

Package ‘INLA’

July 10, 2021

Type Package

Title Full Bayesian Analysis of Latent Gaussian Models using Integrated Nested Laplace Approximations

Description Full Bayesian analysis of latent Gaussian models using Integrated Nested Laplace Approximation. It is a front-end to the inla-program.

Depends R (≥ 3.5),
Matrix ($\geq 1.3-0$),
foreach,
parallel,
sp ($\geq 1.4-2$)

Suggests Deriv,
Ecdat,
HKprocess,
MatrixModels,
Rgraphviz,
deldir,
devtools,
doParallel,
dplyr,
evd,
fields,
ggplot2,
graph,
gridExtra,
knitr,
markdown,
matrixStats,
mlogit,
mpoly,
mvtnorm,
numDeriv,
orthopolynom,
pixmap,
rgdal,
rgeos,
rgl,
rmarkdown,
shiny,
sn,

spdep,
splancs,
symmoments,
testthat,
tools

Imports graphics,
grDevices,
methods,
splines,
stats,
utils

VignetteBuilder knitr

BuildVignettes true

LazyData true

License GPL (>= 2) + file LICENSE

RoxygenNote 7.1.1

Encoding UTF-8

StagedInstall no

Collate 'INLA-package.R'
'agaussian.R'
'ar.R'
'barrier.R'
'binary.R'
'binary.install.R'
'changelog.R'
'collect.results.R'
'compare.results.R'
'coxph.R'
'cpo.R'
'create.data.file.R'
'cut.R'
'cygwin.R'
'debug.graph.R'
'dev.new.R'
'display.matrix.R'
'doc.R'
'expand.dataframe.R'
'export-class.R'
'f.R'
'fgn.R'
'finn.R'
'fmeshier-io.R'
'fmeshier.R'
'graph.convert.R'
'graph.matrix.R'
'group.R'
'hyper.R'
'hyperpar.R'
'idx.R'
'inla.R'

'inlaEnv.R'
'interpret.formula.R'
'jmargin.R'
'jp.R'
'knmodels.R'
'knmodels.sample.R'
'lattice2node.R'
'lincomb.R'
'link-functions.R'
'list-models.R'
'load.R'
'marginal.R'
'mdata.R'
'merge.R'
'mesh.R'
'mesh.components.R'
'meshassessment.R'
'meshbuilder.R'
'model-wrapper.R'
'models-generate.R'
'models.R'
'nmix.lambda.fitted.R'
'obsolete.R'
'options.R'
'os.R'
'over_sp_mesh.R'
'pardo.R'
'pc-alpha.R'
'pc-ar.R'
'pc-bym.R'
'pc-cor0.R'
'pc-cor1.R'
'pc-cormat.R'
'pc-gamma.R'
'pc-gammacount.R'
'pc-gev.R'
'pc-multvar.R'
'pc-prec.R'
'pc-sn.R'
'pc-t.R'
'plot.R'
'posterior.sample.R'
'print.R'
'priors.used.R'
'prune.R'
'q.R'
'qinv.R'
'qreordering.R'
'qsample.R'
'qsolve.R'
'quantile-regression.R'
'read.graph.R'

'remote.R'
 'reorderings.R'
 'rerun.R'
 'residuals.R'
 'rgeneric.R'
 'sampler.R'
 'sandbox.R'
 'scale.model.R'
 'sections.R'
 'set.default.arguments.R'
 'sm.R'
 'spde.common.R'
 'spde1.R'
 'spde2.R'
 'spde3.R'
 'spmesh.R'
 'startup.R'
 'summary.R'
 'surv.R'
 'upgrade.R'
 'utils.R'
 'version.R'

Roxygen list(markdown = TRUE)

Version 21.07.10-1

Date Sat Jul 10 02:54:06 PM +03 2021 (Version_21.07.10-1)

R topics documented:

INLA-package	7
aggregate.gaussian	8
as.inla.mesh.segment	9
BivMetaAnalysis	10
Cancer	11
control.bgev.default	11
control.compute	12
control.expert	13
control.family	14
control.fixed	15
control.group	16
control.hazard	17
control.inla	18
control.lincomb	21
control.link	22
control.mix	23
control.mode	24
control.pardiso.default	24
control.predictor	25
control.results	26
control.update	27
cut	28
debug.graph	29

Drivers	30
Epil	30
extract.groups	31
f	32
fgn	36
geobugs2inla	37
Germany	38
graph2matrix	38
idx	39
inla	40
inla-class	45
inla.ar.pacf2phi	45
inla.as.sparse	46
inla.as.wkt_tree.wkt	47
inla.barrier	48
inla.barrier.pcmatern	49
inla.binary.install	50
inla.changelog	51
inla.collect.results	51
inla.compare.results	52
inla.coxph	53
inla.cpo	54
inla.CRS	55
inla.CRSargs	57
inla.dev.new	58
inla.diameter	59
inla.doc	60
inla.extract.el	60
inla.fmesher.smorg	61
inla.generate.colors	62
inla.get.inlaEnv	63
inla.group	63
inla.has_PROJ6	64
inla.hyperpar	65
inla.hyperpar.sample	66
inla.identical.CRS	67
inla.knmodels	68
inla.knmodels.sample	70
inla.ks.plot	71
inla.list.models	72
inla.load	73
inla.matern.cov	73
inla.mdata	74
inla.merge	75
inla.mesh.1d	77
inla.mesh.1d.A	77
inla.mesh.2d	78
inla.mesh.assessment	80
inla.mesh.assessment	80
inla.mesh.basis	81
inla.mesh.boundary	82
inla.mesh.components	83

<code>inla.mesh.components</code>	84
<code>inla.mesh.create</code>	85
<code>inla.mesh.deriv</code>	87
<code>inla.mesh.fem</code>	88
<code>inla.mesh.lattice</code>	89
<code>inla.mesh.map.lim</code>	90
<code>inla.mesh.project</code>	91
<code>inla.mesh.query</code>	93
<code>inla.mesh.segment</code>	94
<code>inla.models</code>	96
<code>inla.nmix.lambda.fitted</code>	232
<code>inla.nonconvex.hull</code>	234
<code>inla.option</code>	235
<code>inla.over_sp_mesh</code>	237
<code>inla.pardiso</code>	238
<code>inla.priors.used</code>	238
<code>inla.prune</code>	239
<code>inla.qstat</code>	239
<code>inla.reorderings</code>	241
<code>inla.rerun</code>	241
<code>inla.row.kron</code>	242
<code>inla.sample</code>	243
<code>inla.simplify.curve</code>	247
<code>inla.spde.make.A</code>	248
<code>inla.spde.make.block.A</code>	249
<code>inla.spde.make.index</code>	250
<code>inla.spde.models</code>	251
<code>inla.spde.precision</code>	252
<code>inla.spde.result</code>	253
<code>inla.spde.sample</code>	256
<code>inla.spde1.create</code>	256
<code>inla.spde2.generic</code>	258
<code>inla.spde2.matern</code>	260
<code>inla.spde2.matern.sd.basis</code>	263
<code>inla.spde2.pcmatern</code>	264
<code>inla.spTransform</code>	267
<code>inla.sp_get_crs</code>	269
<code>inla.ssh.copy.id</code>	270
<code>inla.stack.remove.unused</code>	270
<code>inla.surv</code>	275
<code>inla.upgrade</code>	276
<code>inla.version</code>	277
<code>inla.wkt_is_geocent</code>	278
<code>joint.marginal</code>	280
<code>jp.define</code>	281
<code>Kidney</code>	282
<code>lattice2node</code>	283
<code>Leuk</code>	284
<code>lines.inla.mesh.segment</code>	285
<code>link</code>	286
<code>make.lincomb</code>	287
<code>marginal</code>	288

meshbuilder	290
Munich	291
nwEngland	292
Oral	292
param2.matern.orig	293
pc.alphaw	294
pc.ar	295
pc.cor0	295
pc.cor1	296
pc.cormat	297
pc.ddof	299
pc.gamma	300
pc.gammacount	301
pc.gevtail	302
pc.multvar	303
pc.prec	304
pc.sn	305
plot.inla	306
plot.inla.CRS	307
plot.inla.mesh	309
plot.inla.trimesh	310
PRborder	311
print.inla	312
PRprec	312
qinv	322
qreordering	323
qsample	324
qsolve	326
read.graph	327
rgeneric.define	329
Salm	330
scale.model	331
Scotland	332
Seeds	333
SPDEtoy	333
summary.inla	334
summary.inla.mesh	335
Surg	335
SurvSim	336
Tokyo	336
Zambia	337
Index	338

Description

Package to perform full Bayesian analysis on generalised additive mixed models using Integrated Nested Laplace Approximations.

Details**Package** INLA**Version** Currently, this package uses a YY.MM.DD versioning system, and is in heavy development. See <https://github.com/hrue/r-inla/> and <https://www.r-inla.org>**License** GPL2See the web-site <https://www.r-inla.org> for further details.**Author(s)**

Havard Rue, Sara Martino, Finn Lindgren, Daniel Simpson and Andrea Riebler

 aggregate.gaussian *Aggregate Gaussian into an equivalent observation*

Description

Aggregate Gaussians observed with the same mean and precision, into an eqv triplet, for use with family="agaussian"

Usage

```
inla.agaussian(y, s = NULL)
```

Arguments

y	Repeated observations. If y is a matrix, then each row represents repeated observations. If y is a list, then each element of the list is a vector of repeated observations. If y is a vector, then the whole vector represents repeated observations. The optional scaling s, must have the same format as y, ie matrix or vector. NA's in y (and s) are removed and not used or counted. If s is given, then the NA-pattern in y and s must be the same.
s	Optional fixed scaling of the precisions. Must be in the same format as y, and have the same NA-pattern. See the documentation for details.

Value

The output is a inla.mdata-object ready for use with family="agaussian". See the example in the documentation.

Author(s)Havard Rue <hrue@r-inla.org>**Examples**

```
A = matrix(1:25,5,5)
inla.agaussian(A)

A[1,-1] = NA
A[2,-(2:3)] = NA
inla.agaussian(A)
```

as.inla.mesh.segment *Convert sp curve objects to inla.mesh.segment objects.*

Description

Convert sp curve objects to inla.mesh.segment objects.

Usage

```
as.inla.mesh.segment(sp, ...)

inla.sp2segment(sp, ...)

## S3 method for class 'SpatialPoints'
as.inla.mesh.segment(sp, reverse = FALSE, grp = NULL, is.bnd = TRUE, ...)

## S3 method for class 'SpatialPointsDataFrame'
as.inla.mesh.segment(sp, ...)

## S3 method for class 'Line'
as.inla.mesh.segment(sp, reverse = FALSE, crs = NULL, ...)

## S3 method for class 'Lines'
as.inla.mesh.segment(sp, join = TRUE, crs = NULL, ...)

## S3 method for class 'SpatialLines'
as.inla.mesh.segment(sp, join = TRUE, grp = NULL, ...)

## S3 method for class 'SpatialLinesDataFrame'
as.inla.mesh.segment(sp, ...)

## S3 method for class 'SpatialPolygons'
as.inla.mesh.segment(sp, join = TRUE, grp = NULL, ...)

## S3 method for class 'SpatialPolygonsDataFrame'
as.inla.mesh.segment(sp, ...)

## S3 method for class 'Polygons'
as.inla.mesh.segment(sp, join = TRUE, crs = NULL, ...)

## S3 method for class 'Polygon'
as.inla.mesh.segment(sp, crs = NULL, ...)
```

Arguments

sp	An sp polygon object of class Polygon, Polygons, SpatialPolygons, or SpatialPolygonsDataFra
...	Additional arguments passed on to other methods.
reverse	Logical, indicating if the line sequence should be traversed backwards.
grp	Group ID specification for each polygon, as used by inla.mesh.segment() , one ID per polygon.

crs An optional CRS or `inla.CRS` object
join If TRUE, join multiple polygons into a single segment (possibly non-simply connected).

Value

A `inla.mesh.segment()` object, or a list of `inla.mesh.segment()` objects.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

`inla.mesh.segment()`

BivMetaAnalysis

Bivariate Meta Analysis

Description

Data are taken from a meta-analysis to compare the utility of three types of diagnostic imaging - lymphangiography (LAG), computed tomography (CT) and magnetic resonance (MR) - to detect lymph node metastases in patients with cervical cancer. The dataset consists of a total of 46 studies: the first 17 for LAG, the following 19 for CT and the last 10 for MR.

Usage

```
BivMetaAnalysis
```

Format

A data frame with 92 observations on the following 9 variables.

N a numeric vector

Y a numeric vector

diid a numeric vector

lag.tp a numeric vector

lag.tn a numeric vector

ct.tp a numeric vector

ct.tn a numeric vector

mr.tp a numeric vector

mr.tn a numeric vector

References

J. Scheidler and H. Hricak and K. K. Yu and L. Subak and M. R. Segal, "Radiological evaluation of lymph node metastases in patients with cervical cancer: a meta-analysis", JAMA 1997

Examples

```
data(BivMetaAnalysis)
```

Cancer *~~ data name/kind ... ~~*

Description

~~ A concise (1-5 lines) description of the dataset. ~~

Format

A data frame with 6690 observations on the following 4 variables.

Y Number of cases

N a numeric vector

Age a numeric vector

region a numeric vector

References

Rue, H and Held, L. (2005) *Gaussian Markov Random Fields - Theory and Applications* Chapman and Hall

`control.bgev.default` *Control variables in control.bgev.default*

Description

Control variables in `control.bgev.default` for use in `inla`

Usage

```
inla.set.control.bgev.default.default(...)
control.bgev.default(beta.ab, q.location, q.mix, q.spread)
```

Arguments

<code>...</code>	Possible arguments
<code>q.location</code>	The quantile level for the location parameter
<code>q.spread</code>	The quantile level for the spread parameter (must be < 0.5)
<code>q.mix</code>	The lower and upper quantile level for the mixing function
<code>beta.ab</code>	The parameters a and b in the Beta mixing function

Value

The `control.bgev`-list is set within the corresponding `control.family`-list as control parameters to the `family="bgev"`. The function `control.bgev.default` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.bgev.default.default` returns a list with all the default values of all parameters within this control statement.

See Also

`control.update`, `control.lincomb`, `control.group`, `control.mix`, `control.link`, `control.expert`, `control.compute`, `control.pardiso.default`, `control.bgev.default`, `control.family`, `control.fixed`, `control.inla`, `control.predictor`, `control.mode`, `control.hazard`, `inla`

<code>control.compute</code>	<i>Control variables in control.compute</i>
------------------------------	---

Description

Control variables in `control.compute` for use in `inla`

Usage

```
inla.set.control.compute.default(...)
control.compute(config, cpo, dic, graph, hyperpar, mlik, openmp.strategy, po, q, return.marginals,
```

Arguments

<code>...</code>	Possible arguments
<code>openmp.strategy</code>	The computational strategy to use: 'small', 'medium', 'large', 'huge', 'default' and 'pardiso'.
<code>hyperpar</code>	A boolean variable if the marginal for the hyperparameters should be computed. Default TRUE.
<code>return.marginals</code>	A boolean variable if the marginals for the latent field should be returned (although it is computed). Default TRUE
<code>return.marginals.predictor</code>	A boolean variable if the marginals for the linear predictor should be returned (although it is computed). Default FALSE
<code>dic</code>	A boolean variable if the DIC-value should be computed. Default FALSE.
<code>mlik</code>	A boolean variable if the marginal likelihood should be computed. Default TRUE.
<code>cpo</code>	A boolean variable if the cross-validated predictive measures (cpo, pit) should be computed (default FALSE)
<code>po</code>	A boolean variable if the predictive ordinate should be computed (default FALSE)
<code>waic</code>	A boolean variable if the Watanabe-Akaike information criteria should be computed (default FALSE)
<code>q</code>	A boolean variable if binary images of the precision matrix, the reordered precision matrix and the Cholesky triangle should be generated. (Default FALSE.)
<code>config</code>	A boolean variable if the internal GMRF approximations be stored. (Default FALSE. EXPERIMENTAL)
<code>smtsp</code>	The sparse-matrix solver, one of 'default', 'taucs', 'band' or 'pardiso' (default <code>inla.getoption("smtsp")</code>). <code>smtsp='pardiso'</code> implies <code>openmp.strategy='pardiso'</code> .
<code>graph</code>	A boolean variable if the graph itself should be returned. (Default FALSE.)

