaussianprior(beta = NULL, omega = NULL, eta = NULL, priors)

Arguments

beta	parameter beta at which prior is to be evaluated
omega	parameter omega at which prior is to be evaluated
eta	parameter eta at which prior is to be evaluated
priors	an object of class mcmcPriors, see ?mcmcPriors

Value

the log of the prior evaluated at the given parameter values

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

`indepGaussianpriorST` *indepGaussianpriorST function*

Description

A function to

Usage

```
indepGaussianpriorST(beta = NULL, omega = NULL, eta = NULL, priors)
```

Arguments

beta	X
omega	X
eta	X
priors	X

Value

...

inference.control *inference.control function*

Description

A function to control inferential settings. This function is used to set parameters for more advanced use of spatSurv.

Usage

```
inference.control(
  gridded = FALSE,
  cellwidth = NULL,
  ext = 2,
  imputation = NULL,
  optimcontrol = NULL,
  hessian = FALSE,
  plotcal = FALSE,
  timeonlyMCMC = FALSE,
  nugget = FALSE,
  savenugget = FALSE,
  split = 0.5,
  logUsigma_priorsmean = 0,
  logUsigma_priorssd = 0.5,
  nis = NULL,
  olinfo = NULL
)
```

Arguments

gridded	logical. Whether to perform computation on a grid. Default is FALSE.
cellwidth	the width of computational cells to use
ext	integer the number of times to extend the computational grid by in order to perform computation. The default is 2.
imputation	for polygonal data, an optional model for inference at the sub-polygonal level, see function <code>imputationModel</code>
optimcontrol	a list of optional arguments to be passed to <code>optim</code> for non-spatial models
hessian	whether to return a numerical hessian. Set this to TRUE for non-spatial models. equal to the number of parameters of the baseline hazard
plotcal	logical, whether to produce plots of the MCMC calibration process, this is a technical option and should only be set to TRUE if poor mixing is evident (the printed <code>h</code> is low), then it is also useful to use a graphics device with multiple plotting windows.
timeonlyMCMC	logical, whether to only time the MCMC part of the algorithm, or whether to include in the reported running time the time taken to calibrate the method (default)

nugget	whether to include a nugget effect in the estimation. Note that only the mean and variance of the nugget effect is returned.
savenugget	whether to save the MCMC chain for the nugget effect
split	how to split the spatial and nugget proposal variance as a the proportion of variance assigned to the spatial effect apriori. Default is 0.5
logUsigma_priormean	prior mean for log standard deviation of nugget effect
logUsigma_priorsd	prior sd for log standard deviation of nugget effect
nis	list of cell counts, each element being a matrix, with attributes "x" and "y" giving grid centroids in x and y directions. Used to impute locations of aggregated data.
olinfo	to be supplied with nis, if continuous inference from aggregated data is required

Value

returns parameters to be used in the function `survspat`

See Also

[survspat](#)

insert

insert function

Description

A function to

Usage

```
insert(pars, idx, val)
```

Arguments

pars	X
idx	X
val	X

Value

...

invtransformweibull *invtransformweibull function*

Description

A function to transform estimates of the (alpha, lambda) parameters of the weibull baseline hazard function, so they are commensurate with R's inbuilt density functions, (shape, scale).

Usage

```
invtransformweibull(x)
```

Arguments

x a vector of paramters

Value

the transformed parameters. For the weibull model, this transforms 'shape' 'scale' (see ?dweibull) to 'alpha' and 'lambda' for the MCMC

is.burnin *is this a burn-in iteration?*

Description

if this mcmc iteration is in the burn-in period, return TRUE

Usage

```
is.burnin(obj)
```

Arguments

obj an mcmc iterator

Value

TRUE or FALSE

is.retain	<i>do we retain this iteration?</i>
-----------	-------------------------------------

Description

if this mcmc iteration is one not thinned out, this is true

Usage

```
is.retain(obj)
```

Arguments

obj an mcmc iterator

Value

TRUE or FALSE

iteration	<i>iteration number</i>
-----------	-------------------------

Description

within a loop, this is the iteration number we are currently doing.

Usage

```
iteration(obj)
```

Arguments

obj an mcmc iterator

Details

get the iteration number

Value

integer iteration number, starting from 1.

logPosterior	<i>logPosterior function</i>
--------------	------------------------------

Description

A function to evaluate the log-posterior of a spatial parametric proportional hazards model. Not intended for general use.

Usage

```
logPosterior(
  surv,
  X,
  beta,
  omega,
  eta,
  gamma,
  priors,
  cov.model,
  u,
  control,
  gradient = FALSE,
  hessian = FALSE
)
```

Arguments

surv	an object of class Surv
X	the design matrix, containing covariate information
beta	parameter beta
omega	parameter omega
eta	parameter eta
gamma	parameter gamma
priors	the priors, an object of class 'mcmcPriors'
cov.model	the spatial covariance model
u	vector of interpoint distances
control	a list containing various control parameters for the MCMC and post-processing routines
gradient	logical whether to evaluate the gradient
hessian	logical whether to evaluate the Hessian

Value

evaluates the log-posterior and the gradient and hessian, if required.

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.

logPosterior_gridded *logPosterior_gridded function*

Description

A function to evaluate the log-posterior of a spatial parametric proportional hazards model using gridded Y. Not intended for general use.

Usage

```
logPosterior_gridded(
  surv,
  X,
  beta,
  omega,
  eta,
  gamma,
  priors,
  cov.model,
  u,
  control,
  gradient = FALSE,
  hessian = FALSE
)
```

Arguments

surv	an object of class <code>Surv</code>
X	the design matrix, containing covariate information
beta	parameter beta
omega	parameter omega
eta	parameter eta
gamma	parameter gamma
priors	the priors, an object of class <code>'mcmcPriors'</code>
cov.model	the spatial covariance model
u	vector of interpoint distances
control	a list containing various control parameters for the MCMC and post-processing routines
gradient	logical whether to evaluate the gradient
hessian	logical whether to evaluate the Hessian

Value

evaluates the log-posterior and the gradient and hessian, if required.

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.

logPosterior_polygonal

logPosterior_polygonal function

Description

A function to evaluate the log-posterior of a spatial parametric proportional hazards model. Not intended for general use.

Usage

```
logPosterior_polygonal(
  surv,
  X,
  beta,
  omega,
  eta,
  gamma,
  priors,
  cov.model,
  u,
  control,
  gradient = FALSE,
  hessian = FALSE
)
```

Arguments

surv	an object of class Surv
X	the design matrix, containing covariate information
beta	parameter beta
omega	parameter omega
eta	parameter eta
gamma	parameter gamma
priors	the priors, an object of class 'mcmcPriors'
cov.model	the spatial covariance model

u	vector of interpoint distances
control	a list containing various control parameters for the MCMC and post-processing routines
gradient	logical whether to evaluate the gradient
hessian	logical whether to evaluate the Hessian

Value

evaluates the log-posterior and the gradient and hessian, if required.

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.

logPosterior_SPDE *logPosterior_SPDE function*

Description

A function to evaluate the log-posterior of a spatial parametric proportional hazards model. Not intended for general use.

Usage

```
logPosterior_SPDE(
  surv,
  X,
  beta,
  omega,
  eta,
  gamma,
  priors,
  cov.model,
  u,
  control,
  gradient = FALSE,
  hessian = FALSE
)
```

Arguments

surv	an object of class Surv
X	the design matrix, containing covariate information
beta	parameter beta
omega	parameter omega

eta	parameter eta
gamma	parameter gamma
priors	the priors, an object of class 'mcmcPriors'
cov.model	the spatial covariance model
u	vector of interpoint distances
control	a list containing various control parameters for the MCMC and post-processing routines
gradient	logical whether to evaluate the gradient
hessian	logical whether to evaluate the Hessian

Value

evaluates the log-posterior and the gradient and hessian, if required.

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.
2. Finn Lindgren, Havard Rue, Johan Lindstrom. An explicit link between Gaussian fields and Gaussian Markov random fields: the stochastic partial differential equation approach. *Journal of the Royal Statistical Society: Series B* 73(4)

loop.mcmc

loop over an iterator

Description

useful for testing progress bars

Usage

```
loop.mcmc(object, sleep = 1)
```

Arguments

object	an mcmc iterator
sleep	pause between iterations in seconds

makehamHaz	<i>makehamHaz function</i>
------------	----------------------------

Description

A function to define a parametric proportional hazards model where the baseline hazard is taken from the Gompertz-Makeham model. This function returns an object inheriting class 'basehazard-spec', list of functions 'distinfo', 'basehazard', 'gradbasehazard', 'hessbasehazard', 'cumbasehazard', 'gradcumbasehazard', 'hesscumbasehazard' and 'densityquantile'

Usage

```
makehamHaz()
```

Details

The `distinfo` function is used to provide basic distribution specific information to other `spatsurv` functions. The user is required to provide the following information in the returned list: `npars`, the number of parameters in this distribution; `parnames`, the names of the parameters; `trans`, the transformation scale on which the priors will be provided; `itrans`, the inverse transformation function that will be applied to the parameters before the hazard, and other functions are evaluated; `jacobian`, the derivative of the inverse transformation function with respect to each of the parameters; and `hessian`, the second derivatives of the inverse transformation function with respect to each of the parameters – note that currently the package `spatsurv` only allows the use of functions where the parameters are transformed independently.

The `basehazard` function is used to evaluate the baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradbasehazard` function is used to evaluate the gradient of the baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hessbasehazard` function is used to evaluate the Hessian of the baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `cumbasehazard` function is used to evaluate the cumulative baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradcumbasehazard` function is used to evaluate the gradient of the cumulative baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hesscumbasehazard` function is used to evaluate the Hessian of the cumulative baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `densityquantile` function is used to return quantiles of the density function. This is NOT REQUIRED for running the MCMC, merely for us in post-processing with the `predict` function where `type` is 'densityquantile'. In the case of the Weibull model for the baseline hazard, it can be shown that the `q`-th quantile is:

Value

an object inheriting class 'basehazardspec'

See Also

[tpowHaz](#), [exponentialHaz](#), [gompertzHaz](#), [weibullHaz](#)

maxlikparamPHsurv *maxlikparamPHsurv function*

Description

A function to get initial estimates of model parameters using maximum likelihood. Not intended for general purpose use.

Usage

```
maxlikparamPHsurv(surv, X, control)
```

Arguments

surv	an object of class Surv
X	the design matrix, containing covariate information
control	a list containing various control parameters for the MCMC and post-processing routines

Value

initial estimates of the parameters

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.

MCE	<i>MCE function</i>
-----	---------------------

Description

A function to compute Monte Carlo expectations from an object inheriting class `mcmcspsurv`

Usage

```
MCE(object, fun)
```

Arguments

<code>object</code>	an object inheriting class <code>mcmcspsurv</code>
<code>fun</code>	a function with arguments <code>beta</code> , <code>omega</code> , <code>eta</code> and <code>Y</code>

Value

the Monte Carlo mean of the function over the posterior.

See Also

[print.mcmcspsurv](#), [quantile.mcmcspsurv](#), [summary.mcmcspsurv](#), [vcov.mcmcspsurv](#), [frailty-lag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspsurv](#), [prior-posterior](#), [posteriorcov](#), [hazardexceedance](#)

<code>mcmcLoop</code>	<i>iterator for MCMC loops</i>
-----------------------	--------------------------------

Description

control an MCMC loop with this iterator

Usage

```
mcmcLoop(N, burnin, thin, trim = TRUE, progressor = mcmcProgressPrint)
```

Arguments

<code>N</code>	number of iterations
<code>burnin</code>	length of burn-in
<code>thin</code>	frequency of thinning
<code>trim</code>	whether to cut off iterations after the last retained iteration
<code>progressor</code>	a function that returns a progress object

mcmcpars	<i>mcmcpars function</i>
----------	--------------------------

Description

A function for setting MCMC options.

Usage

```
mcmcpars(nits, burn, thin, inits = NULL, adaptivescheme = NULL)
```

Arguments

nits	numer of iterations,
burn	length of burnin
thin	thinning parameter eg operated on chain every 'thin' iteration (eg store output or compute some posterior functional)
inits	NOT CURRENTLY IN USE
adaptivescheme	NOT CURRENTLY IN USE

Value

mcmc parameters

mcmcPriors	<i>mcmcPriors function</i>
------------	----------------------------

Description

A function to define priors for the MCMC.

Usage

```
mcmcPriors(
  betaprior = NULL,
  omegaprior = NULL,
  etaprior = NULL,
  call = NULL,
  derivative = NULL
)
```


Arguments

betaprior	prior for beta, the covariate effects
omegaprior	prior for omega, the parameters of the baseline hazard
etaprior	prior for eta, the parameters of the latent field
call	function to evaluate the log-prior e.g. <code>logindepGaussianprior</code>
derivative	function to evaluate the first and second derivatives of the prior

Details

The package `spatsurv` only provides functionality for the built-in Gaussian priors. However, the choice of prior is extensible by the user by creating functions similar to the functions `betapriorGauss`, `omegapriorGauss`, `etapriorGauss`, `indepGaussianprior` and `derivindepGaussianprior`: the first three of which provide a mechanism for storing and retrieving the parameters of the priors; the fourth, a function for evaluating the log of the prior for a given set of parameter values; and the fifth, a function for evaluating the first and second derivatives of the log of the prior. It is assumed that parameters are a priori independent. The user interested in using other priors is encouraged to look at the structure of the five functions mentioned above.

Value

an object of class `mcmcPriors`

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

<code>mcmcProgressNone</code>	<i>null progress monitor</i>
-------------------------------	------------------------------

Description

a progress monitor that does nothing

Usage

```
mcmcProgressNone(mcmcloop)
```

Arguments

<code>mcmcloop</code>	an mcmc loop iterator
-----------------------	-----------------------

Value

a progress monitor

mcmcProgressPrint *printing progress monitor*

Description

a progress monitor that prints each iteration

Usage

```
mcmcProgressPrint(mcmcloop)
```

Arguments

mcmcloop an mcmc loop iterator

Value

a progress monitor

mcmcProgressTextBar *text bar progress monitor*

Description

a progress monitor that uses a text progress bar

Usage

```
mcmcProgressTextBar(mcmcloop)
```

Arguments

mcmcloop an mcmc loop iterator

Value

a progress monitor

midpts	<i>midpts function</i>
--------	------------------------

Description

A function to compute the midpoints of a vector

Usage

```
midpts(x)
```

Arguments

x	a vector
---	----------

Value

the midpoints, a vector of length $\text{length}(x)-1$

multiWayHaz	<i>multiWayHaz function</i>
-------------	-----------------------------

Description

A function to

Usage

```
multiWayHaz(bhlist, bhtime, bhfix, MLimits = NULL)
```

Arguments

bhlist	X
bhtime	X
bhfix	X
MLimits	X

Value

...

neighLocs	<i>neighLocs function</i>
-----------	---------------------------

Description

A function used in the computation of neighbours on non-rectangular grids. Not intended for general use.

Usage

```
neighLocs(coord, cellwidth, order)
```

Arguments

coord	coordinate of interest
cellwidth	a scalar, the width of the grid cells
order	the order of the SPDE approximation: see Lindgren et al 2011 for details

Value

coordinates of centroids of neighbours

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.
2. Finn Lindgren, Havard Rue, Johan Lindstrom. An explicit link between Gaussian fields and Gaussian Markov random fields: the stochastic partial differential equation approach. *Journal of the Royal Statistical Society: Series B* 73(4)

neighOrder	<i>neighOrder function</i>
------------	----------------------------

Description

A function to compute the order of a set of neighbours. Not intended for general use.

Usage

```
neighOrder(neighlocs)
```

Arguments

neighlocs	an object created by the function neighLocs
-----------	---

Value

the neighbour orders

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.
2. Finn Lindgren, Havard Rue, Johan Lindstrom. An explicit link between Gaussian fields and Gaussian Markov random fields: the stochastic partial differential equation approach. *Journal of the Royal Statistical Society: Series B* 73(4)

nextStep	<i>next step of an MCMC chain</i>
----------	-----------------------------------

Description

just a wrapper for nextElem really.

Usage

```
nextStep(object)
```

Arguments

object an mcmc loop object

NonSpatialLogLikelihood_or_gradient	<i>NonSpatialLogLikelihood_or_gradient function</i>
-------------------------------------	---

Description

A function to evaluate the log-likelihood of a non-spatial parametric proportional hazards model. Not intended for general use.

Usage

```
NonSpatialLogLikelihood_or_gradient(
  surv,
  X,
  beta,
  omega,
  control,
  loglikelihood,
  gradient
)
```

Arguments

surv	an object of class Surv
X	the design matrix, containing covariate information
beta	parameter beta
omega	parameter omega
control	a list containing various control parameters for the MCMC and post-processing routines
loglikelihood	logical whether to evaluate the log-likelihood
gradient	logical whether to evaluate the gradient

Value

...

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.

omegapriorGauss	<i>omegapriorGauss function</i>
-----------------	---------------------------------

Description

A function to define Gaussian priors for omega. This function simply stores a vector of means and standard deviations to be passed to the main MCMC function, survspat.

Usage

```
omegapriorGauss(mean, sd)
```

Arguments

mean	the prior mean, a vector of length 1 or more. 1 implies a common mean.
sd	the prior standard deviation, a vector of length 1 or more. 1 implies a common standard deviation.