Value

The function `control.compute` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.compute.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

<code>control.expert</code>	<i>Control variables in control.expert</i>
-----------------------------	--

Description

Control variables in `control.expert` for use in `inla`

Usage

```
inla.set.control.expert.default(...)
control.expert(cpo.idx, cpo.manual, disable.gaussian.check, jp)
```

Arguments

<code>...</code>	Possible arguments
<code>cpo.manual</code>	A boolean variable to decide if the <code>inla</code> -program is to be runned in a manual- <code>cpo</code> -mode. (EXPERT OPTION: DO NOT USE)
<code>cpo.idx</code>	The index/indices of the data point(s) to remove. (EXPERT OPTION: DO NOT USE)
<code>disable.gaussian.check</code>	Disable the check for fast computations with a Gaussian likelihood and identity link (default FALSE)
<code>jp</code>	An object of class <code>inla.jp</code> defining a joint prior

Value

The function `control.expert` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.expert.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.family	<i>Control variables in control.family</i>
----------------	--

Description

Control variables in control.family for use in inla

Usage

```
inla.set.control.family.default(...)
control.family(beta.censor.value, cenpoisson.I, control.bgev, control.link, control.mix, dummy, fi
```

Arguments

...	Possible arguments
dummy	A dummy argument that can be used as a workaround
hyper	Definition of the hyperparameters
initial	(OBSOLETE!) Initial value for the hyperparameter(s) of the likelihood in the internal scale.
prior	(OBSOLETE!) The name of the prior distribution(s) for othe hyperparameter(s).
param	(OBSOLETE!) The parameters for the prior distribution
fixed	(OBSOLETE!) Boolean variable(s) to say if the hyperparameter(s) is fixed or random.
link	(OBSOLETE! Use control.link=list(model=) instead.) The link function to use.
sn.shape.max	Maximum value for the shape-parameter for Skew Normal observations (default 5.0)
gev.scale.xi	(Expert option, do not use unless you know what you are doing.) The internal scaling of the shape-parameter for the GEV distribution. (default 0.1)
control.bgev	See ?control.bgev
cenpoisson.I	The censoring interval for the censored Poisson
beta.censor.value	The censor value for the Beta-likelihood ($0 \leq \text{beta.censor.value} < 1/2$)
variant	This variable is used to give options for various variants of the likelihood, like chosing different parameterisations for example. See the relevant likelihood documentations for options (does only apply to some likelihoods).
control.mix	See ?control.mix
control.link	See ?control.link

Value

The function control.family is used to TAB-complete arguments and returns a list of given arguments. The function inla.set.control.family.default returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.fixed	<i>Control variables in control.fixed</i>
---------------	---

Description

Control variables in control.fixed for use in inla

Usage

```
inla.set.control.fixed.default(...)
control.fixed(cdf, compute, correlation.matrix, expand.factor.strategy, mean, mean.intercept, prec)
```

Arguments

...	Possible arguments
cdf	A list of values to compute the CDF for, for all fixed effects
quantiles	A list of quantiles to compute for all fixed effects
expand.factor.strategy	The strategy used to expand factors into fixed effects based on their levels. The default strategy is us use the model.matrix-function for which NA's are not allowed (expand.factor.strategy="model.matrix") and levels are possible removed. The alternative option (expand.factor.strategy="inla") use an inla-specific expansion which expand a factor into one fixed effects for each level, do allow for NA's and all levels are present in the model. In this case, factors MUST BE factors in the data.frame/list and NOT added as .+factor(x1)+. in the formula only.
mean	Prior mean for all fixed effects except the intercept. Alternatively, a named list with specific means where name=default applies to unmatched names. For example control.fixed=list(mean=list(a=1,b=2,default=0)) assign 'mean=1' to fixed effect 'a' , 'mean=2' to effect 'b' and 'mean=0' to all others. (default 0.0)
mean.intercept	Prior mean for the intercept (default 0.0)
prec	Default precision for all fixed effects except the intercept. Alternatively, a named list with specific means where name=default applies to unmatched names. For example control.fixed=list(prec=list(a=1,b=2,default=0.01)) assign 'prec=1' to fixed effect 'a' , 'prec=2' to effect 'b' and 'prec=0.01' to all others. (default 0.001)
prec.intercept	Default precision the intercept (default 0.0)
compute	Compute marginals for the fixed effects ? (default TRUE)
correlation.matrix	Compute the posterior correlation matrix for all fixed effects? (default FALSE) OOPS: This option will set up appropriate linear combinations and the results are shown as the posterior correlation matrix of the linear combinations. This option will imply control.inla=list(lincomb.derived.correlation.matrix=TRUE).

`remove.names` A vector of names of expanded fixed effects to remove from the model-matrix. This is an expert option, and should only be used if you know what you are doing.

Value

The function `control.fixed` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.fixed.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

<code>control.group</code>	<i>Control variables in control.group</i>
----------------------------	---

Description

Control variables in `control.group` for use in `inla`

Usage

```
inla.set.control.group.default(...)
control.group(adjust.for.con.comp, cyclic, fixed, graph, hyper, initial, model, order, param, prior)
```

Arguments

<code>...</code>	Possible arguments
<code>model</code>	Group model (one of 'exchangable', 'exchangablepos', 'ar1', 'ar', 'rw1', 'rw2', 'besag', or 'iid')
<code>order</code>	Defines the order of the model: for model <code>ar</code> this defines the order <code>p</code> , in $AR(p)$. Not used for other models at the time being.
<code>cyclic</code>	Make the group model cyclic? (Only applies to models 'ar1', 'rw1' and 'rw2')
<code>graph</code>	The graph spesification (Only applies to model 'besag')
<code>scale.model</code>	Scale the intrinsic model (RW1, RW2, BESAG) so the generalized variance is 1. (Default TRUE)
<code>adjust.for.con.comp</code>	Adjust for connected components when <code>scale.model=TRUE</code> ? (default TRUE)
<code>hyper</code>	Definition of the hyperparameter(s)
<code>initial</code>	(OBSOLETE!) The initial value for the group correlation or precision in the internal scale.
<code>fixed</code>	(OBSOLETE!) A boolean variable if the group correction or precision is assumed to be fixed or random.
<code>prior</code>	(OBSOLETE!) The name of the prior distribution for the group correlation or precision in the internal scale
<code>param</code>	(OBSOLETE!) Prior parameters

Value

The function `control.group` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.group.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

<code>control.hazard</code>	<i>Control variables in control.hazard</i>
-----------------------------	--

Description

Control variables in `control.hazard` for use in `inla`

Usage

```
inla.set.control.hazard.default(...)
control.hazard(constr, cutpoints, diagonal, fixed, hyper, initial, model, n.intervals, param, prior)
```

Arguments

<code>...</code>	Possible arguments
<code>model</code>	The model for the baseline hazard model. One of 'rw1', 'rw2' or 'iid'. (Default 'rw1'.)
<code>hyper</code>	The definition of the hyperparameters.
<code>fixed</code>	(OBSOLETE!) A boolean variable; is the precision for 'model' fixed? (Default FALSE.)
<code>initial</code>	(OBSOLETE!) The initial value for the precision.
<code>prior</code>	(OBSOLETE!) The prior distribution for the precision for 'model'
<code>param</code>	(OBSOLETE!) The parameters in the prior distribution
<code>constr</code>	A boolean variable; shall the 'model' be constrained to sum to zero?
<code>diagonal</code>	An extra constant added to the diagonal of the precision matrix
<code>n.intervals</code>	Number of intervals in the baseline hazard. (Default 15)
<code>cutpoints</code>	The cutpoints to use. If not specified they are computed from 'n.intervals' and the maximum length of the interval. (Default NULL)
<code>strata.name</code>	The name of the stratification variable for the baseline hazard in the data.frame
<code>scale.model</code>	Scale the baseline hazard model (RW1, RW2) so the generalized variance is 1. (Default <code>inla.getOption("scale.model.default")</code> .)

Value

The function `control.hazard` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.hazard.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.inla	<i>Control variables in control.inla</i>
--------------	--

Description

Control variables in `control.inla` for use in `inla`

Usage

```
inla.set.control.inla.default(...)
control.inla(adapt.hessian.max.trials, adapt.hessian.mode, adapt.hessian.scale, adaptive.max, adj
```

Arguments

...	Possible arguments
strategy	Character The strategy to use for the approximations; one of 'auto' (default), 'gaussian', 'simplified.laplace', 'laplace' or 'adaptive'
int.strategy	Character The integration strategy to use; one of 'auto' (default), 'ccd', 'grid', 'eb' (empirical bayes), 'user' or 'user.std'
int.design	Matrix Matrix of user-defined integration points and weights. Each row consists theta values and the integration weight. (EXPERIMENTAL!)
interpolator	Character The interpolator used to compute the marginals for the hyperparameters. One of 'auto', 'nearest', 'quadratic', 'weighted.distance', 'ccd', 'ccdintegrate', 'gridsum', 'gaussian'. Default is 'auto'.
fast	Logical If TRUE, then replace conditional modes in the Laplace approximation with conditional expectation (default TRUE)
linear.correction	Logical Default TRUE for the 'strategy = laplace' option.
h	Numerical The step-length for the gradient calculations for the hyperparameters. Default 0.005.
dz	Numerical The step-length in the standardised scale for the integration of the hyperparameters. Default 0.75.
diff.logdens	Numerical The difference of the log.density for the hyperparameters to stop numerical integration using <code>int.strategy='grid'</code> . Default 6.
print.joint.hyper	Logical If TRUE, the store also the joint distribution of the hyperparameters (without any costs). Default TRUE.
force.diagonal	Logical If TRUE, then force the Hessian to be diagonal. (Default FALSE)
skip.configurations	Logical Skip configurations if the values at the main axis are too small. (Default TRUE)
mode.known	Logical If TRUE then no optimisation is done. (Default FALSE.)

adjust.weights	Logical If TRUE then just more accurate integration weights. (Default TRUE.)
tolerance	Numerical The tolerance for the optimisation of the hyperparameters. If set, this is the default value for for 'tolerance.f/2', 'tolerance.g' and 'tolerance.x'; see below.
tolerance.f	Numerical The tolerance for the absolute change in the log posterior in the optimisation of the hyperparameters.
tolerance.g	Numerical The tolerance for the absolute change in the gradient of the log posterior in the optimisation of the hyperparameters.
tolerance.x	Numerical The tolerance for the change in the hyperparameters (root-mean-square) in the optimisation of the hyperparameters.
tolerance.step	Numerical The tolerance for the change in root-mean_squre in the inner Newton-like optimisation of the latent field.
restart	Numerical To improve the optimisation, the optimiser is restarted at the found optimum 'restart' number of times.
optimiser	Character The optimiser to use; one of 'gsl' or 'default'.
verbose	Logical Run in verbose mode? (Default FALSE)
reordering	Character Type of reordering to use. (EXPERT OPTION; one of "AUTO", "DEFAULT", "IDENTITY", "REVERSEIDENTITY", "BAND", "METIS", "GENMMD", "AMD", "MD", "MMD", "AMDBAR", "AMDC", "AMDBARC", or the output from inla.qreordering. Default is 'auto'.)
cpo.diff	Numerical Threshold to define when the cpo-calculations are inaccurate. (EXPERT OPTION.)
npoints	Numerical Number of points to use in the 'strategy=laplace' approximation (default 9)
cutoff	Numerical The cutoff used in the 'strategy=laplace' approximation. (Smaller value is more accurate and more slow.) (default 1e-4)
adapt.hessian.mode	Logical Should optimisation be continued if the Hessian estimate is void? (Default TRUE)
adapt.hessian.max.trials	Numerical Number of steps in the adaptive Hessian optimisation
adapt.hessian.scale	Numerical The scaling of the 'h' after each trial.
adaptive.max	Selecting strategy="adaptive" will chose the default strategy for all fixed effects and model components with length less or equal to adaptive.max, for others, the gaussian strategy will be applied.
huge	Logical If TRUE then try to do some of the internal parallisations differently. Hopefully this will be of benefite for 'HUGE' models. (Default FALSE.) [THIS OPTION IS OBSOLETE AND NOT USED!]
step.len	Numerical The step-length used to compute numerical derivaties of the log-likelihood
stencil	Numerical Number of points in the stencil used to compute the numerical derivaties of the log-likelihood (3, 5, 7 or 9). (default 5)
lincomb.derived.correlation.matrix	Logical If TRUE compute also the correlations for the derived linear combinations, if FALSE do not (Default FALSE)

diagonal	Numerical Expert use only! Add a this value on the diagonal of the joint precision matrix. (default 0.0)
numint.maxfeval	Numerical Maximum number of function evaluations in the the numerical integration for the hyperparameters. (Default 100000.)
numint.relerr	Numerical Relative error requirement in the the numerical integration for the hyperparameters. (Default 1e-5)
numint.abserr	Numerical Absolute error requirement in the the numerical integration for the hyperparameters. (Default 1e-6)
cmn	Numerical The minimum value for the negative Hessian from the likelihood. Increasing this value will stabilise the optimisation but can introduce bias. (Default -Inf)
b.strategy	Character If cmn is used, either keep the linear term (with b.strategy="keep") or skip the contribution by setting the linear term to zero (b.strategy="skip"). The default value is "skip"
step.factor	Numerical The step factor in the Newton-Raphson algorithm saying how large step to take (Default 1.0)
global.node.factor	Numerical The factor which defines the degree required (how many neighbors), as a fraction of n-1, that is required to be classified as a global node and numbered last (whatever the reordering routine says). Here, n, is the size of the graph. (Disabled if larger than 1.) (default 2.0)
global.node.degree	Numerical The degree required (number of neighbors) to be classified as a global node and numbered last (whatever the reordering routine says). (default .Machine\$integer.max)
stupid.search	Logical Enable or disable the stupid-search-algorithm, if the Hessian calculations reveals that the mode is not found. (Default TRUE.)
stupid.search.max.iter	Numerical Maximum number of iterations allowed for the stupid-search-algorithm. (default 1000)
stupid.search.factor	Numerical Factor (≥ 1) to increase the step-length with after each new interaction. (default 1.05)
control.vb	List of arguments for various VB corrections. enable Logical/Character Use this feature? If "auto" this will be selected automatically. strategy Character What to correct, one of "mean", "more.to.come.later...". verbose Logical Be verbose or not. refinement Integer Number of extra possible refinement iterations. max.correct Numerical Bound the allowed correction (strategy="mean": $ \text{diff.mean}/\text{stdev} < \text{max.correct}$). hyperpar.correct Logical Correct the marginal posterior for the hyperparameters? f.enable.limit The size limit to correct for a $f()$ model component.
num.gradient	Character Set the numerical scheme to compute the gradient, one of "forward" or "central" (default).
num.hessian	Character Set the numerical scheme to compute the Hessian, one of "forward" or "central" (default).
optimise.strategy	Character THIS OPTION IS EXPERIMENTAL. Chose the optimiser strategy, one of "plain" or "smart" (default)

use.directions	THIS OPTION IS EXPERIMENTAL. Unless FALSE or NULL, use directions for computing gradient and Hessian, initialised with use.directions if a matrix.
constr.marginal.diagonal	Add stability to $AQ^{-1}A^T$ by adding a small diagonal term. (default $\epsilon^{0.5}$)
improved.simplified.laplace	If TRUE use an experimental improved variant, otherwise, use the standard one.
fast	If stage1 only then try to do all calculations during stage1 by enabling Gaussian strategy with VB correction enabled.

Value

The function `control.inla` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.inla.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.lincomb	<i>Control variables in control.lincomb</i>
-----------------	---

Description

Control variables in `control.lincomb` for use in `inla`

Usage

```
inla.set.control.lincomb.default(...)
control.lincomb(verbose)
```

Arguments

...	Possible arguments
verbose	Use verbose mode for linear combinations if verbose model is set globally. (Default TRUE)

Value

The function `control.lincomb` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.lincomb.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.link	<i>Control variables in control.link</i>
--------------	--

Description

Control variables in control.link for use in inla

Usage

```
inla.set.control.link.default(...)
control.link(a, fixed, hyper, initial, model, order, param, prior, quantile, variant)
```

Arguments

...	Possible arguments
model	The name of the link function/model
order	The order of the link function, where the interpretation of order is model-dependent.
variant	The variant of the link function, where the interpretation of variant is model-dependent.
hyper	Definition of the hyperparameter(s) for the link model chosen
quantile	The quantile for quantile link function
a	The parameter a in the LOGa link
initial	(OBSOLETE!) The initial value(s) for the hyperparameter(s)
fixed	(OBSOLETE!) A boolean variable if hyperparameter(s) is/are fixed or random
prior	(OBSOLETE!) The name of the prior distribution(s) for the hyperparameter(s)
param	(OBSOLETE!) The parameters for the prior distribution(s) for the hyperparameter(s)

Value

The control.link-list is set within the corresponding control.family-list as the link is likelihood-family specific. The function control.link is used to TAB-complete arguments and returns a list of given arguments. The function inla.set.control.link.default returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.mix	<i>Control variables in control.mix</i>
-------------	---

Description

Control variables in control.mix for use in inla

Usage

```
inla.set.control.mix.default(...)
control.mix(fixed, hyper, initial, integrator, model, npoints, param, prior)
```

Arguments

...	Possible arguments
model	The model for the random effect. Currently, only model='gaussian' is implemented
hyper	Definition of the hyperparameter(s) for the random effect model chosen
initial	(OBSOLETE!) The initial value(s) for the hyperparameter(s)
fixed	(OBSOLETE!) A boolean variable if hyperparameter(s) is/are fixed or random
prior	(OBSOLETE!) The name of the prior distribution(s) for the hyperparameter(s)
param	(OBSOLETE!) The parameters for the prior distribution(s) for the hyperparameter(s)
npoints	Number of points used to do the numerical integration (default 101)
integrator	The integration scheme to use (default, quadrature, simpson)

Value

The control.mix -list is set within the corresponding control.family-list a the mixture of the likelihood is likelihood specific. (This option is EXPERIMENTAL.) The function control.mix is used to TAB-complete arguments and returns a list of given arguments. The function inla.set.control.mix.default returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.mode *Control variables in control.mode*

Description

Control variables in control.mode for use in inla

Usage

```
inla.set.control.mode.default(...)
control.mode(fixed, restart, result, theta, x)
```

Arguments

...	Possible arguments
result	Previous result from inla(). Use the theta- and x-mode from this run.
theta	The theta-mode/initial values for theta. This option has preference over result\$mode\$theta.
x	The x-mode/initial values for x. This option has preference over result\$mode\$x.
restart	A boolean variable; should we restart the optimisation from this configuration or fix the mode at this configuration? (Default FALSE.)
fixed	A boolean variable. If TRUE then treat all thetas as known and fixed, and if FALSE then treat all thetas as unknown and random (default).

Value

The function control.mode is used to TAB-complete arguments and returns a list of given arguments. The function inla.set.control.mode.default returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.pardiso.default *Control variables in control.pardiso.default*

Description

Control variables in control.pardiso.default for use in inla

Usage

```
inla.set.control.pardiso.default.default(...)
control.pardiso.default(debug, nrhs, parallel.reordering, verbose)
```

Arguments

...	Possible arguments
verbose	Show detailed output (default FALSE)
debug	Show internal debug output (default FALSE)
parallel.reordering	Do reordering in parallel (default TRUE)
nrhs	Number of right-hand sides to solve for in parallel (default depends on the argument num.threads and -1 will not change it)

Value

Extra options controlling the PARDISO library The function `control.pardiso.default` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.pardiso.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.predictor	<i>Control variables in control.predictor</i>
-------------------	---

Description

Control variables in `control.predictor` for use in `inla`

Usage

```
inla.set.control.predictor.default(...)
control.predictor(A, cdf, compute, cross, fixed, hyper, initial, link, param, precision, prior, quantiles)
```

Arguments

...	Possible arguments
hyper	Definition of the hyperparameters.
fixed	(OBSOLETE!) If the precision for the artificial noise is fixed or not (default TRUE)
prior	(OBSOLETE!) The prior for the artificial noise
param	(OBSOLETE!) Prior parameters for the artificial noise
initial	(OBSOLETE!) The value of the log precision of the artificial noise
compute	A boolean variable; should the marginals for the linear predictor be computed? (Default FALSE.)
cdf	A list of values to compute the CDF for the linear predictor
quantiles	A list of quantiles to compute for the linear predictor

cross	Cross-sum-to-zero constraints with the linear predictor. All linear predictors with the same level of 'cross' are constrained to have sum zero. Use 'NA' for no contribution. 'Cross' has the same length as the linear predictor (including the 'A' matrix extension). (THIS IS AN EXPERIMENTAL OPTION, CHANGES MAY APPEAR.)
A	The observation matrix (matrix or Matrix::sparseMatrix).
precision	The precision for $\eta^* - A\eta$, (default $\exp(15)$)
link	Define the family-connection for unobserved observations (NA). link is integer values which defines the family connection; family[link[idx]] unless is.na(link[idx]) for which the identity-link is used. The link-argument only influence the fitted.values in the result-object. If is.null(link) (default) then the identity-link is used for all missing observations. If the length of link is 1, then this value is replicated with the length of the response vector. If an element of the response vector is !NA then the corresponding entry in link is not used (but must still be a legal value). Setting this variable implies compute=TRUE.
vb.correct	Logical Add this model component to the list of variables to be used for the vb corrected Laplace approximation? If NULL use default choice, otherwise correct if TRUE and do not if FALSE (expert option)

Value

The function `control.predictor` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.predictor.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

control.results	<i>Control variables in control.results</i>
-----------------	---

Description

Control variables in `control.results` for use in `inla`

Usage

```
inla.set.control.results.default(...)
control.results(return.marginals.predictor, return.marginals.random)
```

Arguments

... Possible arguments
return.marginals.random
A boolean variable; read the marginals for the fterms? (Default TRUE)
return.marginals.predictor
A boolean variable; read the marginals for the linear predictor? (Default TRUE)

Value

The function `control.results` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.results.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.results](#), [control.mode](#), [control.hazard](#), [inla](#)

control.update	<i>Control variables in control.update</i>
----------------	--

Description

Control variables in `control.update` for use in `inla`

Usage

```
inla.set.control.update.default(...)
control.update(result)
```

Arguments

...	Possible arguments
result	Update the joint posterior for the hyperparameters from result

Value

The function `control.update` is used to TAB-complete arguments and returns a list of given arguments. The function `inla.set.control.update.default` returns a list with all the default values of all parameters within this control statement.

See Also

[control.update](#), [control.lincomb](#), [control.group](#), [control.mix](#), [control.link](#), [control.expert](#), [control.compute](#), [control.pardiso.default](#), [control.bgev.default](#), [control.family](#), [control.fixed](#), [control.inla](#), [control.predictor](#), [control.mode](#), [control.hazard](#), [inla](#)

cut

*Group-wise model criticism using node-splitting***Description**

This function performs group-wise, cross-validators model assessment for an INLA model using so-called node-splitting (Marshall and Spiegelhalter, 2007; Presanis et al, 2013). The user inputs an object of class `inla` (i.e. a result of a call to `inla()`) as well as a variable name (`split.by`) specifying a grouping: Data points that share the same value of `split.by` are in the same group. The function then checks whether each group is an "outlier", or in conflict with the remaining groups, using the methodology described in Ferkingstad et al (2017). The result is a vector containing a p-value for each group, corresponding to a test for each group i , where the null hypothesis is that group i is consistent with the other groups except i (so a small p-value is evidence that the group is an "outlier"). See Ferkingstad et al (2017) for further details.

Usage

```
inla.cut(result, split.by, mc.cores = NULL, debug=FALSE)
```

Arguments

<code>result</code>	An object of class <code>inla</code> , i.e. a result of a call to <code>inla()</code>
<code>split.by</code>	The name of the variable to group by. Data points that have the same value of <code>split.by</code> are in the same group.
<code>mc.cores</code>	The number of cores to use in <code>parallel::mclapply</code> . If <code>is.null(mc.cores)</code> , then check <code>getOption("mc.cores")</code> and <code>inla.getOption("num.threads")</code> in that order.
<code>debug</code>	Print debugging information if TRUE, default is FALSE

Value

A numeric vector of p-values, corresponding to a test for each group i where the null hypothesis is that group i is consistent with the other groups except i . A small p-value for a group indicates that the group is an "outlier" (in conflict with remaining groups).

This function is EXPERIMENTAL!!!

Author(s)

Egil Ferkingstad <egil.ferkingstad@gmail.com> and Havard Rue <hrue@r-inla.org>

References

- Ferkingstad, E., Held, L. and Rue, H. (2017). Fast and accurate Bayesian model criticism and conflict diagnostics using R-INLA. arXiv preprint arXiv:1708.03272, available at <http://arxiv.org/abs/1708.03272>. Published in *Stat*, 6:331-344 (2017).
- Marshall, E. C. and Spiegelhalter, D. J. (2007). Identifying outliers in Bayesian hierarchical models: a simulation-based approach. *Bayesian Analysis*, 2(2):409-444.
- Presanis, A. M., Ohlssen, D., Spiegelhalter, D. J., De Angelis, D., et al. (2013). Conflict diagnostics in directed acyclic graphs, with applications in Bayesian evidence synthesis. *Statistical Science*, 28(3):376-397.

Examples

```
## See http://www.r-inla.org/examples/case-studies/ferkingstad-2017 and Ferkingstad et al (2017).
```

debug.graph	<i>Debug a graph-file</i>
-------------	---------------------------

Description

Debug a graph specification on file (ascii-mode only), by checking the specification along the way.

Usage

```
inla.debug.graph(graph.file)
```

Arguments

graph.file The filename of the graph (ascii-mode)

Value

If an error is found, then an error message is shown, otherwise the graph-object returned by `inla.read.graph()` is returned.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

`inla.read.graph`

Examples

```
## Not run:  
## cat("3\n 1 1 2\n 2 1 1\n 3 4\n", file="g.dat")  
## g = inla.debug.graph("g.dat")  
## End(Not run)
```

 Drivers

Time series with seasonal effect

Description

Monthly total of car drivers killed or several injured in England from January 1969 to December 1984

Format

A data frame with 204 observations on the following 4 variables.

y Number of deaths

belt Indicator of whether the belt was compulsory to use (1) or not (0)

trend time (in months)

seasonal time (in months)

Details

NB: The last 12 lines of the data set have the first column set to NULL since these data were not observed but we want to predict them.

References

Rue, H and Held, L. (2005) *Gaussian Markov Random Fields - Theory and Applications* Chapman and Hall

Examples

```
data(Drivers)
```

Epil

Repeated measures on Poisson counts

Description

Seizure counts in a randomised trial of anti-convulsant therapy in epilepsy for 59 patients.

Format

A data frame with 236 observations on the following 7 variables.

y Number of seizures

Trt indicator for the presence of treatment

Base 8-week baseline seizure counts

Age Age of the patient

V4 indicator variable for the 4th visit.

rand a numeric vector

Ind indicator for the specific patient

Source

WinBUGS/OpenBUGS Manual Examples Vol I

Examples

```
data(Epil)
```

extract.groups	<i>Extract tagged boundary/internal segments.</i>
----------------	---

Description

Extract boundary or internal segments tagged by group id:s.

Usage

```
extract.groups(...)  
  
## S3 method for class 'inla.mesh.segment'  
extract.groups(segm, groups, groups.new = groups, ...)
```

Arguments

...	Additional arguments, passed on to other methods.
segm	An inla.mesh.segment() object.
groups	The segment groups id:s to extract.
groups.new	Optional vector of group id remapping; groups[k] in the input will be replaced by groups.new[k] in the output.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.segment\(\)](#)

f

*Define general Gaussian models in the INLA formula***Description**

Function used for defining of smooth and spatial terms within `inla` model formulae. The function does not evaluate anything - it exists purely to help set up a model. The function specifies one smooth function in the linear predictor (see [inla.list.models](#)) as

$$w f(x)$$

Usage

```
f(...,
  model = "iid",
  copy=NULL,
  same.as = NULL,
  n=NULL,
  nrep = NULL,
  replicate = NULL,
  ngroup = NULL,
  group = NULL,
  control.group = inla.set.control.group.default(),
  hyper = NULL,
  initial=NULL,
  prior=NULL,
  param = NULL,
  fixed = NULL,
  season.length=NULL,
  constr = NULL,
  extraconstr=list(A=NULL, e=NULL),
  values=NULL,
  cyclic = NULL,
  diagonal = NULL,
  graph=NULL,
  graph.file=NULL,
  cdf=NULL,
  quantiles=NULL,
  Cmatrix=NULL,
  rankdef=NULL,
  Z = NULL,
  nrow = NULL,
  ncol = NULL,
  nu = NULL,
  bvalue = NULL,
  spde.prefix = NULL,
  spde2.prefix = NULL,
  spde2.transform = c("logit", "log", "identity"),
  spde3.prefix = NULL,
  spde3.transform = c("logit", "log", "identity"),
  mean.linear = inla.set.control.fixed.default()$mean,
```

```

prec.linear = inla.set.control.fixed.default()$prec,
compute = TRUE,
of=NULL,
precision = exp(14),
range = NULL,
adjust.for.con.comp = TRUE,
order = NULL,
scale = NULL,
strata = NULL,
rgeneric = NULL,
scale.model = NULL,
args.slm = list(rho.min = NULL, rho.max = NULL,
               X = NULL, W = NULL, Q.beta = NULL),
args.ar1c = list(Z = NULL, Q.beta = NULL),
args.intslope = list(subject = NULL, strata = NULL, covariates = NULL),
vb.correct = NULL,
locations = NULL,
debug = FALSE)

```

Arguments

...	Name of the covariate and, possibly of the weights vector. NB: order counts!!!! The first specified term is the covariate and the second one is the vector of weights (which can be negative).
model	A string indicating the choosen model. The default is iid. See <code>names(inla.models())\$latent</code> for a list of possible alternatives and inla.doc for detailed docs.
copy	TODO
same.as	TODO
n	An optional argument which defines the dimension of the model if this is different from <code>length(sort(unique(covariate)))</code>
nrep	TODO
replicate	We need to write documentation here
ngroup	TODO
group	TODO
control.group	TODO
hyper	Specification of the hyperparameter, fixed or random, initial values, priors and its parameters. See <code>?inla.models</code> for the list of hyparameters for each model and its default options or use <code>inla.doc()</code> for detailed info on the family and supported prior distributions.
initial	THIS OPTION IS OBSOLETE; use hyper!!! Vector indicating the starting values for the optimization algorithm. The length of the vector depends on the number of hyperparamters in the choosen model. If <code>fixed=T</code> the value at which the parameters are fixed is determines through <code>initial</code> . See <code>inla.models()\$latent\$model name'</code> to have info about the choosen model.
prior	THIS OPTION IS OBSOLETE; use hyper!!! Prior distribution(s) for the hyperparameters of the !random model. The default value depends on the type of model, see !www.r-inla.org for a detailed description of the models. See <code>names(inla.models())\$priors</code> for possible prior choices

param	THIS OPTION IS OBSOLETE; use hyper!!! Vector indicating the parameters a and b of the prior distribution for the hyperparameters. The length of the vector depends on the chosen model. See <code>inla.models()\$latent\$model name</code> to have info about the chosen model.
fixed	THIS OPTION IS OBSOLETE; use hyper!!! Vector of boolean variables indicating wheater the hyperparameters of the model are fixed or random. The length of the vector depends on the chosen model See <code>inla.models()\$latent\$model name</code> to have info about the chosen model.
season.length	Length of the seasonal component (ONLY if <code>model="seasonal"</code>)
constr	A boolean variable indicating whater to set a sum to 0 constraint on the term. By default the sum to 0 constraint is imposed on all intrinsic models ("iid","rw1","rw1","besag", etc..).
extraconstr	This argument defines extra linear constraints. The argument is a list with two elements, a matrix A and a vector e , which defines the extra constraint $Ax = e$; for example <code>extraconstr = list(A = A, e=e)</code> . The number of columns of A must correspond to the length of this f-model. Note that this constraint comes additional to the sum-to-zero constraint defined if <code>constr = TRUE</code> .
values	An optional vector giving all values assumed by the covariate for which we want estimated the effect. It must be a numeric vector, a vector of factors or NULL.
cyclic	A boolean specifying wheather the model is cyclical. Only valid for "rw1" and "rw2" models, is <code>cyclic=T</code> then the sum to 0 constraint is removed. For the correct form of the graph file see <i>Martino and Rue (2008)</i> .
diagonal	An extra constant added to the diagonal of the precision matrix.
graph	Defines the graph-object either as a file with a graph-description, an <code>inla.graph</code> -object, or as a (sparse) symmetric matrix.
graph.file	THIS OPTION IS OBSOLETE AND REPLACED BY THE MORE GENERAL ARGUMENT <code>graph</code> . PLEASE CHANGE YOUR CODE. Name of the file containing the graph of the model; see www.r-inla.org/faq .
cdf	A vector of maximum 10 values between 0 and 1 $x(0), x(1), \dots$. The function returns, for each posterior marginal the probabilities $\text{Prob}(X < x(p))$
quantiles	A vector of maximum 10 quantiles, $p(0), p(1), \dots$ to compute for each posterior marginal. The function returns, for each posterior marginal, the values $x(0), x(1), \dots$ such that $\text{Prob}(X < x(p)) = p$
Cmatrix	The specification of the precision matrix for the generic, generic3 or z models (up to a scaling constant). <code>Cmatrix</code> is either a (dense) matrix, a matrix created using <code>Matrix::sparseMatrix()</code> , or a filename which stores the non-zero elements of <code>Cmatrix</code> , in three columns: i, j and Q_{ij} . In case of the generic3 model, it is a list of such specifications.
rankdef	A number defining the rank deficiency of the model, with sum-to-zero constraint and possible extra-constraints taken into account. See details.
Z	The matrix for the z-model
nrow	Number of rows for 2d-models
ncol	Number of columns for 2d-models
nu	Smoothing parameter for the Matern2d-model, possible values are $c(0, 1, 2, 3)$

bvalue	TODO
spde.prefix	TODO
spde2.prefix	TODO
spde2.transform	TODO
spde3.prefix	TODO
spde3.transform	TODO
mean.linear	Prior mean for the linear component, only used if model="linear"
prec.linear	Prior precision for the linear component, only used if model="linear"
compute	A boolean variable indicating whether the marginal posterior distribution for the nodes in the f() model should be computed or not. This is usefull for large models where we are only interested in some posterior marginals.
of	TODO
precision	The precision for the artificial noise added when creating a copy of a model and others.
range	A vector of size two giving the lower and upper range for the scaling parameter beta in the model COPY, CLINEAR, MEC and MEB. If low = high then the identity mapping is used.
adjust.for.con.comp	If TRUE (default), adjust some of the models (currently: besag, bym, bym2 and besag2) if the number of connected components in graph is larger than 1. If FALSE, do nothing.
order	Defines the order of the model: for model ar this defines the order p, in AR(p). Not used for other models at the time being.
scale	A scaling vector. Its meaning depends on the model.
strata	Currently not in use
rgeneric	A object of class inla.rgeneric which defines the model. (EXPERIMENTAL!)
scale.model	Logical. If TRUE then scale the RW1 and RW2 and BESAG and BYM and BESAG2 and RW2D models so the their (generlized) variance is 1. Default value is inla.getOption("scale.model.default")
args.slm	Required arguments to the model="slm"; see the documentation for further details.,
args.ar1c	Required arguments to the model="ar1c"; see the documentation for further details.,
args.intslope	A list with the subject (factor), strata (factor) and covariates (numeric) for the intslope model; see the documentation for further details.,
vb.correct	Add this model component to the list of variables to be used for the vb corrected Laplace approximation? If NULL use default choice, otherwise correct if TRUE and do not if FALSE. (expert option),
locations	A matrix with locations for the model dmatern. This also defines n.
debug	Enable local debug output

Details

There is no default value for rankdef, if it is not defined by the user then it is computed by the rank deficiency of the prior model (for the generic model, the default is zero), plus 1 for the sum-to-zero constraint if the prior model is proper, plus the number of extra constraints. **Oops:** This can be wrong, and then the user must define the rankdef explicitly.