Value

an object of class "omegapriorGauss"

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

omegapriorGaussST	<i>omegapriorGaussST function</i>
-------------------	-----------------------------------

Description

A function to

Usage

```
omegapriorGaussST(basehaz, fmean, fsd, taumean, tausd, thetamean, thetasd)
```

Arguments

basehaz	X
fmean	X
fsd	X
taumean	X
tausd	X
thetamean	X
thetasd	X

Value

...

optifix	<i>optifix function</i>
---------	-------------------------

Description

optifix. Optimise with fixed parameters

Usage

```
optifix(
  par,
  fixed,
  fn,
  gr = NULL,
  ...,
  method = c("Nelder-Mead", "BFGS", "CG", "L-BFGS-B", "SANN"),
  lower = -Inf,
  upper = Inf,
  control = list(),
  hessian = FALSE
)
```

Arguments

par	X
fixed	X
fn	X
gr	X
...	X
method	X
lower	X
upper	X
control	X
hessian	X

Details

its like optim, but with fixed parameters.

specify a second argument 'fixed', a vector of TRUE/FALSE values. If TRUE, the corresponding parameter in fn() is fixed. Otherwise its variable and optimised over.

The return thing is the return thing from optim() but with a couple of extra bits - a vector of all the parameters and a vector copy of the 'fixed' argument.

Written by Barry Rowlingson <b.rowlingson@lancaster.ac.uk> October 2011

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and must retain the text: "Originally written by Barry Rowlingson" in comments.

Value

...

plot.FFTgrid	<i>plot.FFTgrid function</i>
--------------	------------------------------

Description

A function to

Usage

```
## S3 method for class 'FFTgrid'
plot(x, y = NULL, ...)
```

Arguments

x	X
y	X
...	X

Value

...

`plotsurv`*plotsurv function*

Description

A function to produce a 2-D plot of right censored spatial survival data.

Usage

```
plotsurv(
  spp,
  ss,
  maxcex = 1,
  transform = identity,
  background = NULL,
  eventpt = 19,
  eventcol = "red",
  censpt = "+",
  censcol = "black",
  xlim = NULL,
  ylim = NULL,
  xlab = NULL,
  ylab = NULL,
  add = FALSE,
  ...
)
```

Arguments

<code>spp</code>	A spatial points data frame
<code>ss</code>	A Surv object (with right-censoring)
<code>maxcex</code>	maximum size of dots default is equivalent to setting <code>cex</code> equal to 1
<code>transform</code>	optional transformation to apply to the data, a function, for example <code>'sqrt'</code>
<code>background</code>	a background object to plot default is null, which gives a blank background note that if non-null, the parameters <code>xlim</code> and <code>ylim</code> will be derived from this object.
<code>eventpt</code>	The type of point to illustrate events, default is 19 (see <code>?pch</code>)
<code>eventcol</code>	the colour of events, default is black
<code>censpt</code>	The type of point to illustrate events, default is "+" (see <code>?pch</code>)
<code>censcol</code>	the colour of censored observations, default is red
<code>xlim</code>	optional x-limits of plot, default is to choose this automatically

<code>ylim</code>	optional y-limits of plot, default is to choose this automatically
<code>xlab</code>	label for x-axis
<code>ylab</code>	label for y-axis
<code>add</code>	logical, whether to add the survival plot on top of an existing plot, default is FALSE, which produces a plot in a new device
<code>...</code>	other arguments to pass to plot

Value

Plots the survival data non-censored observations appear as dots and censored observations as crosses. The size of the dot is proportional to the observed time.

<code>polyadd</code>	<i>polyadd function</i>
----------------------	-------------------------

Description

A function to add two polynomials in the form of vectors of coefficients. The first element of the vector being the constant (order 0) term

Usage

```
polyadd(poly1, poly2)
```

Arguments

<code>poly1</code>	a vector of coefficients for the first polynomial of length degree plus 1
<code>poly2</code>	a vector of coefficients for the second polynomial of length degree plus 1

Value

the coefficients of the sum of `poly1` and `poly2`

polymult	<i>polymult function</i>
----------	--------------------------

Description

A function to multiply two polynomials in the form of vectors of coefficients. The first element of the vector being the constant (order 0) term

Usage

```
polymult(poly1, poly2)
```

Arguments

poly1	a vector of coefficients for the first polynomial of length degree plus 1
poly2	a vector of coefficients for the second polynomial of length degree plus 1

Value

the coefficients of the product of poly1 and poly2

posteriorcov	<i>posteriorcov function</i>
--------------	------------------------------

Description

A function to produce a plot of the posterior covariance function with upper and lower quantiles.

Usage

```
posteriorcov(  
  x,  
  probs = c(0.025, 0.5, 0.975),  
  rmax = NULL,  
  n = 100,  
  plot = TRUE,  
  bw = FALSE,  
  corr = FALSE,  
  ...  
)
```

Arguments

<code>x</code>	an object of class <code>mcmcspatsurv</code>
<code>probs</code>	vector of probabilities to be fed to quantile function
<code>rmax</code>	maximum distance in space to compute this distance up to
<code>n</code>	the number of points at which to evaluate the posterior covariance.
<code>plot</code>	whether to plot the result
<code>bw</code>	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
<code>corr</code>	logical whether to return the correlation function, default is FALSE i.e. returns the covariance function
<code>...</code>	other arguments to be passed to <code>matplot</code> function

Value

produces a plot of the posterior spatial covariance function.

See Also

[print.mcmcspatsurv](#), [quantile.mcmcspatsurv](#), [summary.mcmcspatsurv](#), [vcov.mcmcspatsurv](#), [frailty-lag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspatsurv](#), [prior-posterior](#), [MCE](#), [hazardexceedance](#)

`predict.mcmcspatsurv` *predict.mcmcspatsurv* function

Description

A function to produce predictions from MCMC output. These could include quantiles of the individual density, survival or hazard functions or quantiles of the density function (if available analytically).

Usage

```
## S3 method for class 'mcmcspatsurv'
predict(
  object,
  type = "density",
  t = NULL,
  n = 110,
  indx = NULL,
  probs = c(0.025, 0.5, 0.975),
  plot = TRUE,
  pause = TRUE,
  bw = FALSE,
  ...
)
```

Arguments

object	an object of class mcmcspsurv
type	can be "density", "hazard", "survival" or "densityquantile". Default is "density". Note that "densityquantile" is not always analytically tractable for some choices of baseline hazard function.
t	optional vector of times at which to compute the quantiles, Default is NULL, in which case a uniformly spaced vector of length n from 0 to the maximum time is used
n	the number of points at which to compute the quantiles if t is NULL
indx	the index number of a particular individual or vector of indices of individuals for which the quantiles should be produced
probs	vector of probabilities
plot	whether to plot the result
pause	logical whether to pause between plots, the default is TRUE
bw	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
...	other arguments, not used here

Value

the required predictions

See Also

[print.mcmcspsurv](#), [quantile.mcmcspsurv](#), [summary.mcmcspsurv](#), [vcov.mcmcspsurv](#), [frailty-lag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

print.mcmc

print.mcmc function

Description

print method print an mcmc iterator's details

Usage

```
## S3 method for class 'mcmc'
print(x, ...)
```

Arguments

x	a mcmc iterator
...	other args

```
print.mcmcsratsurv    print.mcmcsratsurv function
```

Description

A function to print summary tables from an MCMC run

Usage

```
## S3 method for class 'mcmcsratsurv'
print(x, probs = c(0.5, 0.025, 0.975), digits = 3, scientific = -3, ...)
```

Arguments

<code>x</code>	an object inheriting class <code>mcmcsratsurv</code>
<code>probs</code>	vector of quantiles to return
<code>digits</code>	see help file <code>?format</code>
<code>scientific</code>	see help file <code>?format</code>
<code>...</code>	additional arguments, not used here

Value

prints summary tables to the console

See Also

[quantile.mcmcsratsurv](#), [summary.mcmcsratsurv](#), [vcov.mcmcsratsurv](#), [frailtylag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcsratsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

```
print.mlspatsurv    print.mlspatsurv function
```

Description

A function to print summary tables from an MCMC run

Usage

```
## S3 method for class 'mlspatsurv'
print(x, probs = c(0.5, 0.025, 0.975), digits = 3, scientific = -3, ...)
```

Arguments

x an object inheriting class `mcmcspatsurv`
probs vector of quantiles to return
digits see help file `?format`
scientific see help file `?format`
... additional arguments, not used here

Value

prints summary tables to the console

See Also

[quantile.mcmcspatsurv](#), [summary.mcmcspatsurv](#), [vcov.mcmcspatsurv](#), [frailtylag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspatsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

`print.textSummary` *print.textSummary function*

Description

A function to print summary tables from an MCMC run

Usage

```
## S3 method for class 'textSummary'  
print(x, ...)
```

Arguments

x an object inheriting class `textSummary`
... additional arguments, not used here

Value

prints a text summary of 'x' to the console

priorposterior *priorposterior function*

Description

A function to produce plots of the prior (which shows as a red line) and posterior (showing as a histogram)

Usage

```
priorposterior(  
  x,  
  breaks = 30,  
  ylab = "Density",  
  main = "",  
  pause = TRUE,  
  bw = FALSE,  
  ...  
)
```

Arguments

x	an object inheriting class <code>mcmcspatsurv</code>
breaks	see <code>?hist</code>
ylab	optional y label
main	optional title
pause	logical whether to pause between plots, the default is <code>TRUE</code>
bw	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
...	other arguments passed to the <code>hist</code> function

Value

plots of the prior (red line) and posterior (histogram).