Value

TODO

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla](#), [hyperpar.inla](#)

fgn	<i>Return the coefficients in the 3-component AR(1) mixture representing FGN(H)</i>
-----	---

Description

This function will return the coefficients in the 3-component AR(1) mixture representing FGN(H)

Usage

```
inla.fgn(H, K=4L, lag.max = NULL, approx = TRUE)
```

Arguments

H	The Hurst coefficient ($0 < H < 1$), or a vector of those
K	The number of components in representation, must be 3L or 4L
lag.max	Integer. If positive integer, return the coefficients implicitly as the ACF from 0 to lag.max
approx	Logical. If lag.max is an positive integer and approx is FALSE, then return the true ACF instead of the approximated one.

Value

`inla.fgn` returns a named matrix. If `is.null(lag.max)`, then first column is H, columns $1+1:K$ are lag one correlations (or phi's), and columns $1+K+1:K$ are the weights. If `lag.max > 0`, then return the ACFs in columns $2+(0:lag.max)$, for the H in column 1, either the approximated ones or the true ones.

This function is EXPERIMENTAL!!!

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
r = c(inla.fgn(0.7))
r_m = inla.fgn(seq(0.6, 0.8, by=0.01))
```

`geobugs2inla`*INLA utility functions*

Description

Various utility functions for INLA

Usage

```
inla.geobugs2inla(adj, num, graph.file="graph.dat")
```

Arguments

<code>adj</code>	A vector listing the ID numbers of the adjacent areas for each area. This is a sparse representation of the full adjacency matrix for the study region, and can be generated using the Adjacency Tool from the Map menu in GeoBUGS.
<code>num</code>	A vector of length N (the total number of areas) giving the number of neighbours n_i for each area.
<code>graph.file</code>	Name of the file of the new graph in the INLA format.

Value

The return value is the name of the graph-file created.

Note

These are all the same function, and the two different names are due to backward-compatibility

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla](#), [inla.surv](#), [hyperpar.inla](#)

Germany	<i>Disease Mapping</i>
---------	------------------------

Description

Cases of Oral cavity cancer in Germany from 1986-1990

Format

A data frame with 544 observations on the following 4 variables.

region Region of Germany

E Fixed quantity which accounts for number of people in the district (offset)

Y Number of cases

x covariate measuring smoking consumption

References

Rue, H and Held, L. (2005) *Gaussian Markov Random Fields - Theory and Applications* Chapman and Hall

Examples

```
data(Germany)
```

graph2matrix	<i>Construct a neighbour-matrix from a graph</i>
--------------	--

Description

Construct a neighbour-matrix from a graph and display it

Usage

```
inla.graph2matrix(graph, ...)
inla.spy(graph, ..., reordering = NULL, factor = 1.0, max.dim = NULL)
```

Arguments

graph	An inla.graph-object, a (sparse) symmetric matrix, a filename containing the graph, or a list or collection of characters and/or numbers defining the graph.
reordering	A possible reordering. Typical the one obtained from a inla-call, result\$misc\$reordering, or the result of inla.qreordering.
factor	A scaling of the inla.graph-object to reduce the size.
max.dim	Maximum dimension of the inla.graph-object plotted; if missing(factor) and max.dim is set, then factor is computed automatically to give the given max.dim.
...	Additional arguments to inla.read.graph()

Value

`inla.graph2matrix` returns a sparse symmetric matrix where the non-zero pattern is defined by the graph. The `inla.spy` function, plots a binary image of a graph. The `reordering` argument is typically the reordering used by `inla`, found in `result$misc$reordering`.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla.read.graph](#), [inla.qreordering](#)

Examples

```
n = 50
Q = matrix(0, n, n)
idx = sample(1:n, 2*n, replace=TRUE)
Q[idx, idx] = 1
diag(Q) = 1
g = inla.read.graph(Q)
QQ = inla.graph2matrix(g)
inla.spy(QQ)
print(all.equal(as.matrix(Q), as.matrix(QQ)))

g.file = inla.write.graph(g)
inla.dev.new()
inla.spy(g.file)
inla.spy(g.file, reordering = inla.qreordering(g))

g = inla.read.graph(g.file)
inla.dev.new()
inla.spy(g)

inla.dev.new()
inla.spy(3, 1, "1 2 2 1 1 3 0")
inla.dev.new()
inla.spy(3, 1, "1 2 2 1 1 3 0", reordering = 3:1)
```

 idx

Convert indexes

Description

Convert indexes given by to triplet '(idx, group, replicate)' to the (one-dimensional) index used in the grouped and replicated model

Usage

```
inla.idx(idx, n = max(idx),
         group = rep(1, length(idx)), ngroup = max(group),
         replicate = rep(1, length(idx)), nrep = max(replicate))
```

Arguments

idx	The index within the basic model. (Legal values from '1' to 'n'.)
n	The length 'n' of the basic model.
group	The index within group. (Legal values from '1' to 'ngroup'.)
ngroup	Number of groups.
replicate	The index within replication. (Legal values from '1' to 'nrep'.)
nrep	Number of replications.

Value

inla.idx returns indexes in the range '1' to 'n*ngroup*nrep' representing where the triplet '(idx,group,replicate)' is stored internally in the full grouped and replicated model.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
##TODO
```

inla

Bayesian analysis of structured additive models

Description

inla performs a full Bayesian analysis of additive models using Integrated Nested Laplace approximation

Usage

```
inla(
  formula,
  family = "gaussian",
  contrasts = NULL,
  data,
  quantiles=c(0.025, 0.5, 0.975),
  E = NULL,
  offset=NULL,
  scale = NULL,
  weights = NULL,
  Ntrials = NULL,
  strata = NULL,
  link.covariates = NULL,
  verbose = FALSE,
  lincomb = NULL,
  selection = NULL,
  control.compute = list(),
  control.predictor = list(),
  control.family = list(),
```

```

control.inla = list(),
control.fixed = list(),
control.mode = list(),
control.expert = list(),
control.hazard = list(),
control.lincomb = list(),
control.update = list(),
control.pardiso = list(),
only.hyperparam = FALSE,
inla.call = inla.getOption("inla.call"),
inla.arg = inla.getOption("inla.arg"),
num.threads = inla.getOption("num.threads"),
blas.num.threads = inla.getOption("blas.num.threads"),
keep = inla.getOption("keep"),
working.directory = inla.getOption("working.directory"),
silent = inla.getOption("silent"),
inla.mode = c("classic", "twostage", "experimental"),
debug = inla.getOption("debug"),
.parent.frame = parent.frame()
)

```

Arguments

formula	A inla formula like $y \sim 1 + z + f(\text{ind}, \text{model}="iid") + f(\text{ind2}, \text{weights}, \text{model}="ar1")$. This is much like the formula for a glm except that smooth or spatial terms can be added to the right hand side of the formula. See <code>f</code> for full details and the web site www.r-inla.org for several worked out examples. Each smooth or spatial term specified through <code>f</code> should correspond to separate column of the data frame data. The response variable, <code>y</code> can be a univariate response variable, a list or the output of the function <code>inla.surf</code> for survival analysis models.
family	A string indicating the likelihood family. The default is <code>gaussian</code> with identity link. See <code>names(inla.models())\$likelihood</code> for a list of possible alternatives and use inla.doc for detailed docs for individual families.
contrasts	Optional contrasts for the fixed effects; see <code>?lm</code> or <code>?glm</code> for details.
data	A data frame or list containing the variables in the model. The data frame MUST be provided
quantiles	A vector of quantiles, $p(0), p(1), \dots$ to compute for each posterior marginal. The function returns, for each posterior marginal, the values $x(0), x(1), \dots$ such that $\text{Prob}(X < x(p)) = p$
E	Known component in the mean for the Poisson likelihoods defined as $E_i \exp(\eta_i)$ <p>where</p> η_i <p>is the linear predictor. If not provided it is set to <code>rep(1, n.data)</code>.</p>
offset	This argument is used to specify an a-priori known and fixed component to be included in the linear predictor during fitting. This should be <code>NULL</code> or a numeric vector of length either one or equal to the number of cases. One or more

offset() terms can be included in the formula instead or as well, and if both are used, they are combined into a common offset. If the A-matrix is used in the linear predictor statement `control.predictor`, then the offset given in this argument is added to `eta*`, the linear predictor related to the observations, as $\eta^* = A \eta + \text{offset}$, whereas an offset in the formula is added to `eta`, the linear predictor related to the formula, as $\eta = \dots + \text{offset.formula}$. So in this case, the offset defined here and in the formula has a different meaning and usage.

<code>scale</code>	Fixed (optional) scale parameters of the precision for Gaussian and Student-T response models. Default value is <code>rep(1, n.data)</code> .
<code>weights</code>	Fixed (optional) weights parameters of the likelihood, so the <code>log-likelihood[i]</code> is changed into <code>weights[i]*log-likelihood[i]</code> . Default value is <code>rep(1, n.data)</code> . WARNING: The normalizing constant for the likelihood is NOT recomputed, so ALL marginals (and the marginal likelihood) must be interpreted with great care.
<code>Ntrials</code>	A vector containing the number of trials for the binomial likelihood and variants, or the number of required successes for the <code>nbinomial2</code> likelihood. Default value is <code>rep(1, n.data)</code> .
<code>strata</code>	Fixed (optional) strata indicators for <code>tstrata</code> likelihood model.
<code>link.covariates</code>	A vector or matrix with covariates for link functions
<code>verbose</code>	Boolean indicating if the <code>inla</code> -program should run in a verbose mode (default FALSE).
<code>lincomb</code>	Used to define linear combination of nodes in the latent field. The posterior distribution of such linear combination is computed by the <code>inla</code> function. See www.r-inla.org/faq for examples of how to define such linear combinations.
<code>selection</code>	This is a similar argument to the one in <code>inla.posterior.sample</code> and follow the same format. This argument allows to define a subset of the latent field for which to compute an approximated joint distribution. It will appear in <code>result\$selection</code> . See also <code>?inla.rjmgarginal</code> .
<code>control.compute</code>	See <code>?control.compute</code>
<code>control.predictor</code>	See <code>?control.predictor</code>
<code>control.family</code>	See <code>?control.family</code>
<code>control.inla</code>	See <code>?control.inla</code>
<code>control.fixed</code>	See <code>?control.fixed</code>
<code>control.mode</code>	See <code>?control.mode</code>
<code>control.expert</code>	See <code>?control.expert</code>
<code>control.hazard</code>	See <code>?control.hazard</code>
<code>control.lincomb</code>	See <code>?control.lincomb</code>
<code>control.update</code>	See <code>?control.update</code>
<code>control.pardiso</code>	See <code>?control.pardiso</code>
<code>only.hyperparam</code>	A boolean variable saying if only the hyperparameters should be computed. This option is mainly used internally. (TODO: This option should not be located here, change it!)

<code>inla.call</code>	The path to, or the name of, the <code>inla</code> -program. This program is installed together with the R-package, but, for example, a native compiled version can be used instead to improve the performance.
<code>inla.arg</code>	A string indicating ALL arguments to the ' <code>inla</code> ' program and do not include default arguments. (OOPS: This is an expert option!)
<code>num.threads</code>	Maximum number of threads the <code>inla</code> -program will use, or as ' <code>A:B</code> ' defining the number threads in the outer (A) and inner (B) layer for nested parallelism.
<code>blas.num.threads</code>	The absolute value of <code>blas.num.threads</code> is the maximum number of threads the the <code>openblas/mklblas</code> will use (if available). Value is ignored if ≤ 0 (then environment variables are used).
<code>keep</code>	A boolean variable indicating that the working files (ini file, data files and results files) should be kept. If TRUE and no <code>working.directory</code> is specified the working files are stored in a directory called " <code>inla</code> ".
<code>working.directory</code>	A string giving the name of a non-existing directory where to store the working files.
<code>silent</code>	If equal to 1L or TRUE, then the <code>inla</code> -program would be "silent". If equal to 2L, then suppress also error messages from the <code>inla</code> -program.
<code>inla.mode</code>	Run <code>inla</code> in classic-mode, twostage-mode or the new experimental-mode? Not all features are available in the experimental-mode and this mode is really work-in-progress at the moment!!!! Default is to use the mode set by <code>inla.getOption("inla.mode")</code> which is default classic-mode.
<code>debug</code>	If TRUE, then enable some debug output.
<code>.parent.frame</code>	Internal use only

Value

`inla` returns an object of class "`inla`". This is a list containing at least the following arguments:

<code>summary.fixed</code>	Matrix containing the mean and standard deviation (plus, possibly quantiles and cdf) of the the fixed effects of the model.
<code>marginals.fixed</code>	A list containing the posterior marginal densities of the fixed effects of the model.
<code>summary.random</code>	List of matrices containing the mean and standard deviation (plus, possibly quantiles and cdf) of the the smooth or spatial effects defined through <code>f()</code> .
<code>marginals.random</code>	A list containing the posterior marginal densities of the random effects defined through <code>f</code> .
<code>summary.hyperpar</code>	A matrix containing the mean and sd (plus, possibly quantiles and cdf) of the hyperparameters of the model
<code>marginals.hyperpar</code>	A list containing the posterior marginal densities of the hyperparameters of the model.
<code>summary.linear.predictor</code>	A matrix containing the mean and sd (plus, possibly quantiles and cdf) of the linear predictors η in the model