See Also

[print.mcmcspatsurv](#), [quantile.mcmcspatsurv](#), [summary.mcmcspatsurv](#), [vcov.mcmcspatsurv](#), [frailty-lag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspatsurv](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

proposalVariance *proposalVariance function*

Description

A function to compute an approximate scaling matrix for the MCMC algorithm. Not intended for general use.

Usage

```
proposalVariance(  
  X,  
  surv,  
  betahat,  
  omegahat,  
  Yhat,  
  priors,  
  cov.model,  
  u,  
  control  
)
```

Arguments

X	the design matrix, containing covariate information
surv	an object of class Surv
betahat	an estimate of beta
omegahat	an estimate of omega
Yhat	an estimate of Y
priors	the priors
cov.model	the spatial covariance model
u	a vector of pairwise distances
control	a list containing various control parameters for the MCMC and post-processing routines

Value

an estimate of eta and also an approximate scaling matrix for the MCMC

proposalVariance_gridded

proposalVariance_gridded function

Description

A function to compute an approximate scaling matrix for the MCMC algorithm. Not intended for general use.

Usage

```
proposalVariance_gridded(
  X,
  surv,
  betahat,
  omegahat,
  Yhat,
  priors,
  cov.model,
  u,
  control
)
```

Arguments

X	the design matrix, containing covariate information
surv	an object of class Surv
betahat	an estimate of beta
omegahat	an estimate of omega
Yhat	an estimate of Y
priors	the priors
cov.model	the spatial covariance model
u	a vector of pairwise distances
control	a list containing various control parameters for the MCMC and post-processing routines

Value

an estimate of eta and also an approximate scaling matrix for the MCMC

proposalVariance_polygonal
proposalVariance_polygonal function

Description

A function to compute an approximate scaling matrix for the MCMC algorithm. Not intended for general use.

Usage

```
proposalVariance_polygonal(  
  X,  
  surv,  
  betahat,  
  omegahat,  
  Yhat,  
  priors,  
  cov.model,  
  u,  
  control  
)
```

Arguments

X	the design matrix, containing covariate information
surv	an object of class Surv
betahat	an estimate of beta
omegahat	an estimate of omega
Yhat	an estimate of Y
priors	the priors
cov.model	the spatial covariance model
u	a vector of pairwise distances
control	a list containing various control parameters for the MCMC and post-processing routines

Value

an estimate of eta and also an approximate scaling matrix for the MCMC

`proposalVariance_SPDE` *proposalVariance_SPDE function*

Description

A function to compute an approximate scaling matrix for the MCMC algorithm. Not intended for general use.

Usage

```
proposalVariance_SPDE(
  X,
  surv,
  betahat,
  omegahat,
  Yhat,
  priors,
  cov.model,
  u,
  control
)
```

Arguments

<code>X</code>	the design matrix, containing covariate information
<code>surv</code>	an object of class <code>Surv</code>
<code>betahat</code>	an estimate of beta
<code>omegahat</code>	an estimate of omega
<code>Yhat</code>	an estimate of Y
<code>priors</code>	the priors
<code>cov.model</code>	the spatial covariance model
<code>u</code>	a vector of pairwise distances
<code>control</code>	a list containing various control parameters for the MCMC and post-processing routines

Value

an estimate of eta and also an approximate scaling matrix for the MCMC

PsplineHaz

PsplineHaz function**Description**

A function to define a parametric proportional hazards model where the baseline hazard is modelled by a basis spline and where the coefficients of the model follow a partially improper random walk prior. This function returns an object inheriting class 'basehazardspec', list of functions 'distinfo', 'basehazard', 'gradbasehazard', 'hessbasehazard', 'cumbasehazard', 'gradcumbasehazard', 'hesscumbasehazard' and 'densityquantile'

Usage

```
PsplineHaz(times, knots = quantile(times), degree = 3, MLinits = NULL)
```

Arguments

times	vector of survival times (both censored and uncensored)
knots	vector of knots in ascending order, must include minimum and maximum values of 'times'
degree	degree of the spline basis, default is 3
MLinits	optional starting values for the non-spatial maximisation routine using optim. Note that we are working with the log of the parameters. Default is -10 for each parameter.

Details

The `distinfo` function is used to provide basic distribution specific information to other `spatsurv` functions. The user is required to provide the following information in the returned list: `nparams`, the number of parameters in this distribution; `parnames`, the names of the parameters; `trans`, the transformation scale on which the priors will be provided; `itrans`, the inverse transformation function that will be applied to the parameters before the hazard, and other functions are evaluated; `jacobian`, the derivative of the inverse transformation function with respect to each of the parameters; and `hessian`, the second derivatives of the inverse transformation function with respect to each of the parameters – note that currently the package `spatsurv` only allows the use of functions where the parameters are transformed independently.

The `basehazard` function is used to evaluate the baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradbasehazard` function is used to evaluate the gradient of the baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hessbasehazard` function is used to evaluate the Hessian of the baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `cumbasehazard` function is used to evaluate the cumulative baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradcumbasehazard` function is used to evaluate the gradient of the cumulative baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hesscumbasehazard` function is used to evaluate the Hessian of the cumulative baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `densityquantile` function is used to return quantiles of the density function. This is NOT REQUIRED for running the MCMC, merely for us in post-processing with the `predict` function where `type` is `'densityquantile'`. In the case of the Weibull model for the baseline hazard, it can be shown that the `q`-th quantile is:

Value

an object inheriting class `'basehazardspec'`

See Also

[exponentialHaz](#), [gompertzHaz](#), [makehamHaz](#), [weibullHaz](#)

psplineprior

psplineprior function

Description

A function for evaluating the log of an independent Gaussian prior for a given set of parameter values.

Usage

```
psplineprior(beta = NULL, omega = NULL, eta = NULL, priors)
```

Arguments

<code>beta</code>	parameter <code>beta</code> at which prior is to be evaluated
<code>omega</code>	parameter <code>omega</code> at which prior is to be evaluated
<code>eta</code>	parameter <code>eta</code> at which prior is to be evaluated
<code>priors</code>	an object of class <code>mcmcPriors</code> , see <code>?mcmcPriors</code>

Value

the log of the prior evaluated at the given parameter values

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

psplineRWprior *psplineRWprior function*

Description

A function to define Gaussian priors for omega. This function simply stores a vector of means and standard deviations to be passed to the main MCMC function, `survspat`.

Usage

```
psplineRWprior(taumean, tausd, basehaz, order = 2)
```

Arguments

taumean	the prior mean, a vector of length 1 or more. 1 implies a common mean.
tausd	the prior standard deviation, a vector of length 1 or more. 1 implies a common standard deviation.
basehaz	an object inheriting class "basehazardspec", specifically, this function was used for such objects created by a call to the function <code>PsplineHaz</code>
order	the order of the random walk, default is 2

Value

an object of class "omegapriorGauss"

See Also

[survspat](#), [betapriorGauss](#), [omegapriorGauss](#), [etapriorGauss](#), [indepGaussianprior](#), [derivindepGaussianprior](#)

 QuadApprox

QuadApprox function

Description

A function to compute the second derivative of a function (of several real variables) using a quadratic approximation on a grid of points defined by the list `argRanges`. Also returns the local maximum.

Usage

```
QuadApprox(fun, npts, argRanges, plot = FALSE, ...)
```

Arguments

<code>fun</code>	a function
<code>npts</code>	integer number of points in each direction
<code>argRanges</code>	a list of ranges on which to construct the grid for each parameter
<code>plot</code>	whether to plot the quadratic approximation of the posterior (for two-dimensional parameters only)
<code>...</code>	other arguments to be passed to <code>fun</code>

Value

a 2 by 2 matrix containing the curvature at the maximum and the (x,y) value at which the maximum occurs

 quantile.mcmcspatsurv *quantile.mcmcspatsurv function*

Description

A function to extract quantiles of the parameters from an mcmc run

Usage

```
## S3 method for class 'mcmcspatsurv'
quantile(x, probs = c(0.025, 0.5, 0.975), ...)
```

Arguments

<code>x</code>	an object inheriting class <code>mcmcspatsurv</code>
<code>probs</code>	vector of probabilities
<code>...</code>	other arguments to be passed to the function, not used here

Value

quantiles of model parameters

See Also

[print.mcmcspsurv](#), [summary.mcmcspsurv](#), [vcov.mcmcspsurv](#), [frailtylag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

quantile.mlspatsurv *quantile.mlspatsurv function*

Description

A function to extract quantiles of the parameters from an mcmc run

Usage

```
## S3 method for class 'mlspatsurv'  
quantile(x, probs = c(0.025, 0.5, 0.975), ...)
```

Arguments

x	an object inheriting class mcmcspsurv
probs	vector of probabilities
...	other arguments to be passed to the function, not used here

Value

quantiles of model parameters

See Also

[print.mcmcspsurv](#), [summary.mcmcspsurv](#), [vcov.mcmcspsurv](#), [frailtylag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

randompars *randompars function*

Description

A function to return the mcmc chains for the spatially correlated frailties

Usage

```
randompars(x)
```

Arguments

x an object of class mcmcpatsurv

Value

the Y mcmc chains

See Also

[print.mcmcpatsurv](#), [quantile.mcmcpatsurv](#), [summary.mcmcpatsurv](#), [vcov.mcmcpatsurv](#), [frailty-lag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [baselinehazard](#), [predict.mcmcpatsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

reconstruct.bs *reconstruct.bs function*

Description

Generic function for reconstructing B-spline covariate effects. See `?reconstruct.bs.mcmcpatsurv` and `?reconstruct.bs.coxph`

Usage

```
reconstruct.bs(mod, ...)
```

Arguments

mod an object
... additional arguments

Value

method reconstruct.bs

reconstruct.bs.coxph *reconstruct.bs.coxph function*

Description

When `bs(varname)` has been used in the formula of a coxph model, this function can be used to reconstruct the predicted relative risk of that parameter over time.

Usage

```
## S3 method for class 'coxph'
reconstruct.bs(
  mod,
  varname,
  fun = NULL,
  probs = c(0.025, 0.975),
  bw = FALSE,
  xlab = NULL,
  ylab = NULL,
  plot = TRUE,
  ...
)
```

Arguments

<code>mod</code>	model output, created by function <code>survspat</code>
<code>varname</code>	name of the variable modelled by a B-spline
<code>fun</code>	optional function to feed in. Default is to plot relative risk against the covariate of interest. Useful choices include "identity" (but with no quotes), which plots the non-linear effect on the scale of the linear predictor.
<code>probs</code>	upper and lower quantiles for confidence regions to plot> The default is <code>c(0.025,0.975)</code> .
<code>bw</code>	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
<code>xlab</code>	label for x axis, there is a sensible default
<code>ylab</code>	label for y axis, there is a sensible default
<code>plot</code>	logical, whether to plot the effect of <code>varname</code> over time
<code>...</code>	other arguments to be passed to the plotting function.

Value

median, upper and lower confidence bands for the effect of `varname` over time; the function also produces a plot.

```
reconstruct.bs.mcmcspatsurv
      reconstruct.bs.mcmcspatsurv function
```

Description

When `bs(varname)` has been used in the formula of a model, this function can be used to reconstruct the posterior relative risk of that parameter over time.

Usage

```
## S3 method for class 'mcmcspatsurv'
reconstruct.bs(
  mod,
  varname,
  probs = c(0.025, 0.975),
  bw = FALSE,
  xlab = NULL,
  ylab = NULL,
  plot = TRUE,
  ...
)
```

Arguments

<code>mod</code>	model output, created by function <code>survspat</code>
<code>varname</code>	name of the variable modelled by a B-spline
<code>probs</code>	upper and lower quantiles for confidence regions to plot> The default is <code>c(0.025,0.975)</code> .
<code>bw</code>	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
<code>xlab</code>	label for x axis, there is a sensible default
<code>ylab</code>	label for y axis, there is a sensible default
<code>plot</code>	logical, whether to plot the effect of <code>varname</code> over time
<code>...</code>	other arguments to be passed to the plotting function.

Value

median, upper and lower confidence bands for the effect of `varname` over time; the function also produces a plot.

resetLoop	<i>reset iterator</i>
-----------	-----------------------

Description

call this to reset an iterator's state to the initial

Usage

```
resetLoop(obj)
```

Arguments

obj	an mcmc iterator
-----	------------------

residuals.mcmcspatsurv	<i>residuals.mcmcspatsurv function</i>
------------------------	--

Description

A function to compute Cox-Snell / modeified Cox-Snell / Martingale or Deviance residuals

Usage

```
## S3 method for class 'mcmcspatsurv'  
residuals(object, type = "Cox-Snell", ...)
```

Arguments

object	an object produced by the function survspat
type	type of residuals to return. Possible choices are 'Cox-Snell', 'modified-Cox-Snell', 'Martingale' or 'deviance'.
...	other arguments (not used here)

Value

the residuals

rootWeibullHaz	<i>rootWeibullHaz function</i>
----------------	--------------------------------

Description

A function to define a parametric proportional hazards model where the baseline hazard is taken from the Weibull model. This function returns an object inheriting class 'basehazardspec', list of functions 'distinfo', 'basehazard', 'gradbasehazard', 'hessbasehazard', 'cumbasehazard', 'gradcumbasehazard', 'hesscumbasehazard' and 'densityquantile'

Usage

```
rootWeibullHaz(MLinits = NULL)
```

Arguments

MLinits initial values for optim, default is NULL

Details

The `distinfo` function is used to provide basic distribution specific information to other `spatsurv` functions. The user is required to provide the following information in the returned list: `npars`, the number of parameters in this distribution; `parnames`, the names of the parameters; `trans`, the transformation scale on which the priors will be provided; `itrans`, the inverse transformation function that will be applied to the parameters before the hazard, and other functions are evaluated; `jacobian`, the derivative of the inverse transformation function with respect to each of the parameters; and `hessian`, the second derivatives of the inverse transformation function with respect to each of the parameters – note that currently the package `spatsurv` only allows the use of functions where the parameters are transformed independently.

The `basehazard` function is used to evaluate the baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradbasehazard` function is used to evaluate the gradient of the baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hessbasehazard` function is used to evaluate the Hessian of the baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `cumbasehazard` function is used to evaluate the cumulative baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradcumbasehazard` function is used to evaluate the gradient of the cumulative baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hesscumbasehazard` function is used to evaluate the Hessian of the cumulative baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `densityquantile` function is used to return quantiles of the density function. This is NOT REQUIRED for running the MCMC, merely for us in post-processing with the `predict` function where `type` is `'densityquantile'`. In the case of the Weibull model for the baseline hazard, it can be shown that the q -th quantile is:

Value

an object inheriting class `'basehazardspec'`

See Also

[tpwHaz](#), [exponentialHaz](#), [gompertzHaz](#), [makehamHaz](#)

`setTxtProgressBar2` *set the progress bar*

Description

update a text progress bar. See `help(txtProgressBar)` for more info.

Usage

```
setTxtProgressBar2(pb, value, title = NULL, label = NULL)
```

Arguments

<code>pb</code>	text progress bar object
<code>value</code>	new value
<code>title</code>	ignored
<code>label</code>	text for end of progress bar

`setupHazard` *setupHazard function*

Description

A function to set up the baseline hazard, cumulative hazard and derivative functions for use in evaluating the log posterior. This function is not intended for general use.

Usage

```
setupHazard(dist, pars, grad = FALSE, hess = FALSE)
```

Arguments

dist	an object of class 'basehazardspec'
pars	parameters with which to create the functions necessary to evaluate the log posterior
grad	logical, whether to create gradient functions for the baseline hazard and cumulative hazard
hess	logical, whether to create hessian functions for the baseline hazard and cumulative hazard

Value

a list of functions used in evaluating the log posterior

setupPrecMatStruct *setupPrecMatStruct function*

Description

A function to set up the computational grid and precision matrix structure for SPDE models.

Usage

```
setupPrecMatStruct(shape, cellwidth, no)
```

Arguments

shape	an object of class SpatialPolygons or SpatialPolygonsDataFrame
cellwidth	a scalar, the width of the grid cells
no	the order of the SPDE approximation: see Lindgren et al 2011 for details

Value

the computational grid and a function for constructing the precision matrix

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.
2. Finn Lindgren, Havard Rue, Johan Lindstrom. An explicit link between Gaussian fields and Gaussian Markov random fields: the stochastic partial differential equation approach. *Journal of the Royal Statistical Society: Series B* 73(4)

showGrid	<i>showGrid function</i>
----------	--------------------------

Description

A function to show the grid that will be used for a given cellwidth

Usage

```
showGrid(dat, cellwidth, ext = 2, boundingbox = NULL)
```

Arguments

dat	any spatial data object whose bounding box can be computed using the function <code>bbox</code> .
cellwidth	an initial suggested cellwidth
ext	the extension parameter for the FFT transform, set to 2 by default
boundingbox	optional bounding box over which to construct computational grid, supplied as an object on which the function <code>'bbox'</code> returns the bounding box

Value

a plot showing the grid and the data. Ideally the data should only just fit inside the grid.

simsurv	<i>simsurv function</i>
---------	-------------------------

Description

A function to simulate spatial parametric proportional hazards model. The function works by simulating candidate survival times using MCMC in parallel for each individual based on each individual's covariates and the common parameter effects, beta.