<code>marginals.linear.predictor</code>	If <code>compute=TRUE</code> in <code>control.predictor</code> , a list containing the posterior marginals of the linear predictors η in the model.
<code>summary.fitted.values</code>	A matrix containing the mean and sd (plus, possibly quantiles and cdf) of the fitted values $g^{-1}(\eta)$ obtained by transforming the linear predictors by the inverse of the link function. This quantity is only computed if <code>marginals.fitted.values</code> is computed. Note that if an observation is NA then the identity link is used. You can manually transform a marginal using <code>inla.marginal.transform()</code> or set the argument <code>link</code> in the <code>control.predictor</code> -list; see <code>?control.predictor</code>
<code>marginals.fitted.values</code>	If <code>compute=TRUE</code> in <code>control.predictor</code> , a list containing the posterior marginals of the fitted values $g^{-1}(\eta)$ obtained by transforming the linear predictors by the inverse of the link function. Note that if an observation is NA then the identity link is used. You can manually transform a marginal using <code>inla.marginal.transform()</code> or set the argument <code>link</code> in the <code>control.predictor</code> -list; see <code>?control.predictor</code>
<code>summary.lincomb</code>	If <code>lincomb != NULL</code> a list of matrices containing the mean and sd (plus, possibly quantiles and cdf) of all linear combinations defined.
<code>marginals.lincomb</code>	If <code>lincomb != NULL</code> a list of posterior marginals of all linear combinations defined.
<code>selection</code>	Provide the approximated joint distribution for the selection
<code>dic</code>	If <code>dic=TRUE</code> in <code>control.compute</code> , the deviance information criteria and effective number of parameters, otherwise NULL
<code>cpo</code>	If <code>cpo=TRUE</code> in <code>control.compute</code> , a list of three elements: <code>cpo\$cpo</code> are the values of the conditional predictive ordinate (CPO), <code>cpo\$pit</code> are the values of the probability integral transform (PIT) and <code>cpo\$failure</code> indicates whether some assumptions are violated. In short, if <code>cpo\$failure[i] > 0</code> then some assumption is violated, the higher the value (maximum 1) the more seriously.
<code>po</code>	If <code>po=TRUE</code> in <code>control.compute</code> , a list of one elements: <code>po\$po</code> are the values of the predictive ordinate (CPO) ($\pi(y_i y)$)
<code>waic</code>	If <code>waic=TRUE</code> in <code>control.compute</code> , a list of two elements: <code>waic\$waic</code> is the Watanabe-Akaike information criteria, and <code>waic\$p.eff</code> is the estimated effective number of parameters
<code>mlik</code>	If <code>mlik=TRUE</code> in <code>control.compute</code> , the log marginal likelihood of the model (using two different estimates), otherwise NULL
<code>neffp</code>	Expected effective number of parameters in the model. The standard deviation of the expected number of parameters and the number of replicas for parameter are also returned
<code>mode</code>	A list of two elements: <code>mode\$theta</code> is the computed mode of the hyperparameters and <code>mode\$x</code> is the mode of the latent field given the modal value of the hyperparameters.
<code>call</code>	The matched call.
<code>formula</code>	The formula supplied
<code>nhyper</code>	The number of hyperparameters in the model
<code>cpu.used</code>	The cpu time used by the <code>inla</code> function

Author(s)

Havard Rue <hrue@r-inla.org> and Sara Martino

References

Rue, H. and Martino, S. and Chopin, N. (2009) *Approximate Bayesian Inference for latent Gaussian models using Integrated Nested Laplace Approximations*, *JRSS-series B (with discussion)*, vol 71, no 2, pp 319-392. Rue, H and Held, L. (2005) *Gaussian Markov Random Fields - Theory and Applications* Chapman and Hall

See Also

[f](#), [inla.hyperpar](#)

Examples

```
## Not run:
##See the web page \url{www.r-inla.org} for a series of worked out examples

## End(Not run)
```

inla-class	<i>inla estimation object class</i>
------------	-------------------------------------

Description

The inla class is defined in the INLA package

See Also

[inla](#)

inla.ar.pacf2phi	<i>Convert between parameterizations for the AR(p) model</i>
------------------	--

Description

These functions convert between the AR(p) coefficients `phi`, the partial autocorrelation coefficients `pacf` and the autocorrelation function `acf`. The `phi`-parameterization is the same as used for arima-models in R; see `?arima` and the parameter-vector `a` in `Details`.

Usage

```
inla.ar.pacf2phi(pac)

inla.ar.phi2pacf(phi)

inla.ar.phi2acf(phi, lag.max = length(phi))

inla.ar.pacf2acf(pac, lag.max = length(pac))
```

Arguments

pac	The partial autocorrelation coefficients
phi	The AR(p) parameters phi
lag.max	The maximum lag to compute the ACF for

Value

- `inla.ar.pacf2phi` returns phi for given pacf.
- `inla.ar.phi2pacf` returns pac for given phi.
- `inla.ar.phi2acf` returns acf for given phi.
- `inla.ar.pacf2acf` returns acf for given pacf.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
pac <- runif(5)
phi <- inla.ar.pacf2phi(pac)
pac2 <- inla.ar.phi2pacf(phi)
print(paste("Error:", max(abs(pac2 - pac))))
print("Correlation matrix (from pac)")
print(toeplitz(inla.ar.pacf2acf(pac)))
print("Correlation matrix (from phi)")
print(toeplitz(inla.ar.phi2acf(phi)))
```

inla.as.sparse	<i>Convert a matrix or sparse matrix into the sparse formate used by INLA</i>
----------------	---

Description

Convert a matrix or sparse matrix into the sparse format used by INLA (`dgTMatrix`)

Usage

```
inla.as.sparse(...)
inla.as.dgTMatrix(A, unique = TRUE, na.rm = FALSE, zeros.rm = FALSE)
```

Arguments

...	The arguments. The matrix or sparse matrix, and the additional arguments
A	The matrix
unique	Logical. If TRUE, then ensure that the internal representation is unique and there are no duplicated entries. (Do not change this unless you know what you are doing.)
na.rm	Replace NA's in the matrix with zeros.
zeros.rm	Remove zeros in the matrix.

Value

inla.as.sparse and inla.as.dgTMatrix is the same function. The returned value is a sparse matrix in the dgTMatrix-format.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
A = matrix(1:9, 3, 3)
inla.as.sparse(A)
```

inla.as.wkt_tree.wkt *Internal WKT handling*

Description

Conversion between WKT and a tree representation

Usage

```
inla.as.wkt_tree.wkt(x, ...)  
  
inla.as.wkt.wkt_tree(x, pretty = FALSE, ...)  
  
inla.wkt_tree_get_item(x, item, duplicate = 1)  
  
inla.wkt_tree_set_item(x, item_tree, duplicate = 1)
```

Arguments

x	A WKT2 string, or a wkt_tree list structure
...	Unused
item	character vector with item labels identifying a parameter item entry.
duplicate	For items that have more than one match, duplicate indicates the index number of the desired version. Default: 1
item_tree	An item tree identifying a parameter item entry

inla.barrier

*Functions for defining the Barrier models***Description**

Functions for defining Barrier models as an inla rgeneric model

Usage

```
inla.barrier.pcmatern(mesh, barrier.triangles, prior.range,
                    prior.sigma, range.fraction=0.2)
inla.barrier.polygon(mesh, barrier.triangles, Omega=NULL)
inla.barrier.q(fem, ranges, sigma=1)
inla.barrier.fem(mesh, barrier.triangles, Omega=NULL)
```

Arguments

mesh	The mesh to build the model on, from inla.mesh.2d
barrier.triangles	The numerical ids of the triangles that make up the barrier area
prior.range	2 parameters (range0, Prange) for the prior spatial range. If Prange is NA, then range0 is used as a fixed range value (not tested).
prior.sigma	2 parameters (sig0, Psig) for the prior marginal standard deviation sigma. If Psig is NA, then sig0 is used as a fixed sigma value (not tested).
range.fraction	The length of the spatial range inside the barrier area, as a fraction of the range parameter.
Omega	Advanced option for creating a set of permeable barriers (not documented)

Details

This model is described in the ArXiv preprint arXiv:1608.03787. For examples, see <https://haakonbakka.bitbucket.io/btopic107.html>.

Value

inla.barrier.pcmatern gives the (rgeneric) model object for fitting the model in INLA, inla.barrier.polygon gives the polygon around the barrier (mainly for plotting), inla.barrier.q is an internal method producing the Q matrix from a result of inla.barrier.fem, inla.barrier.fem is an internal method producing the Finite Element matrices.

Author(s)

Haakon Bakka <bakka@r-inla.org>

See Also

inla.spde2.pcmatern

inla.barrier.pcmatern *Functions for defining the Barrier models*

Description

Functions for defining Barrier models as an inla rgeneric model

Usage

```
inla.barrier.pcmatern(
  mesh,
  barrier.triangles,
  prior.range,
  prior.sigma,
  range.fraction = 0.2
)
```

```
inla.barrier.polygon(mesh, barrier.triangles, Omega = NULL)
```

```
inla.barrier.q(fem, ranges, sigma = 1, envir = NULL)
```

```
inla.barrier.fem(mesh, barrier.triangles, Omega = NULL)
```

Arguments

mesh	The mesh to build the model on, from inla.mesh.2d
barrier.triangles	The numerical ids of the triangles that make up the barrier area
prior.range	2 parameters (range0,Prange) for the prior spatial range. If Prange is NA, then range0 is used as a fixed range value (not tested).
prior.sigma	2 parameters (sig0,Psig) for the prior marginal standard deviation sigma. If Psig is NA, then sig0 is used as a fixed sigma value (not tested).
range.fraction	The length of the spatial range inside the barrier area, as a fraction of the range parameter.
Omega	Advanced option for creating a set of permeable barriers (not documented)
fem	represents the Barrier model or the Different Terrains (DT) model, by containing all the needed matrices to solve the SPDE
ranges, sigma	the hyperparameters that determine Q
envir	the environment used for caching (with optimize=TRUE), if any

Details

This model is described in the ArXiv preprint arXiv:1608.03787. For examples, see <https://haakonbakkagit.github.io/btopic128.html>

- inla.barrier.pcmatern This function creates the model component used in inla(...)
- inla.barrier.polygon This function constructs SpatialPolygons for the different subdomains (areas)

- `inla.barrier.q`: This function computes a specific precision matrix
- `inla.barrier.fem` This function computes the Finite Element matrices that are needed to compute the precision matrix `Q` later

Value

- `inla.barrier.pcmatern` gives the (rgeneric) model object for fitting the model in INLA
- `inla.barrier.polygon` gives the polygon around the barrier (mainly for plotting)
- `inla.barrier.q` is an internal method producing the `Q` matrix from a result of `inla.barrier.fem`,
- `inla.barrier.fem` is an internal method producing the Finite Element matrices.

Author(s)

Haakon Bakka <bakka@r-inla.org>

See Also

`inla.spde2.pcmatern`

`inla.binary.install` *Install alternative binary builds*

Description

Install alternative binary builds.

Usage

```
inla.binary.install(os = c("CentOS Linux-6", "CentOS Linux-7", "CentOS Linux-8",
                          "CentOS Stream-8", "Fedora-33", "Fedora-34", "Manjaro Linux",
                          "Ubuntu-16.04", "Ubuntu-18.04", "Ubuntu-20.04"),
                   verbose = TRUE, md5.check = TRUE)
```

Arguments

<code>os</code>	If <code>os</code> is given, install binary build for this <code>os</code> . If <code>os</code> is not given, chose <code>os</code> interactively among available builds.
<code>verbose</code>	Logical. Verbose output if TRUE
<code>md5.check</code>	Logical. If TRUE, stop if <code>md5-checksum-file</code> is not present or <code>md5-checksum</code> fail. If FALSE, ignore <code>md5-checksum</code> check.

Details

Install a new binary for `os` unless `missing(os)`, for which the `os` is chosen interactively among the available builds.

Value

Return TRUE if installation was successful and FALSE if not.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
## Not run:
  inla.binary.install()
  inla.binary.install(os = "CentOS Linux-7")

## End(Not run)
```

inla.changelog	<i>inla.changelog</i>
----------------	-----------------------

Description

List the recent changes in the inla-program and its R-interface

Usage

```
inla.changelog()
```

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla](#)

inla.collect.results	<i>Collect results from a inla-call</i>
----------------------	---

Description

inla.collect.results collect results from a inla-call

Usage

```
inla.collect.results(
  results.dir,
  debug=FALSE,
  only.hyperparam=FALSE,
  file.log = NULL,
  file.log2 = NULL)
```

Arguments

results.dir	The directory where the results of the inla run are stored
debug	Logical. If TRUE some debugging information are printed
only.hyperparam	Binary variable indicating wheather only the results for the hyperparameters should be collected
file.log	Character. The filename, if any, of the logfile for the internal calculations
file.log2	Character. The filename, if any, of the logfile2 for the internal calculations

Details

This function is mainly used inside `inla` to collect results after running the `inla` function. It can also be used to collect results into R after having runned a `inla` section outside R.

Value

The function returns an object of class "inla", see the help file for `inla` for details.

`inla.compare.results` *Compare INLA and MCMC results*

Description

A small utility to compare INLA and MCMC results (OBSOLETE)

Usage

```
inla.compare.results(dir.inla = NULL, dir.mcmc = NULL)
```

Arguments

<code>dir.inla</code>	The directory with the INLA results
<code>dir.mcmc</code>	The directory with the MCMC results

Value

Return nothing. This is an interactive function.

This function is OBSOLETE

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
## See demo("Tokyo-compare")
```

inla.coxph	<i>Convert a Cox proportional hazard model into Poisson regression</i>
------------	--

Description

Tools to convert a Cox proportional hazard model into Poisson regression

Usage

```
inla.coxph(formula, data, control.hazard = list(), tag="", debug=FALSE)
inla.rbind.data.frames(...)
```

Arguments

formula	The formula for the coxph model where the response must be a <code>inla.surv</code> -object.
data	All the data used in the formula, as a list.
control.hazard	Control the model for the baseline-hazard; see <code>?control.hazard</code> .
tag	An optional tag added to the names of the new variables created (to make them unique when combined with several calls of <code>inla.coxph</code> . Note that <code>E..coxph</code> is not included, as its usually merged into one vector over different expansions.
debug	Print debug-information
...	Data.frames to be <code>rbind</code> -ed, padding with NA.

Value

`inla.coxph` returns a list of new expanded variables to be used in the `inla`-call. Note that element `data` and `data.list` needs to be merged into a `list` to be passed as the `data` argument. See the example for details.

`inla.rbind.data.frames` returns the `rbinded` `data.frames` padded with NAs. There is a better implementation in `dplyr::bind_rows`, which is used if package `dplyr` is installed.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
## How the cbind.data.frames works:
df1 = data.frame(x=1:2, y=2:3, z=3:4)
df2 = data.frame(x=3:4, yy=4:5, zz=5:6)
inla.rbind.data.frames(df1, df2)

## Standard example of how to convert a coxph into a Poisson regression
n = 1000
x = runif(n)
lambda = exp(1+x)
y = rexp(n, rate=lambda)
event = rep(1,n)
data = list(y=y, event=event, x=x)
y.surv = inla.surv(y, event)
```

```

intercept1 = rep(1, n)
p = inla.coxph(y.surv ~ -1 + intercept1 + x,
              list(y.surv = y.surv, x=x, intercept1 = intercept1))

r = inla(p$formula,
        family = p$family,
        data=c(as.list(p$data), p$data.list),
        E = p$E)
summary(r)

## How to use this in a joint model
intercept2 = rep(1, n)
y = 1 + x + rnorm(n, sd=0.1)
df = data.frame(intercept2, x, y)

## new need to cbind the data.frames, and then add the list-part of
## the data
df.joint = c(as.list(inla.rbind.data.frames(p$data, df)), p$data.list)
df.joint$Y = cbind(df.joint$y.coxph, df.joint$y)

## merge the formulas, recall to add '-1' and to use the new joint
## reponse 'Y'
formula = update(p$formula, Y ~ intercept2 -1 + .)

rr = inla(formula,
          family = c(p$family, "gaussian"),
          data = df.joint,
          E = df.joint$E.coxph)

```

inla.cpo

Improved estimates for the CPO/PIT-values

Description

Improve the estimates of the CPO/PIT-values by recomputing the model-fit by removing data-points.

Usage

```

inla.cpo(result,
         force = FALSE,
         mc.cores = NULL,
         verbose = TRUE,
         recompute.mode = TRUE)

```

Arguments

result	An object of class <code>inla</code> , ie a result of a call to <code>inla()</code>
force	If <code>TRUE</code> , then recompute all CPO/PIT values and not just those with <code>result\$cpo\$failure > 0</code> .
mc.cores	The number of cores to use in <code>parallel::mclapply</code> . If <code>is.null(mc.cores)</code> , then check <code>getOption("mc.cores")</code> and <code>inla.getOption("num.threads")</code> in that order.

verbose Run in verbose mode?
recompute.mode Should be mode (and the integration points) be recomputed when a data-point is removed or not?

Value

The object returned is the same as `result` but the new improved estimates of the CPO/PIT values replaced.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla](#)

Examples

```
n = 10
y = rnorm(n)
r = inla(y ~ 1, data = data.frame(y), control.compute = list(cpo=TRUE))

rr = inla.cpo(r, force=TRUE)
```

inla.CRS

Create a coordinate reference system object

Description

Creates either a CRS object or an `inla.CRS` object, describing a coordinate reference system

Usage