Usage

```
simsurv(
  X = cbind(age = runif(100, 5, 50), sex = rbinom(100, 1, 0.5), cancer = rbinom(100, 1,
    0.2)),
  beta = c(0.0296, 0.0261, 0.035),
  omega = 1,
  dist = exponentialHaz(),
  coords = matrix(runif(2 * nrow(X)), nrow(X), 2),
  cov.parameters = c(1, 0.1),
  cov.model = covmodel(model = "exponential", pars = NULL),
  mcmc.control = mcmcpars(nits = 1e+05, burn = 10000, thin = 90),
  savechains = TRUE
)
```

Arguments

X	a matrix of covariate information
beta	the parameter effects
omega	vector of parameters for the baseline hazard model
dist	the distribution choice: exp or weibull at present
coords	matrix with 2 columns giving the coordinates at which to simulate data
cov.parameters	a vector: the parameters for the covariance function
cov.model	an object of class covmodel, see ?covmodel
mcmc.control	mcmc control paramters, see ?mcmcpars
savechains	save all chains? runs faster if set to FALSE, but then you'll be unable to conduct convergence/mixing diagnostics

Value

in list element 'survtimes', a vector of simulated survival times (the last simulated value from the MCMC chains) in list element 'T' the MCMC chains

See Also

[covmodel](#), [survspat](#), [tpowHaz](#), [exponentialHaz](#), [gompertzHaz](#), [makehamHaz](#), [weibullHaz](#)

spatialpars

spatialpars function

Description

A function to return the mcmc chains for the spatial covariance function parameters

Usage

```
spatialpars(x)
```

Arguments

x	an object of class mcmcparsurv
---	--------------------------------

Value

the eta mcmc chains

See Also

[print.mcmcparsurv](#), [quantile.mcmcparsurv](#), [summary.mcmcparsurv](#), [vcov.mcmcparsurv](#), [frailty-lag1](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcparsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

spatsurvVignette	<i>spatsurvVignette function</i>
------------------	----------------------------------

Description

Display the introductory vignette for the spatsurv package.

Usage

```
spatsurvVignette()
```

Value

displays the vignette by calling browseURL

SPDE	<i>SPDE function</i>
------	----------------------

Description

A function to declare and evaluate an SPDE covariance function.

Usage

```
SPDE(ord)
```

Arguments

ord	the order of the model to be used, currently an integer between 1 and 3. See Lindgren 2011 paper.
-----	---

Value

an covariance function based on the SPDE model

See Also

[ExponentialCovFct](#), [covmodel](#), [CovarianceFct](#)

SPDEprec

SPDEprec function

Description

A function to used in entering elements into the precision matrix of an SPDE model. Not intended for general use.

Usage

SPDEprec(a, ord)

Arguments

a parameter a, see Lindgren et al 2011.
ord the order of the SPDE model, see Lindgren et al 2011.

Value

a function used for creating the precision matrix

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. *Journal of Statistical Software*, 77(4), 1-32, doi:10.18637/jss.v077.i04.
2. Finn Lindgren, Havard Rue, Johan Lindstrom. An explicit link between Gaussian fields and Gaussian Markov random fields: the stochastic partial differential equation approach. *Journal of the Royal Statistical Society: Series B* 73(4)

SpikedExponentialCovFct

SpikedExponentialCovFct function

Description

A function to declare and also evaluate a spiked exponential covariance function. This is an exponential covariance function with a nugget.

Usage

SpikedExponentialCovFct()

Value

the spiked exponential covariance function

See Also

[ExponentialCovFct](#), [covmodel](#), [CovarianceFct](#)

spplot1

spplot1 function

Description

A function to provide spplot-like plotting capability but NOT using trellis graphics. This function also acts as an interface for fast plotting of `SpatialPolygonsDataFrame` or `SpatialPixelsDataFrame` objects using leaflet HTML plotting capabilities to get zoomable plots with real-world context: transformation to the correct projection is done automatically.

Usage

```
spplot1(  
  x,  
  what,  
  palette = brewer.pal(5, "Oranges"),  
  breaks = NULL,  
  legpos = "topleft",  
  fun = identity,  
  include.lowest = TRUE,  
  bty = "n",  
  bg = NULL,  
  printlegend = TRUE,  
  bw = FALSE,  
  useLeaflet = FALSE,  
  urltemplate = urlTemplate("Stamen_Toner"),  
  fillOpacity = 0.5,  
  legendOpacity = 0.5,  
  OSMbg = NULL,  
  leafletLegend = TRUE,  
  alpha.f = 0.5,  
  plotinorder = FALSE,  
  legendText = NULL,  
  legendFun = NULL,  
  ...  
)
```

Arguments

x	a <code>SpatialPolygonsDataFrame</code> or a <code>SpatialPointsDataFrame</code>
what	the name of the variable to plot
palette	the palette, can either be a vector of names of colours, or a vector of colours produced for example by the <code>brewer.pal</code> function.

breaks	optional breaks for the legend, a vector of length 1 + length(palette)
legpos	the position of the legend, options are 'topleft', 'topright', 'bottomleft', 'bottomright'
fun	an optional function of the data to plot, default is the identity function
include.lowest	see ?cut
bty	see ?legend
bg	see ?legend
printlegend	logical: print the legend?
bw	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
useLeaflet	whether to use leaflet to produce a zoomable map this requires the leaflet package, available by issuing the command "devtools::install_github('rstudio/leaflet')"
urltemplate	template for leaflet map background, default is urlTemplate('Stamen-Toner'), but any valid web address for leaflet templates will work here. See ?urlTemplate.
fillOpacity	see ?addPolygons
legendOpacity	see opacity argument in function addLegend
OSMbg	optional OpenStreetMap background to add to plot, obtain this using the function getBackground
leafletLegend	logical, display the leaflet legend?
alpha.f	point transparency, see ?adjustcolor, default is 0.5
plotinorder	whether to plot in order of the size of the variable being plotted, useful for overlapping windows where small counts may obscure big counts
legendText	X
legendFun	X
...	other arguments to be passed to plot

Details

See <http://leaflet-extras.github.io/leaflet-providers/preview/> for examples of leaflet templates.

Instructions on installing the leaflet R package are available from <https://rstudio.github.io/leaflet/>

Value

either produces a plot or if useLeaflet is TRUE, returns a leaflet map widget to which further layers can be added

See Also

[urlTemplate](#), [getBackground](#), [brewer.pal](#)

spplot_compare	<i>spplot_compare function</i>
----------------	--------------------------------

Description

A function to compare two `SpatialPolgonsDataFrame` or `SpatialPointsDataFrame` objects using a unified legend for the variable of interest in both

Usage

```
spplot_compare(
  x,
  y,
  what,
  what1 = what,
  palette = brewer.pal(9, "Oranges"),
  legpos = "topleft",
  border = NA,
  fun = identity,
  t1 = "",
  t2 = "",
  bw = FALSE,
  ...
)
```

Arguments

<code>x</code>	a <code>SpatialPolgonsDataFrame</code> or a <code>SpatialPointsDataFrame</code>
<code>y</code>	a <code>SpatialPolgonsDataFrame</code> or a <code>SpatialPointsDataFrame</code>
<code>what</code>	the name of the variable from <code>x</code> to plot
<code>what1</code>	the name of the variable from <code>y</code> to plot. default is to plot the variable of the same name
<code>palette</code>	the palette, can either be a vector of names of colours, or a vector of colours produced for example by the <code>brewer.pal</code> function.
<code>legpos</code>	the position of the legend, options are 'topleft', 'topright', 'bottomleft', 'bottomright'
<code>border</code>	see <code>?spplot</code>
<code>fun</code>	an optional function of the data to plot, default is the identity function
<code>t1</code>	title for the plot of <code>x</code>
<code>t2</code>	title for the plot of <code>y</code>
<code>bw</code>	Logical. Plot in black/white/greyscale? Default is to produce a colour plot. Useful for producing plots for journals that do not accept colour plots.
<code>...</code>	other arguments to be passed to the plot function

Value

produces a plot comparing $x[[\text{what}]]$ and $y[[\text{what1}]]$

Summarise

Summarise function

Description

A function to completely summarise the output of an object of class `mcmcspatsurv`.

Usage

```
Summarise(
  obj,
  digits = 3,
  scientific = -3,
  inclIntercept = FALSE,
  printmode = "LaTeX",
  displaymode = "console",
  ...
)
```

Arguments

<code>obj</code>	an object produced by a call to <code>lgcpPredictSpatialPlusPars</code> , <code>lgcpPredictAggregateSpatialPlusPars</code> , <code>lgcpPredictSpatioTemporalPlusPars</code> or <code>lgcpPredictMultitypeSpatialPlusPars</code>
<code>digits</code>	see the option "digits" in <code>?format</code>
<code>scientific</code>	see the option "scientific" in <code>?format</code>
<code>inclIntercept</code>	logical: whether to summarise the intercept term, default is <code>FALSE</code> .
<code>printmode</code>	the format of the text to return, can be 'LaTeX' (the default) or 'text' for plain text.
<code>displaymode</code>	default is 'console' alternative is 'rstudio'
<code>...</code>	other arguments passed to the function "format"

Value

A text summary, that can be pasted into a LaTeX document and later edited.

summary.mcmc	<i>summary.mcmc function</i>
--------------	------------------------------

Description

summary of an mcmc iterator print out values of an iterator and reset it. DONT call this in a loop that uses this iterator - it will reset it. And break.

Usage

```
## S3 method for class 'mcmc'
summary(object, ...)
```

Arguments

object	an mcmc iterator
...	other args

summary.mcmcpatsurv	<i>summary.mcmcpatsurv function</i>
---------------------	-------------------------------------

Description

A function to return summary tables from an MCMC run

Usage

```
## S3 method for class 'mcmcpatsurv'
summary(object, probs = c(0.5, 0.025, 0.975), ...)
```

Arguments

object	an object inheriting class mcmcpatsurv
probs	vector of quantiles to return
...	additional arguments

Value

summary tables to the console

See Also

[print.mcmcpatsurv](#), [quantile.mcmcpatsurv](#), [vcov.mcmcpatsurv](#), [frailtylag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcpatsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

surv3d

*Spatial Survival Plot in 3D***Description**

Do a 3d plot of spatial survival data

Usage

```
surv3d(
  spp,
  ss,
  lwd = 2,
  lcol = "black",
  lalpha = 1,
  pstyle = c("point", "text"),
  psize = c(20, 10),
  pcol = c("red", "black"),
  ptext = c("X", ""),
  palpha = 1,
  title = "Spatial Survival",
  basegrid = TRUE,
  baseplane = TRUE
)
```

Arguments

spp	A spatial points data frame
ss	A Surv object (with right-censoring)
lwd	Line width for stems
lcol	Line colour for stems
lalpha	Opacity for stems
pstyle	Point style "point" or "text"
psize	Vector of length 2 for uncensored/censored points size
pcol	Vector of length 2 for uncensored/censored points colours
ptext	Vector of length 2 for uncensored/censored text characters
palpha	Opacity for points/text
title	Main title for plot
basegrid	add a grid at t=0
baseplane	add a plane at t=0

Details

Uses rgl graphics to make a spinny zoomy plot

Value

nothing

Author(s)

Barry S Rowlingson

Examples

```
## Not run:
require(sp)
require(survival)
d = data.frame(
  x=runif(40)*1.5,
  y = runif(40),
  age=as.integer(20+30*runif(40)),
  sex = sample(c("M", "F"), 40, TRUE)
)
coordinates(d)~x+y
d$surv = Surv(as.integer(5+20*runif(40)), runif(40)>.9)
clear3d(); surv3d(d, d$surv, baseplane=TRUE, basegrid=TRUE)
clear3d(); surv3d(d, d$surv, baseplane=TRUE, basegrid=TRUE, pstyle="t", lalpha=0.5, lwd=3, palpha=1)

## End(Not run)
```

survival_PP

survival_PP function

Description

A function to compute an individual's survival function

Usage

```
survival_PP(inputs)
```

Arguments

inputs inputs for the function including the model matrix, frailties, fixed effects and the parameters of the baseline hazard derived from this model

Value

the survival function for the individual

survspat *survspat function*

Description

A function to run a Bayesian analysis on censored spatial survival data assuming a proportional hazards model using an adaptive Metropolis-adjusted Langevin algorithm.

Usage

```
survspat(
  formula,
  data,
  dist,
  cov.model,
  mcmc.control,
  priors,
  shape = NULL,
  ids = list(shpid = NULL, dataid = NULL),
  control = inference.control(grided = FALSE),
  boundingbox = NULL
)
```

Arguments

formula	the model formula in a format compatible with the function <code>flexsurvreg</code> from the <code>flexsurv</code> package
data	a <code>SpatialPointsDataFrame</code> object containing the survival data as one of the columns OR for polygonal data a <code>data.frame</code> , in which case, the argument <code>shape</code> must also be supplied
dist	choice of distribution function for baseline hazard. Current options are: <code>exponentialHaz</code> , <code>weibullHaz</code> , <code>gompertzHaz</code> , <code>makehamHaz</code> , <code>tpowHaz</code>
cov.model	an object of class <code>covmodel</code> , see <code>?covmodel</code> <code>?ExponentialCovFct</code> or <code>?SpikedExponentialCovFct</code>
mcmc.control	mcmc control parameters, see <code>?mcmcpars</code>
priors	an object of class <code>Priors</code> , see <code>?mcmcPriors</code>
shape	when <code>data</code> is a <code>data.frame</code> , this can be a <code>SpatialPolygonsDataFrame</code> , or a <code>SpatialPointsDataFrame</code> , used to model spatial variation at the small region level. The regions are the polygons, or they represent the (possibly weighted) centroids of the polygons.
ids	named list entry <code>shpid</code> character string giving name of variable in <code>shape</code> to be matched to variable <code>dataid</code> in <code>data</code> . <code>dataid</code> is the second entry of the named list.
control	additional control parameters, see <code>?inference.control</code>
boundingbox	optional bounding box over which to construct computational grid, supplied as an object on which the function <code>'bbox'</code> returns the bounding box

Value

an object inheriting class 'mcmcsurv' for which there exist methods for printing, summarising and making inference from.

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. Journal of Statistical Software, 77(4), 1-32, doi:10.18637/jss.v077.i04.

See Also

[tpowHaz](#), [exponentialHaz](#), [gompertzHaz](#), [makehamHaz](#), [weibullHaz](#), [covmodel](#), [linkExponentialCovFct](#), [SpikedExponentialCovFct](#), [mcmcpars](#), [mcmcPriors](#), [inference.control](#)

survspatNS

survspatNS function

Description

A function to perform maximum likelihood inference for non-spatial survival data.

Usage

```
survspatNS(formula, data, dist, control = inference.control())
```

Arguments

formula	the model formula in a format compatible with the function flexsurvreg from the flexsurv package
data	a SpatialPointsDataFrame object containing the survival data as one of the columns
dist	choice of distribution function for baseline hazard. Current options are: exponentialHaz, weibullHaz, gompertzHaz, makehamHaz, tpowHaz
control	additional control parameters, see ?inference.control

Value

an object inheriting class 'mcmcsurv' for which there exist methods for printing, summarising and making inference from.

References

1. Benjamin M. Taylor and Barry S. Rowlingson (2017). spatsurv: An R Package for Bayesian Inference with Spatial Survival Models. Journal of Statistical Software, 77(4), 1-32, doi:10.18637/jss.v077.i04.

See Also

[tpowHaz](#), [exponentialHaz](#), [gompertzHaz](#), [makehamHaz](#), [weibullHaz](#), [covmodel](#), [linkExponentialCovFct](#), [SpikedExponentialCovFct](#), [mcmcpars](#), [mcmcPriors](#), [inference.control](#)

textSummary	<i>textSummary function</i>
-------------	-----------------------------

Description

A function to print a text description of the inferred parameters beta and eta from a call to the function `lgcpPredictSpatialPlusPars`, `lgcpPredictAggregateSpatialPlusPars`, `lgcpPredictSpatioTemporalPlusPars` or `lgcpPredictMultitypeSpatialPlusPars`

Usage

```
textSummary(
  obj,
  digits = 3,
  scientific = -3,
  inclIntercept = FALSE,
  printmode = "LaTeX",
  ...
)
```

Arguments

<code>obj</code>	an object produced by a call to <code>lgcpPredictSpatialPlusPars</code> , <code>lgcpPredictAggregateSpatialPlusPars</code> , <code>lgcpPredictSpatioTemporalPlusPars</code> or <code>lgcpPredictMultitypeSpatialPlusPars</code>
<code>digits</code>	see the option "digits" in <code>?format</code>
<code>scientific</code>	see the option "scientific" in <code>?format</code>
<code>inclIntercept</code>	logical: whether to summarise the intercept term, default is FALSE.
<code>printmode</code>	the format of the text to return, can be 'LaTeX' (the default) or 'text' for plain text.
<code>...</code>	other arguments passed to the function "format"

Value

A text summary, that can be pasted into a LaTeX document and later edited.

timevaryingPL	<i>timevaryingPL function</i>
---------------	-------------------------------

Description

A function to

Usage

```
timevaryingPL(  
  formula,  
  t0,  
  t,  
  delta,  
  dist,  
  data,  
  ties = "Efron",  
  optimcontrol = NULL  
)
```

Arguments

formula	a formula of the form 'S ~ coef1 + coef2' etc the object S will be created
t0	X
t	X
delta	censoring indicator a vector of 1 for an event and 0 for censoring
dist	X
data	X
ties	X default is Efron
optimcontrol	X

Value

...

 tpowHaz

tpowHaz function

Description

A function to define a parametric proportional hazards model where the baseline hazard is taken from the 'powers of t' model. This function returns an object inheriting class 'basehazardspec', list of functions 'distinfo', 'basehazard', 'gradbasehazard', 'hessbasehazard', 'cumbasehazard', 'gradcumbasehazard', 'hesscumbasehazard' and 'densityquantile'

Usage

```
tpowHaz(powers)
```

Arguments

powers a vector of powers of t. These are powers are treated as fixed in estimation routines and it is assumed that the log cumulative baseline hazard is a linear combination of these powers of t

Details

The `distinfo` function is used to provide basic distribution specific information to other `spatsurv` functions. The user is required to provide the following information in the returned list: `npars`, the number of parameters in this distribution; `parnames`, the names of the parameters; `trans`, the transformation scale on which the priors will be provided; `itrans`, the inverse transformation function that will be applied to the parameters before the hazard, and other functions are evaluated; `jacobian`, the derivative of the inverse transformation function with respect to each of the parameters; and `hessian`, the second derivatives of the inverse transformation function with respect to each of the parameters – note that currently the package `spatsurv` only allows the use of functions where the parameters are transformed independently.