```
inla.CRS(  
  projargs = NULL,  
  doCheckCRSArgs = TRUE,  
  args = NULL,  
  oblique = NULL,  
  SRS_string = NULL,  
  ...  
)  
  
inla.wkt_predef()
```

Arguments

projargs	Either 1) a projection argument string suitable as input to <code>sp::CRS</code> , or 2) an existing CRS object, or 3) a shortcut reference string to a predefined projection; run <code>names(inla.wkt_predef())</code> for valid predefined projections.
doCheckCRSArgs	default TRUE, must be set to FALSE by package developers including CRS in an S4 class definition to avoid uncontrollable loading of the <code>rgdal</code> namespace.
args	An optional list of name/value pairs to add to and/or override the PROJ4 arguments in <code>projargs</code> . <code>name=value</code> is converted to <code>"name=value"</code> , and <code>name=NA</code> is converted to <code>"name"</code> .
oblique	Vector of length at most 4 of rotation angles (in degrees) for an oblique projection, all values defaulting to zero. The values indicate (longitude, latitude, orientation, orbit), as explained in the Details section below.
SRS_string	a WKT2 string defining the coordinate system; see <code>sp::CRS</code> . This takes precedence over <code>projargs</code> .
...	Additional parameters. Not currently in use.

Details

The first two elements of the `oblique` vector are the (longitude, latitude) coordinates for the oblique centre point. The third value (orientation) is a counterclockwise rotation angle for an observer looking at the centre point from outside the sphere. The fourth value is the quasi-longitude (orbit angle) for a rotation along the oblique observers equator.

Simple oblique: `oblique=c(0,45)`

Polar: `oblique=c(0,90)`

Quasi-transversal: `oblique=c(0,0,90)`

Satellite orbit viewpoint: `oblique=c(lon0-time*v1,0,orbitangle,orbit0+time*v2)`, where `lon0` is the longitude at which a satellite orbit crosses the equator at `time=0`, when the satellite is at an angle `orbit0` further along in its orbit. The orbital angle relative to the equatorial plane is `orbitangle`, and `v1` and `v2` are the angular velocities of the planet and the satellite, respectively. Note that "forward" from the satellite's point of view is "to the right" in the projection.

When `oblique[2]` or `oblique[3]` are non-zero, the resulting projection is only correct for perfect spheres.

Value

Either an `sp::CRS` object or an `inla.CRS` object, depending on if the coordinate reference system described by the parameters can be expressed with a pure `sp::CRS` object or not.

An S3 `inla.CRS` object is a list, usually (but not necessarily) containing at least one element:

`crs` The basic `sp::CRS` object

`inla.wkt_predef` returns a WKT2 string defining a projection

`inla.wkt_predef` returns a WKT2 string defining a projection

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[sp::CRS\(\)](#), [crs_wkt\(\)](#), [inla.sp_get_crs\(\)](#) [plot.CRS\(\)](#), [inla.identical.CRS\(\)](#)

Examples

```
if (require(rgdal)) {
  crs1 <- inla.CRS("longlat_globe")
  crs2 <- inla.CRS("lambert_globe")
  crs3 <- inla.CRS("mollweide_norm")
  crs4 <- inla.CRS("hammer_globe")
  crs5 <- inla.CRS("sphere")
  crs6 <- inla.CRS("globe")
}
## Not run:
names(inla.wkt_predef())

## End(Not run)

## Not run:
names(inla.wkt_predef())

## End(Not run)
```

inla.CRSargs

Show expanded CRS arguments

Description

Wrapper for `sp::CRS` and `inla.CRS` objects to extract the coordinate reference system argument string. Should no longer be used with PROJ6/rgdal3; see [inla.crs_get_wkt\(\)](#)

Usage

```
inla.CRSargs(x, ...)

inla.as.CRSargs.list(x, ...)

inla.as.list.CRSargs(x, ...)

inla.as.list.CRS(x, ...)

inla.as.CRS.list(x, ...)
```

Arguments

`x` An `sp::CRS` or `inla.CRS` object (for `inla.CRSargs` and `inla.as.list.CRS`), a character string (for `inla.as.list.CRSargs`), or a list (for `inla.as.CRS.list` and `inla.as.CRSargs.list`).

`...` Additional arguments passed on to other methods.

Details

- `inla.as.CRSargs.list`: CRS proj4 string for name=value pair list
- `inla.as.list.CRSargs`: List of name=value pairs from CRS proj4 string

Value

For `inla.CRSargs` and `inla.as.CRSargs.list`, a character string with PROJ.4 arguments.

For `inla.as.list.CRS` and `inla.as.list.CRSargs`, a list of name/value pairs.

For `inla.as.CRS.list`, a CRS or `inla.CRS` object.

Author(s)

Finn Lindgren finn.lindgren@gmail.com

See Also

[rgdal::CRSargs\(\)](#), [inla.CRS\(\)](#)

Examples

```
if (require(rgdal)) {  
  crs0 <- inla.CRS("longlat")  
  p4s <- inla.CRSargs(crs0)  
  lst <- inla.as.list.CRSargs(p4s)  
  crs1 <- inla.as.CRS.list(lst)  
  lst$a <- 2  
  crs2 <- inla.CRS(p4s, args = lst)  
  print(inla.CRSargs(crs0))  
  print(inla.CRSargs(crs1))  
  print(inla.CRSargs(crs2))  
}
```

`inla.dev.new`

Opens a new device

Description

Open a new device using `dev.new` unless using RStudio

Usage

```
inla.dev.new(...)
```

Arguments

... Optional arguments to `dev.new`

Value

The value of `dev.new` if not running RStudio, otherwise NULL

Author(s)

Havard Rue <hrue@r-inla.org>

inla.diameter	<i>Diameter of a point set</i>
---------------	--------------------------------

Description

Find an upper bound to the convex hull of a point set

Usage

```
inla.diameter(x, ...)

## Default S3 method:
inla.diameter(x, manifold = "", ...)

## S3 method for class 'inla.mesh.1d'
inla.diameter(x, ...)

## S3 method for class 'inla.mesh'
inla.diameter(x, ...)

## S3 method for class 'inla.mesh.segment'
inla.diameter(x, ...)

## S3 method for class 'inla.mesh.lattice'
inla.diameter(x, ...)
```

Arguments

x	A point set as an $n \times d$ matrix, or an <code>inla.mesh</code> related object.
...	Additional parameters passed on to other methods.
manifold	Character string specifying the manifold type. Default is to treat the point set with Euclidean R^d metrics. Use <code>manifold="S2"</code> for great circle distances on the unit sphere (this is set automatically for <code>inla.mesh</code> objects).

Details

- `inla.diameter.default` Calculate upper bound for the diameter of a point set, by encapsulating in a circular domain.

Value

A scalar, upper bound for the diameter of the convex hull of the point set.

Author(s)

Finn Lindgren finn.lindgren@gmail.com

Examples

```
inla.diameter(matrix(c(0, 1, 1, 0, 0, 0, 1, 1), 4, 2))
```

inla.doc

View documentation

Description

View documentation of latent, prior and likelihood models.

Usage

```
inla.doc(what, section, verbose=FALSE)
```

Arguments

what	What to view documentation about; name of latent model, name of prior, etc. (A regular expression.)
section	An optional section, like <code>names(inla.models())</code> , to look for the documentation. If missing, all sections are used.
verbose	Logical If TRUE then run in verbose mode

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

www.r-inla.org

Examples

```
## Not run: inla.doc("rw2")
## Not run: inla.doc("gaussian", section = "prior")
```

inla.extract.el

Extract elements by matching name from container objects.

Description

Extract elements by wildcard name matching from a `data.frame`, `list`, or `matrix`.

Usage

```
inla.extract.el(M, ...)  
  
## S3 method for class 'matrix'  
inla.extract.el(M, match, by.row = TRUE, ...)  
  
## S3 method for class 'data.frame'  
inla.extract.el(M, match, by.row = TRUE, ...)  
  
## S3 method for class 'list'  
inla.extract.el(M, match, ...)
```

Arguments

M	A container object.
...	Additional arguments, not used.
match	A regex defining the matching criterion.
by.row	If TRUE, extract data by row, otherwise by column.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

inla.fmesher.smorg *Compute various mesh related quantities.*

Description

Low level function for computing finite element matrices, spherical harmonics, B-splines, and point mappings with barycentric triangle coordinates.

Usage

```
inla.fmesher.smorg(  
  loc,  
  tv,  
  fem = NULL,  
  aniso = NULL,  
  gradients = FALSE,  
  sph0 = NULL,  
  sph = NULL,  
  bspline = NULL,  
  points2mesh = NULL,  
  splitlines = NULL,  
  output = NULL,  
  keep = FALSE  
)
```

Arguments

loc	3-column triangle vertex coordinate matrix.
tv	3-column triangle vertex index matrix.
fem	Maximum finite element matrix order to be computed.
aniso	A two-element list with γ and v for an anisotropic operator $\nabla \cdot H \nabla$, where $H = \gamma I + vv^\top$
gradients	When TRUE, calculate derivative operator matrices dx, dy, and dz.
sph0	Maximal order of rotationally invariant spherical harmonics.
sph	Maximal order of general spherical harmonics.
bspline	Rotationally invariant B-splines on a sphere. 3-vector with number of basis functions n, basis degree degree, and a logical; TRUE uniform knot angles, FALSE for uniform spacing in $\sin(\textit{latitude})$.
points2mesh	3-column matrix with points to be located in the mesh.
splitlines	A list with elements loc (3-column coordinate matrix) and idx (2-column index matrix) describing line segments that are to be split into sub-segments at triangle boundaries.
output	Names of objects to be included in the output, if different from defaults.
keep	When TRUE, for debugging purposes keep the fmasher I/O files on disk.

Value

A list of generated named quantities.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

inla.generate.colors *Generate text RGB color specifications.*

Description

Generates a tex RGB color specification matrix based on a color palette.

Usage

```
inla.generate.colors(
  color,
  color.axis = NULL,
  color.n = 512,
  color.palette = cm.colors,
  color.truncate = FALSE,
  alpha = NULL
)
```

Arguments

color	character, matrix or vector
color.axis	The min/max limit values for the color mapping.
color.n	The number of colors to use in the color palette.
color.palette	A color palette function.
color.truncate	If TRUE, truncate the colors at the color axis limits.
alpha	Transparency/opaqueness values.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

inla.get.inlaEnv	<i>Return the internal environment used by INLA</i>
------------------	---

Description

A function which return the internal environment used by INLA

Usage

```
inla.get.inlaEnv()
```

Value

This function returns the internal environment used by INLA to keep internal variables.

Author(s)

Havard Rue <hrue@r-inla.org>

inla.group	<i>Group or cluster covariates</i>
------------	------------------------------------

Description

inla.group group or cluster covariates so to reduce the number of unique values

Usage

```
inla.group(x, n = 25, method = c("cut", "quantile"), idx.only = FALSE)
```

Arguments

x	The vector of covariates to group.
n	Number of classes or bins to group into.
method	Group either using bins with equal length intervals (method = "cut"), or equal distance in the 'probability' scale using the quantiles (method = "quantile").
idx.only	Option to return the index only and not the method.

Value

`inla.group` return the new grouped covariates where the classes are set to the median of all the covariates belonging to that group.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[f](#)

Examples

```
## this gives groups 3 and 8
x = 1:10
x.group = inla.group(x, n = 2)

## this is the intended use, to reduce the number of unique values in
## the of first argument of f()
n = 100
x = rnorm(n)
y = x + rnorm(n)
result = inla(y ~ f(inla.group(x, n = 20), model = "iid"), data=data.frame(y=y,x=x))
```

inla.has_PROJ6

PROJ6 detection

Description

Detect whether PROJ6 is available for INLA

Usage

```
inla.has_PROJ6()
```

```
inla.not_for_PROJ6(fun)
```

```
inla.not_for_PROJ4(fun)
```

```
inla.fallback_PROJ6(fun)
```

```
inla.requires_PROJ6(fun)
```

Arguments

`fun` The name of the calling function

Details

`inla.has_PROJ6` is called to check if PROJ6&GDAL3 are available.

Value

For `inla.has_PROJ6`, logical; TRUE if PROJ6 is available, FALSE otherwise

Functions

- `inla.not_for_PROJ6`: Called to warn about using old PROJ4 features even though PROJ6 is available
- `inla.not_for_PROJ4`: Called to give an error when calling methods that are only available for PROJ6
- `inla.fallback_PROJ6`: Called to warn about falling back to using old PROJ4 methods when a PROJ6 method hasn't been implemented
- `inla.requires_PROJ6`: Called to give an error when PROJ6 is required but not available

Examples

```
inla.has_PROJ6()
```

<code>inla.hyperpar</code>	<i>Improved estimates for the hyperparameters</i>
----------------------------	---

Description

Improve the estimates of the posterior marginals for the hyperparameters of the model using the grid integration strategy.

Usage

```
inla.hyperpar(
  result,
  skip.configurations = TRUE,
  verbose = FALSE,
  dz = 0.75,
  diff.logdens = 15,
  h = NULL,
  restart = FALSE,
  quantiles = NULL,
  keep = FALSE)
```

Arguments

<code>result</code>	An object of class <code>inla</code> , ie a result of a call to <code>inla()</code>
<code>skip.configurations</code>	A boolean variable; skip configurations if the values at the main axis are too small. (Default TRUE)
<code>verbose</code>	Boolean indicating whether the <code>inla</code> program should run in a verbose mode.
<code>dz</code>	Step length in the standardized scale used in the construction of the grid, default 0.75.
<code>diff.logdens</code>	The difference of the log density for the hyperparameters to stop numerical integration using <code>int.strategy='grid'</code> . Default 15

h	The step-length for the gradient calculations for the hyperparameters. Default 0.01.
restart	A boolean defining wheather the optimizer should start again to ind the mode or if it should use the mode contained in the object
quantiles	A vector of quantiles, to compute for each posterior marginal.
keep	A boolean variable indicating the working files (ini file, data files and results files) should be kept

Value

The object returned is the same as object but the estimates of the hyperparameters are replaced by improved estimates.

Note

This function might take a long time or if the number of hyperparameters in the model is large. If it complains and says I cannot get enough memory, try to increase the value of the argument dz or decrease diff.logdens.

Author(s)

Havard Rue <hrue@r-inla.org>

References

See the references in inla

See Also

[inla](#)

inla.hyperpar.sample *Produce samples from the approximated joint posterior for the hyperparameters*

Description

Produce samples from the approximated joint posterior for the hyperparameters

Usage

```
inla.hyperpar.sample(n, result, intern=FALSE, improve.marginals = FALSE)
```

Arguments

n	Integer. Number of samples required.
result	An inla-object, f.ex the output from an inla-call.
intern	Logical. If TRUE then produce samples in the internal scale for the hyperparameter, if FALSE then produce samples in the user-scale. (For example log-precision (intern) and precision (user-scale))
improve.marginals	Logical. If TRUE, then improve the samples taking into account possible better marginal estimates for the hyperparameters in result.

Value

A matrix where each sample is a row. The contents of the column is described in the rownames.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
n = 100
r = inla(y ~ 1 + f(idx), data = data.frame(y=rnorm(n), idx = 1:n))
ns = 500
x = inla.hyperpar.sample(ns, r)

rr = inla.hyperpar(r)
xx = inla.hyperpar.sample(ns, rr, improve.marginals=TRUE)
```

inla.identical.CRS *Test CRS and inla.CRS for equality*

Description

Wrapper for identical, optionally testing only the CRS part of two objects

Usage

```
inla.identical.CRS(crs0, crs1, crsonly = FALSE)
```

Arguments

crs0	A CRS or inla.CRS object.
crs1	A CRS or inla.CRS object.
crsonly	Logical. If TRUE, only the CRS part of a inla.CRS object is compared.

Author(s)

Finn Lindgren finn.lindgren@gmail.com

See Also

[inla.CRS\(\)](#)

Examples

```
crs0 <- inla.CRS("longlat")
crs1 <- inla.CRS("longlat", oblique = c(0, 90))
print(c(
  inla.identical.CRS(crs0, crs0),
  inla.identical.CRS(crs0, crs1),
  inla.identical.CRS(crs0, crs1, crsonly = TRUE)
))
```

inla.knmodels

Spacetime interaction models

Description

It implements the models in Knorr-Held, L. (2000) with three different constraint approaches: sum-to-zero, contrast or diagonal add.

Usage

```
inla.knmodels(
  formula,
  data,
  progress=FALSE,
  control.st=list(
    t=NULL,
    s=NULL,
    st=NULL,
    graph=NULL,
    type=c(paste(1:4), paste0(2:4, 'c'), paste0(2:4, 'd')),
    diagonal=1e-5,
    ...)
)
```

Arguments

formula	The formula specifying the other model components, without the spacetime interaction term. The spacetime interaction term will be added accordingly to the specification in the <code>control.st</code> argument. See <code>inla</code>
progress	If it is to be shown the model fitting progress. Useful if more than one interaction type is being fitted.
control.st	Named list of arguments to control the spacetime interaction. It should contain: <ul style="list-style-type: none"> time to be used as the index set for the main temporal effect which will be considered for the constraints when it is the case. space to be used as the index set for the main spatial effect which will be considered for the constraints when it is the case. spacetime to be the index set for the spacetime interaction effect. graph to be the graph for the spatial neighbor structure to be used in a <code>f</code> term for the main spatial random effect term or for building the spacetime interaction model. type to specify the spacetime interaction type. 1 to 4 corresponds to the four interaction types in Knorr-Held, L. (2000) with all the needed sum-to-zero constraints. 2c, 3c and 4c are the contrast version considering the first time or space constrained to be equal to zero. 2d, 3d and 4d are the corresponding versions when considering the diagonal add approach. diagonal to be the value to be added to the diagonal when using the diagonal add approach. timeref to specify the time point to be the reference time in the contrast parametrization.

spaceref to specify the area to be the reference for the contrast parametrization.

... where additional arguments can be passed to **f** function. Specification of the hyperparameter, fixed or random, initial value, prior and its parameters for the spacetime interaction. See `?inla.models` and look for `generic0`. By default we scale it and use the PC-prior to set the prior using the `pc.prec` prior with `param = c(0.5, 0.5)`. See documentation with `?inla.doc("pc.prec")`.

... Arguments to be passed to the `inla` function.

Value

`inla.knmodels` returns an object of class "inla". or a list of objects of this class if it is asked to compute more than one interaction type at once. Note: when the model type is 2c, 3c, 4c, 2d, 3d or 4d, it also includes linear combinations summary.

Author(s)

Elias T. Krainski

See Also

`inla.knmodels.sample` to sample from

Examples

```
### define space domain as a grid
grid <- SpatialGrid(GridTopology(c(0,0), c(1, 1), c(4, 5)))
(n <- nrow(xy <- coordinates(grid)))

### build a spatial neighborhood list
jj <- lapply(1:n, function(i)
  which(sqrt((xy[i,1]-xy[,1])^2 + (xy[i,2]-xy[,2])^2)==1))

### build the spatial adjacency matrix
graph <- sparseMatrix(rep(1:n, sapply(jj, length)),
  unlist(jj), x=1, dims=c(n, n))

### some random data at 10 time point
dat <- inla.knmodels.sample(graph, m=10, tau.t=2, tau.s=2, tau.st=3)
str(dat)
sapply(dat$x, summary)

nd <- length(dat$x$eta)
dat$e <- runif(nd, 0.9, 1.1)*rgamma(n, 40, 2)
dat$y <- rpois(nd, dat$e*exp(dat$x$eta-3))
summary(dat$y)

### fit the type 4 considering three different approaches
tgraph <- sparseMatrix(i=c(2:10, 1:9), j=c(1:9, 2:10), x=1)
res <- inla.knmodels(y ~ f(time, model='bym2', graph=tgraph) +
  f(space, model='bym2', graph=graph),
  data=dat, family='poisson', E=dat$E, progress=TRUE,
  control.st=list(time=time, space=space,
    spacetime=spacetime, graph=graph, type=c(4, '4c', '4d')),
  control.compute=list(dic=TRUE, waic=TRUE, cpo=TRUE))
sapply(res, function(x)
  c(dic=x$dic$dic, waic=x$waic$waic, cpo=-sum(log(x$cpo$cpo))))
```

inla.knmodels.sample *Spacetime interaction models sampler function*

Description

It implements the sampling method for the models in Knorr-Held, L. (2000) considering the algorithm 3.1 in Rue & Held (2005) book.

Usage

```
inla.knmodels.sample(
  graph,
  m,
  type=4,
  intercept=0,
  tau.t=1,
  phi.t=0.7,
  tau.s=1,
  phi.s=0.7,
  tau.st=1,
  ev.t=NULL,
  ev.s=NULL)
```

Arguments

graph	
m	Time dimension.
type	Integer from 1 to 4 to identify one of the four interaction type.
intercept	A constant to be added to the linear predictor
tau.t	Precision parameter for the main temporal effect.
phi.t	Mixing parameter in the bym2 model assumed for the main temporal effect.
tau.s	Precision parameter for the main spatial effect.
phi.s	Mixing parameter in the bym2 model assumed for the main spatial effect.
tau.st	Precision parameter for the spacetime effect.
ev.t	Eigenvalues and eigenvectors of the temporal precision matrix structure.
ev.s	Eigenvalues and eigenvectors of the spatial precision matrix structure.

Value

A list with the following elements

time	The time index for each observation, with length equals $m*n$.
space	The spatial index for each observation, with length equals $m*n$.
spacetime	The spacetime index for each observation, with length equals $m*n$.
x	A list with the following elements
t.iid	The unstructured main temporal effect part.
t.str	The structured main temporal effect part.

t	The main temporal effect with length equals 2m.
s.iid	The unstructured main spatial effect part.
s.str	The structured main spatial effect part.
s	The main spatial effect with length equals 2n.
st	The spacetime interaction effect with length equals m*n.
eta	The linear predictor with length equals n*m.

Author(s)

Elias T. Krainski

See Also[inla.knmodels](#) for model fitting

`inla.ks.plot`*Kolmogorov-Smirnov Test Plots*

Description

Illustrate a one-sample Kolmogorov-Smirnov test by plotting the empirical distribution deviation.

Usage

```
inla.ks.plot(x, y, diff=TRUE, ...)
```

Arguments

x	a numeric vector of data values.
y	a cumulative distribution function such as 'pnorm'.
diff	logical, indicating if the normalised difference should be plotted. If FALSE, the absolute distribution functions are plotted.
...	additional arguments for ks.test , ignored in the plotting. In particular, only two-sided tests are illustrated.

Details

In addition to the (normalised) empirical distribution deviation, lines for the K-S test statistic are drawn, as well as \pm two standard deviations around the expectation under the null hypothesis.

Value

A list with class "hctest", as generated by [ks.test](#)

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also[ks.test](#)

Examples

```
## Check for N(0,1) data
data = rowSums(matrix(runif(100*12)*2-1,100,12))/2
inla.ks.plot(data, pnorm)

## Not run:
## Check the goodness-of-fit of cross-validated predictions
result = inla(..., control.predictor=list(cpo=TRUE))
inla.ks.plot(result$pit, punif)

## End(Not run)
```

inla.list.models	<i>List available model components, likelihoods, priors, etc</i>
------------------	--

Description

List available model components, likelihoods, priors, etc. To read specific documentation for the individual elements, use [inla.doc](#).

Usage

```
inla.list.models(section = names(inla.models()), ...)
```

Arguments

section	The section(s) to list, missing section will list all sections. <code>names(inla.models())</code> lists available sections.
...	Additional argument to <code>cat</code>

Details

The list is `cat`'ed with ... arguments.

This function is EXPERIMENTAL.

Value

Nothing is returned

Author(s)

Havard Rue

Examples

```
## Not run:
inla.list.models("likelihood")
inla.list.models(c("prior", "group"))
inla.list.models(file=file("everything.txt"))

#Show detailed doc for a specific prior/likelihood/latent model
inla.doc("binomial")

## End(Not run)
```

inla.load	<i>Load or source a file</i>
-----------	------------------------------

Description

Load or source a file: (internal use)

Usage

```
inla.load(filename, debug = TRUE)
```

Arguments

filename	The name of the file to be loaded, alternatively, sourced.
debug	Logical. Turn on/off debug information.

Details

Try to load the file into the global environment, if that fail, try to source the file into the global environment.

Value

None

Author(s)

Havard Rue <hrue@r-inla.org>

inla.matern.cov	<i>Numerical evaluation of Matern and related covariance functions.</i>
-----------------	---

Description

Calculates covariance and correlation functions for Matern models and related oscillating SPDE models, on R^d and on the sphere, S^2 .

Usage

```
inla.matern.cov(
  nu,
  kappa,
  x,
  d = 1,
  corr = FALSE,
  norm.corr = FALSE,
  theta,
  epsilon = 1e-08
)

inla.matern.cov.s2(nu, kappa, x, norm.corr = FALSE, theta = 0)
```

Arguments

nu	The Matern smoothness parameter.
kappa	The spatial scale parameter.
x	Distance values.
d	Space dimension; the domain is R^d .
corr	If TRUE, calculate correlations, otherwise calculate covariances. Only used for pure Matern models (i.e. with $\theta = 0$).
norm.corr	If TRUE, normalise by the estimated variance, giving approximate correlations.
theta	Oscillation strength parameter.
epsilon	Tolerance for detecting points close to distance zero.

Details

On R^d , the models are *defined* by the spectral density given by

$$S(w) = \frac{1}{(2\pi)^d (\kappa^4 + 2\kappa^2 \cos(\pi\theta) |w|^2 + |w|^4)^{(\nu+d/2)/2}}$$

On S^2 , the models are *defined* by the spectral coefficients

$$S(k) = \frac{2k + 1}{4\pi (\kappa^4 + 2\kappa^2 \cos(\pi\theta) k(k + 1) + k^2(k + 1)^2)^{(\nu+1)/2}}$$

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

inla.mdata

Create an mdata-object for INLA

Description

This defines an mdata-object for matrix valued response-families

Usage

```
inla.mdata(y, ...)
is.inla.mdata(object)
as.inla.mdata(object)
```

Arguments

y	The response vector/matrix
...	Additional vectors/matrices of same length as y
object	Any R-object
x	An mdata object

Value

An object of class `inla.mdata`. There is method for `print`.

`is.inla.mdata` returns TRUE if object inherits from class `inla.mdata`, otherwise FALSE.

`as.inla.mdata` returns an object of class `inla.mdata`

Note

It is often required to set `Y=inla.mdata(...)` and then define the formula as `Y~...`, especially when used with `inla.stack`.

Author(s)

Havard Rue

See Also

[inla](#)

inla.merge

Merge a mixture of inla-objects

Description

Merge a mixture of inla-objects

Usage

```
## S3 method for class 'inla'
merge(x, y, ..., prob = rep(1, length(loo)),
      mc.cores = NULL, verbose = FALSE)
inla.merge(loo, prob = rep(1, length(loo)), verbose = FALSE)
```

Arguments

<code>x</code>	An inla-object to be merged
<code>y</code>	An inla-object to be merged
<code>...</code>	Additional inla-objects to be merged
<code>loo</code>	List of inla-objects to be merged
<code>prob</code>	The mixture of (possibly unnormalized) probabilities
<code>mc.cores</code>	The number of cores to use in <code>parallel::mclapply</code> . If <code>is.null(mc.cores)</code> , then check <code>getOption("mc.cores")</code> and <code>inla.getOption("num.threads")</code> in that order.
<code>verbose</code>	Turn on verbose-output or not

Details

The function `merge.inla` implements method `merge` for `inla`-objects. `merge.inla` is a wrapper for the function `inla.merge`. The interface is slightly different, `merge.inla` is more tailored for interactive use, whereas `inla.merge` is better in general code.

`inla.merge` is intended for merging a mixture of `inla`-objects, each run with the same formula and settings, except for a set of hyperparameters that are fixed to different values. Using this function, we can then integrate over these hyperparameters using (unnormalized) integration weights `prob`. The main objects to be merged, are the summary statistics and marginal densities (like for hyperparameters, fixed, random, etc). Not all entries in the object can be merged, and by default these are inherited from the first object in the list, while some are just set to `NULL`. Those objects that are merged, will be listed if run with option `verbose=TRUE`.

Note that merging hyperparameter in the user-scale is prone to discretization error in general, so it is more stable to convert the marginal of the hyperparameter from the merged internal scale to the user-scale. (This is not done by this function.)

Value

A merged `inla`-object.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
set.seed(123)
n = 100
y = rnorm(n)
y[1:10] = NA
x = rnorm(n)
z1 = runif(n)
z2 = runif(n)*n
idx = 1:n
idx2 = 1:n
lc1 = inla.make.lincomb(idx = c(1, 2, 3))
names(lc1) = "lc1"
lc2 = inla.make.lincomb(idx = c(0, 1, 2, 3))
names(lc2) = "lc2"
lc3 = inla.make.lincomb(idx = c(0, 0, 1, 2, 3))
names(lc3) = "lc3"
lc = c(lc1, lc2, lc3)
rr = list()
for (logprec in c(0, 1, 2))
  rr[[length(rr)+1]] = inla(y ~ 1 + x + f(idx, z1) + f(idx2, z2),
    lincomb = lc,
    control.family = list(hyper = list(prec = list(initial = logprec))),
    control.predictor = list(compute = TRUE, link = 1),
    data = data.frame(y, x, idx, idx2, z1, z2))
r = inla.merge(rr, prob = seq_along(rr), verbose=TRUE)
summary(r)
```

inla.mesh.1d

Function space definition objects for 1D SPDE models.

Description

Create a 1D mesh specification `inla.mesh.1d` object, that defines a function space for 1D SPDE models.

Usage

```
inla.mesh.1d(
  loc,
  interval = range(loc),
  boundary = NULL,
  degree = 1,
  free.clamped = FALSE,
  ...
)
```

```
inla.mesh.1d.fem(mesh)
```

Arguments

<code>loc</code>	B-spline knot locations.
<code>interval</code>	Interval domain endpoints.
<code>boundary</code>	Boundary condition specification. Valid conditions are <code>c('neumann', 'dirichlet', 'free', 'cyclic')</code> . Two separate values can be specified, one applied to each endpoint.
<code>degree</code>	The B-spline basis degree. Supported values are 0, 1, and 2.
<code>free.clamped</code>	If TRUE, for 'free' boundaries, clamp the basis functions to the interval endpoints.
<code>...</code>	Additional option, currently unused.
<code>mesh</code>	An <code>inla.mesh.1d</code> object

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

inla.mesh.1d.A

Mapping matrix for 1D meshes

Description

Calculates barycentric coordinates and weight matrices for `inla.mesh.1d()` objects.

Usage

```
inla.mesh.1d.A(  
  mesh,  
  loc,  
  weights = NULL,  
  derivatives = NULL,  
  method = c("linear", "nearest", "quadratic")  
)
```

Arguments

mesh	An <code>inla.mesh.1d()</code> object.
loc	Coordinate values.
weights	Weights to be applied to the A matrix rows.
derivatives	If TRUE, also compute derivative weight matrices dA and d2A.
method	Interpolation method. If not specified for <code>inla.mesh.1d.A</code> (recommended), it is determined by the mesh basis function properties.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

inla.mesh.2d

High-quality triangulations

Description

Create a triangle mesh based on initial point locations, specified or automatic boundaries, and mesh quality parameters.

Usage

```
inla.mesh.2d(  
  loc = NULL,  
  loc.domain = NULL,  
  offset = NULL,  
  n = NULL,  
  boundary = NULL,  
  interior = NULL,  
  max.edge = NULL,  
  min.angle = NULL,  
  cutoff = 1e-12,  
  max.n.strict = NULL,  
  max.n = NULL,  
  plot.delay = NULL,  
  crs = NULL  
)
```

Arguments

<code>loc</code>	Matrix of point locations to be used as initial triangulation nodes. Can alternatively be a <code>SpatialPoints</code> or <code>SpatialPointsDataFrame</code> object.
<code>loc.domain</code>	Matrix of point locations used to determine the domain extent. Can alternatively be a <code>SpatialPoints</code> or <code>SpatialPointsDataFrame</code> object.
<code>offset</code>	The automatic extension distance. One or two values, for an inner and an optional outer extension. If negative, interpreted as a factor relative to the approximate data diameter (default=-0.10???)
<code>n</code>	The number of initial nodes in the automatic extensions (default=16)
<code>boundary</code>	A list of one or two <code>inla.mesh.segment()</code> objects describing domain boundaries.
<code>interior</code>	An <code>inla.mesh.segment()</code> object describing desired interior edges.
<code>max.edge</code>	The largest allowed triangle edge length. One or two values.
<code>min.angle</code>	The smallest allowed triangle angle. One or two values. (Default=21)
<code>cutoff</code>	The minimum allowed distance between points. Point at most as far apart as this are replaced by a single vertex prior to the mesh refinement step.
<code>max.n.strict</code>	The maximum number of vertices allowed, overriding <code>min.angle</code> and <code>max.edge</code> (default=-1, meaning no limit). One or two values, where the second value gives the number of additional vertices allowed for the extension.
<code>max.n</code>	The maximum number of vertices allowed, overriding <code>max.edge</code> only (default=-1, meaning no limit). One or two values, where the second value gives the number of additional vertices allowed for the extension.
<code>plot.delay</code>	On Linux (and Mac if appropriate X11 libraries are installed), specifying a non-negative numeric value activates a rudimentary plotting system in the underlying <code>fmesh</code> program, showing the triangulation algorithm at work, with waiting time factor <code>plot.delay</code> between each step. On all systems, specifying any negative value activates displaying the result after each step of the multi-step domain extension algorithm.
<code>crs</code>	An optional CRS or <code>inla.CRS</code> object

Value

An `inla.mesh` object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

`inla.mesh.create()`, `inla.delaunay()`, `inla.nonconvex.hull()`

Examples