The `basehazard` function is used to evaluate the baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradbasehazard` function is used to evaluate the gradient of the baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hessbasehazard` function is used to evaluate the Hessian of the baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `cumbasehazard` function is used to evaluate the cumulative baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradcumbasehazard` function is used to evaluate the gradient of the cumulative baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hesscumbasehazard` function is used to evaluate the Hessian of the cumulative baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `densityquantile` function is used to return quantiles of the density function. This is NOT REQUIRED for running the MCMC, merely for us in post-processing with the `predict` function where `type` is `'densityquantile'`. In the case of the Weibull model for the baseline hazard, it can be shown that the `q`-th quantile is:

Value

an object inheriting class `'basehazardspec'`

See Also

[exponentialHaz](#), [gompertzHaz](#), [makehamHaz](#), [weibullHaz](#)

transformweibull	<i>transformweibull function</i>
------------------	----------------------------------

Description

A function to back-transform estimates of the parameters of the weibull baseline hazard function, so they are commensurate with R's inbuilt density functions. Transforms from (shape, scale) to (alpha, lambda)

Usage

```
transformweibull(x)
```

Arguments

`x` a vector of paramters

Value

the transformed parameters. For the weibull model, this is the back-transform from `'alpha'` and `'lambda'` to `'shape'` `'scale'` (see `?dweibull`).

TwoWayHazAdditive	<i>TwoWayHazAdditive function</i>
-------------------	-----------------------------------

Description

A function to

Usage

```
TwoWayHazAdditive(bhlist, bhtime, bhfix, MLimits = NULL)
```

Arguments

bhlist	X
bhtime	X
bhfix	X
MLimits	X

Value

...

txtProgressBar2	<i>A text progress bar with label</i>
-----------------	---------------------------------------

Description

This is the base txtProgressBar but with a little modification to implement the label parameter for style=3. For full info see txtProgressBar

Usage

```
txtProgressBar2(
  min = 0,
  max = 1,
  initial = 0,
  char = "=",
  width = NA,
  title = "",
  label = "",
  style = 1
)
```

Arguments

min	min value for bar
max	max value for bar
initial	initial value for bar
char	the character (or character string) to form the progress bar.
width	progress bar width
title	ignored
label	text to put at the end of the bar
style	bar style

urlTemplate	<i>urlTemplate function</i>
-------------	-----------------------------

Description

A function to return a url for a leaflet template for use as map backgrounds with the spplot1 function.

Usage

```
urlTemplate(name = "Stamen_Toner")
```

Arguments

name	name of the template to use, the default is 'Stamen_Toner'
------	--

Details

Possible templates are: OpenStreetMap_Mapnik, OpenStreetMap_BlackAndWhite, OpenStreetMap_DE, OpenStreetMap_France, OpenStreetMap_HOT, OpenTopoMap, Thunderforest_OpenCycleMap, Thunderforest_Transport, Thunderforest_Landscape, Thunderforest_Outdoors, OpenMapSurfer_Roads, OpenMapSurfer_Grayscale, Hydda_Full, Hydda_Base, MapQuestOpen_OSM, Stamen_Toner, Stamen_TonerBackground, Stamen_TonerLite, Stamen_Watercolor, Stamen_Terrain, Stamen_TerrainBackground, Stamen_TopOSMRelief, Esri_WorldStreetMap, Esri_WorldTopoMap, Esri_WorldImagery, Esri_WorldTerrain, Esri_WorldShadedRelief, Esri_WorldPhysical, Esri_OceanBasemap, Esri_NatGeoWorldMap, Esri_WorldGrayCanvas, Acetate_all, Acetate_terrain, HERE_satelliteDay, HERE_hybridDayMobile, HERE_hybridDay

See <http://leaflet-extras.github.io/leaflet-providers/preview/> for other leaflet templates

Value

url for the leaflet template

vcov.mcmcsurv *vcov.mcmcsurv function*

Description

A function to return the variance covariance matrix of the parameters beta, omega and eta

Usage

```
## S3 method for class 'mcmcsurv'
vcov(object, ...)
```

Arguments

object an object inheriting class mcmcsurv
 ... other arguments, not used here

Value

the variance covariance matrix of the parameters beta, omega and eta

See Also

[print.mcmcsurv](#), [quantile.mcmcsurv](#), [summary.mcmcsurv](#), [frailtylag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

vcov.mlspatsurv *vcov.mlspatsurv function*

Description

A function to return the variance covariance matrix of the parameters beta, omega and eta

Usage

```
## S3 method for class 'mlspatsurv'
vcov(object, ...)
```

Arguments

object an object inheriting class mcmcsurv
 ... other arguments, not used here

Value

the variance covariance matrix of the parameters beta, omega and eta

See Also

[print.mcmcspsurv](#), [quantile.mcmcspsurv](#), [summary.mcmcspsurv](#), [frailtylag1](#), [spatialpars](#), [hazardpars](#), [fixedpars](#), [randompars](#), [baselinehazard](#), [predict.mcmcspsurv](#), [priorposterior](#), [posteriorcov](#), [MCE](#), [hazardexceedance](#)

weibullHaz

weibullHaz function

Description

A function to define a parametric proportional hazards model where the baseline hazard is taken from the Weibull model. This function returns an object inheriting class 'basehazardspec', list of functions 'distinfo', 'basehazard', 'gradbasehazard', 'hessbasehazard', 'cumbasehazard', 'gradcumbasehazard', 'hesscumbasehazard' and 'densityquantile'

Usage

```
weibullHaz(MLinits = NULL)
```

Arguments

MLinits initial values for optim, default is NULL

Details

The `distinfo` function is used to provide basic distribution specific information to other `spatsurv` functions. The user is required to provide the following information in the returned list: `npars`, the number of parameters in this distribution; `parnames`, the names of the parameters; `trans`, the transformation scale on which the priors will be provided; `itrans`, the inverse transformation function that will be applied to the parameters before the hazard, and other functions are evaluated; `jacobian`, the derivative of the inverse transformation function with respect to each of the parameters; and `hessian`, the second derivatives of the inverse transformation function with respect to each of the parameters – note that currently the package `spatsurv` only allows the use of functions where the parameters are transformed independently.

The `basehazard` function is used to evaluate the baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradbasehazard` function is used to evaluate the gradient of the baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hessbasehazard` function is used to evaluate the Hessian of the baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `cumbasehazard` function is used to evaluate the cumulative baseline hazard function for the distribution of interest. It returns a function that accepts as input a vector of times, `t` and returns a vector.

The `gradcumbasehazard` function is used to evaluate the gradient of the cumulative baseline hazard function with respect to the parameters, this typically returns a vector. It returns a function that accepts as input a vector of times, `t`, and returns a matrix.

The `hesscumbasehazard` function is used to evaluate the Hessian of the cumulative baseline hazard function. It returns a function that accepts as input a vector of times, `t` and returns a list of hessian matrices corresponding to each `t`.

The `densityquantile` function is used to return quantiles of the density function. This is NOT REQUIRED for running the MCMC, merely for us in post-processing with the `predict` function where `type` is `'densityquantile'`. In the case of the Weibull model for the baseline hazard, it can be shown that the `q`-th quantile is:

Value

an object inheriting class `'basehazardspec'`

See Also

[tpowHaz](#), [exponentialHaz](#), [gompertzHaz](#), [makehamHaz](#)

YfromGamma

YfromGamma function

Description

A function to change Gammas (white noise) into Ys (spatially correlated noise). Used in the MALA algorithm.

Usage

```
YfromGamma(Gamma, invrootQeigs, mu)
```

Arguments

<code>Gamma</code>	Gamma matrix
<code>invrootQeigs</code>	inverse square root of the eigenvectors of the precision matrix
<code>mu</code>	parameter of the latent Gaussian field

Value

Y

YFromGamma_SPDE	<i>YFromGamma_SPDE function</i>
-----------------	---------------------------------

Description

A function to go from Gamma to Y

Usage

```
YFromGamma_SPDE(gamma, U, mu)
```

Arguments

gamma	Gamma
U	upper Cholesky matrix
mu	the mean

Value

the value of Y for the given Gamma

References

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2. Finn Lindgren, Havard Rue, Johan Lindstrom. An explicit link between Gaussian fields and Gaussian Markov random fields: the stochastic partial differential equation approach. *Journal of the Royal Statistical Society: Series B* 73(4)

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