```
loc <- matrix(runif(10 * 2), 10, 2)

if (require("splancs")) {
  boundary <- list(
    inla.nonconvex.hull(loc, 0.1, 0.15),
```

```

        inla.nonconvex.hull(loc, 0.2, 0.2)
      )
      offset <- NULL
    } else {
      boundary <- NULL
      offset <- c(0.1, 0.2)
    }
    mesh <- inla.mesh.2d(loc, boundary = boundary, offset = offset, max.edge = c(0.05, 0.1))

    plot(mesh)

```

inla.mesh.assessment *Interactive mesh building and diagnostics*

Description

Assess the finite element approximation errors in a mesh for interactive R sessions. More detailed assessment tools are in [meshbuilder\(\)](#).

Usage

```
inla.mesh.assessment(mesh, spatial.range, alpha = 2, dims = c(500, 500))
```

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

inla.mesh.2d, inla.mesh.create, meshbuilder

Examples

```

bnd <- inla.mesh.segment(cbind(
  c(0, 10, 10, 0, 0),
  c(0, 0, 10, 10, 0)
), bnd = TRUE)
mesh <- inla.mesh.2d(boundary = bnd, max.edge = 1)
out <- inla.mesh.assessment(mesh, spatial.range = 3, alpha = 2)

```

inla.mesh.assessment *Interactive mesh building and diagnostics*

Description

Assess the finite element approximation errors in a mesh for interactive R sessions. More detailed assessment tools are in [meshbuilder](#).

Usage

```
inla.mesh.assessment(mesh)
```

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

inla.mesh.2d, inla.mesh.create, meshbuilder

Examples

```
bnd <- inla.mesh.segment(cbind(c(0, 10, 10, 0, 0),
                              c(0, 0, 10, 10, 0)), bnd = TRUE)
mesh <- inla.mesh.2d(boundary = bnd, max.edge = 1)
out <- inla.mesh.assessment(mesh, spatial.range = 3, alpha = 2)
```

inla.mesh.basis

Basis functions for inla.mesh

Description

Calculate basis functions on a 1d or 2d [inla.mesh\(\)](#)

Usage

```
inla.mesh.basis(
  mesh,
  type = "b.spline",
  n = 3,
  degree = 2,
  knot.placement = "uniform.area",
  rot.inv = TRUE,
  boundary = "free",
  free.clamped = TRUE,
  ...
)
```

Arguments

mesh	An <code>inla.mesh.1d</code> or <code>inla.mesh</code> object.
type	<code>b.spline</code> (default) for B-spline basis functions, <code>sph.harm</code> for spherical harmonics (available only for meshes on the sphere)
n	For B-splines, the number of basis functions in each direction (for 1d meshes n must be a scalar, and for planar 2d meshes a 2-vector). For spherical harmonics, n is the maximal harmonic order.
degree	Degree of B-spline polynomials. See inla.mesh.1d() .
knot.placement	For B-splines on the sphere, controls the latitudinal placements of knots. <code>"uniform.area"</code> (default) gives uniform spacing in $\sin(\text{latitude})$, <code>"uniform.latitude"</code> gives uniform spacing in latitudes.
rot.inv	For spherical harmonics on a sphere, <code>rot.inv=TRUE</code> gives the rotationally invariant subset of basis functions.

boundary	Boundary specification, default is free boundaries. See inla.mesh.1d() for more information.
free.clamped	If TRUE and boundary is "free", the boundary basis functions are clamped to 0/1 at the interval boundary by repeating the boundary knots.
...	

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.1d\(\)](#) [inla.mesh.2d\(\)](#)

Examples

```
n <- 100
loc <- matrix(runif(n * 2), n, 2)
mesh <- inla.mesh.2d(loc, max.edge = 0.05)
basis <- inla.mesh.basis(mesh, n = c(4, 5))

proj <- inla.mesh.projector(mesh)
image(proj$x, proj$y, inla.mesh.project(proj, basis[, 7]))

if (require(rgl)) {
  plot(mesh, rgl = TRUE, col = basis[, 7], draw.edges = FALSE, draw.vertices = FALSE)
}
```

inla.mesh.boundary *Constraint segment extraction for inla.mesh*

Description

Constructs an list of `inla.mesh.segment` object from boundary or interior constraint information in an [inla.mesh\(\)](#) object.

Usage

```
inla.mesh.boundary(mesh, grp = NULL)
```

```
inla.mesh.interior(mesh, grp = NULL)
```

Arguments

mesh	An <code>inla.mesh</code> object.
grp	Group indices to extract. If NULL, all boundary/interior constrain groups are extracted.

Value

A list of `inla.mesh.segment` objects.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.segment\(\)](#), [inla.mesh.create\(\)](#), [inla.mesh.create.helper\(\)](#)

Examples

```
loc <- matrix(runif(100 * 2) * 1000, 100, 2)
mesh <- inla.mesh.create.helper(points.domain = loc, max.edge = c(50, 500))
boundary <- inla.mesh.boundary(mesh)
interior <- inla.mesh.interior(mesh)
```

inla.mesh.components *Compute connected mesh subsets*

Description

Compute subsets of vertices and triangles in an inla.mesh object that are connected by edges.

Usage

```
inla.mesh.components(mesh)
```

Value

A list with elements `vertex` and `triangle`, vectors of integer labels for which connected component they belong, and `info`, a `data.frame` with columns

<code>component</code>	Connected component integer label.
<code>nV</code>	The number of vertices in the component.
<code>nT</code>	The number of triangles in the component.
<code>area</code>	The surface area associated with the component. Component labels are not comparable across different meshes, but some ordering stability is guaranteed by initiating each component from the lowest numbered triangle whenever a new component is initiated.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.2d\(\)](#), [inla.mesh.create\(\)](#)

Examples

```

# Construct two simple meshes:
loc <- matrix(c(0, 1, 0, 1), 2, 2)
mesh1 <- inla.mesh.2d(loc = loc, max.edge = 0.1)
bnd <- inla.nonconvex.hull(loc, 0.3)
mesh2 <- inla.mesh.2d(boundary = bnd, max.edge = 0.1)

# Compute connectivity information:
conn1 <- inla.mesh.components(mesh1)
conn2 <- inla.mesh.components(mesh2)
# One component, simply connected mesh
conn1$info
# Two disconnected components
conn2$info

# Extract the subset mesh for the largest component:
# (Note: some information is lost, such as fixed segments,
# and boundary edge labels.)
maxi <- conn2$info$component[which.max(conn2$info$area)]
mesh3 <- inla.mesh.create(
  loc = mesh2$loc,
  tv = mesh2$graph$tv[conn2$triangle == maxi, , drop = FALSE]
)

```

inla.mesh.components *Compute connected mesh subsets*

Description

Compute subsets of vertices and triangles in an inla.mesh object that are connected by edges.

Usage

```
inla.mesh.components(mesh)
```

Value

A list with elements `vertex` and `triangle`, vectors of integer labels for which connected component they belong, and `info`, a data.frame with columns

<code>component</code>	Connected component integer label.
<code>nV</code>	The number of vertices in the component.
<code>nT</code>	The number of triangles in the component.
<code>area</code>	The surface area associated with the component. Component labels are not comparable across different meshes, but some ordering stability is guaranteed by initiating each component from the lowest numbered triangle whenever a new component is initiated.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

inla.mesh.2d, inla.mesh.create

Examples

```
# Construct two simple meshes:
loc <- matrix(c(0,1,0,1), 2, 2)
mesh1 <- inla.mesh.2d(loc = loc, max.edge=0.1)
bnd <- inla.nonconvex.hull(loc, 0.3)
mesh2 <- inla.mesh.2d(boundary = bnd, max.edge=0.1)

# Compute connectivity information:
conn1 <- inla.mesh.components(mesh1)
conn2 <- inla.mesh.components(mesh2)
# One component, simply connected mesh
conn1$info
# Two disconnected components
conn2$info

# Extract the subset mesh for the largest component:
# (Note: some information is lost, such as fixed segments,
# and boundary edge labels.)
maxi <- conn2$info$component[which.max(conn2$info$area)]
mesh3 <- inla.mesh.create(loc = mesh2$loc,
                        tv = mesh2$graph$tv[conn2$triangle == maxi,,drop=FALSE])
```

inla.mesh.create

Low level function for high-quality triangulations

Description

Create a constrained refined Delaunay triangulation (CRDT) for a set of spatial locations.

Usage

```
inla.mesh.create(
  loc = NULL,
  tv = NULL,
  boundary = NULL,
  interior = NULL,
  extend = (missing(tv) || is.null(tv)),
  refine = FALSE,
  lattice = NULL,
  globe = NULL,
  cutoff = 1e-12,
  plot.delay = NULL,
  data.dir,
  keep = (!missing(data.dir) && !is.null(data.dir)),
  timings = FALSE,
  quality.spec = NULL,
  crs = NULL
)

inla.delaunay(loc, ...)
```

Arguments

loc	Matrix of point locations. Can alternatively be a <code>SpatialPoints</code> or <code>SpatialPointsDataFrame</code> object.
tv	A triangle-vertex index matrix, specifying an existing triangulation.
boundary	A list of <code>inla.mesh.segment</code> objects, generated by <code>inla.mesh.segment()</code> , specifying boundary constraint segments.
interior	A list of <code>inla.mesh.segment</code> objects, generated by <code>inla.mesh.segment()</code> , specifying interior constraint segments.
extend	logical or list specifying whether to extend the data region, with parameters list("n") the number of edges in the extended boundary (default=8) list("offset") the extension distance. If negative, interpreted as a factor relative to the approximate data diameter (default=-0.10) Setting to FALSE is only useful in combination lattice or boundary.
refine	logical or list specifying whether to refine the triangulation, with parameters list("min.angle") the minimum allowed interior angle in any triangle. The algorithm is guaranteed to converge for min.angle at most 21 (default=21) list("max.edge") the maximum allowed edge length in any triangle. If negative, interpreted as a relative factor in an ad hoc formula depending on the data density (default=Inf) list("max.n.strict") the maximum number of vertices allowed, overriding min.angle and max.edge (default=-1, meaning no limit) list("max.n") the maximum number of vertices allowed, overriding max.edge only (default=-1, meaning no limit)
lattice	An <code>inla.mesh.lattice</code> object, generated by <code>inla.mesh.lattice()</code> , specifying points on a regular lattice.
globe	Subdivision resolution for a semi-regular spherical triangulation with equidistant points along equidistant latitude bands.
cutoff	The minimum allowed distance between points. Point at most as far apart as this are replaced by a single vertex prior to the mesh refinement step.
plot.delay	On Linux (and Mac if appropriate X11 libraries are installed), specifying a numeric value activates a rudimentary plotting system in the underlying <code>fmesher</code> program, showing the triangulation algorithm at work.
data.dir	Where to store the <code>fmesher</code> data files. Defaults to <code>tempdir()</code> if <code>keep</code> is FALSE, otherwise <code>"inla.mesh.data"</code> .
keep	TRUE if the data files should be kept in <code>data.dir</code> or deleted afterwards. Defaults to true if <code>data.dir</code> is specified, otherwise false. Warning: If <code>keep</code> is false, <code>data.dir</code> and its contents will be deleted (unless set to <code>tempdir()</code>).
timings	If TRUE, obtain timings for the mesh construction.
quality.spec	List of vectors of per vertex <code>max.edge</code> target specification for each location in <code>loc</code> , <code>boundary/interior (segm)</code> , and <code>lattice</code> . Only used if refining the mesh.
crs	An optional CRS or <code>inla.CRS</code> object
...	Optional parameters passed on to <code>inla.mesh.create</code> .

Details

`inla.mesh.create` generates triangular meshes on subsets of R^2 and S^2 . Use the higher level wrapper function `inla.mesh.2d()` for greater control over mesh resolution and coarser domain extensions.

`inla.delaunay` is a wrapper function for obtaining the convex hull of a point set and calling `inla.mesh.create` to generate the classical Delaunay triangulation.

Value

An `inla.mesh` object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.2d\(\)](#), [inla.mesh.1d\(\)](#), [inla.mesh.segment\(\)](#), [inla.mesh.lattice\(\)](#), [inla.mesh.query\(\)](#)

Examples

```
loc <- matrix(runif(10 * 2), 10, 2)

mesh <- inla.delaunay(loc)
plot(mesh)

mesh <- inla.mesh.create(loc,
  interior = inla.mesh.segment(idx = 1:2),
  extend = TRUE,
  refine = list(max.edge = 0.1)
)
plot(mesh)

loc2 <- matrix(c(0, 1, 1, 0, 0, 0, 1, 1), 4, 2)
mesh2 <- inla.mesh.create(
  loc = loc,
  boundary = inla.mesh.segment(loc2),
  interior = inla.mesh.segment(idx = 1:2),
  quality.spec = list(seg = 0.2, loc = 0.05),
  refine = list(min.angle = 26)
)
plot(mesh2)
```

`inla.mesh.deriv`*Directional derivative matrices for functions on meshes.*

Description

Calculates directional derivative matrices for functions on `inla.mesh()` objects.

Usage

```
inla.mesh.deriv(mesh, loc)
```

Arguments

mesh	An <code>inla.mesh()</code> object.
loc	Coordinates where the derivatives should be evaluated.

Value

A	The projection matrix, $u(\text{loc}_i) = \sum_j A_{ij} w_j$
dx, dy, dz	Derivative weight matrices, $du/dx(\text{loc}_i) = \sum_j dx_{ij} w_j$, etc.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

inla.mesh.fem	<i>Finite element matrices</i>
---------------	--------------------------------

Description

Constructs finite element matrices for `inla.mesh()` and `inla.mesh.1d()` objects.

Usage

```
inla.mesh.fem(mesh, order = 2)
```

Arguments

mesh	An <code>inla.mesh()</code> or <code>inla.mesh.1d()</code> object.
order	The model order.

Value

A list of sparse matrices based on basis functions ψ_i :

c0	$c0[i,j] = \langle \psi_i, 1 \rangle$
c1	$c1[i,j] = \langle \psi_i, \psi_j \rangle$
g1	$g1[i,j] = \langle \text{grad } \psi_i, \text{grad } \psi_j \rangle$
g2	$g2 = g1 * c0^{-1} * g1$
gk	$gk = g1 * (c0^{-1} * g1)^{(k-1)}$, up to and including $k=\text{order}$

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

inla.mesh.lattice *Lattice grids for inla.mesh*

Description

Construct a lattice grid for `inla.mesh()`

Usage

```
inla.mesh.lattice(
  x = seq(0, 1, length.out = 2),
  y = seq(0, 1, length.out = 2),
  z = NULL,
  dims = if (is.matrix(x)) { dim(x) } else { c(length(x), length(y)) },
  units = NULL,
  crs = NULL
)
```

Arguments

<code>x</code>	vector or grid matrix of x-values
<code>y</code>	vector of grid matrix of y-values
<code>z</code>	if x is a matrix, a grid matrix of z-values
<code>dims</code>	the size of the grid, length 2 vector
<code>units</code>	One of <code>c("default", "longlat", "longsinlat")</code> .
<code>crs</code>	An optional CRS or <code>inla.CRS</code> object

Value

An `inla.mesh.lattice` object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh\(\)](#)

Examples

```
lattice <- inla.mesh.lattice(seq(0, 1, length.out = 17), seq(0, 1, length.out = 10))

## Use the lattice "as-is", without refinement:
mesh <- inla.mesh.create(lattice = lattice, boundary = lattice$segm)
mesh <- inla.mesh.create(lattice = lattice, extend = FALSE)
plot(mesh)

## Refine the triangulation, with limits on triangle angles and edges:
mesh <- inla.mesh.create(
  lattice = lattice,
```

```

    refine = list(max.edge = 0.08),
    extend = FALSE
  )
plot(mesh)

## Add an extension around the lattice, but maintain the lattice edges:
mesh <- inla.mesh.create(
  lattice = lattice,
  refine = list(max.edge = 0.08),
  interior = lattice$segm
)
plot(mesh)

## Only add extension:
mesh <- inla.mesh.create(lattice = lattice, refine = list(max.edge = 0.08))
plot(mesh)

```

inla.mesh.map.lim *Coordinate mappings for inla.mesh projections.*

Description

Calculates coordinate mappings for inla.mesh projections.

Usage

```

inla.mesh.map.lim(
  loc = NULL,
  projection = c("default", "longlat", "longsinlat", "mollweide")
)

inla.mesh.map(
  loc,
  projection = c("default", "longlat", "longsinlat", "mollweide"),
  inverse = TRUE
)

```

Arguments

loc	Coordinates to be mapped.
projection	The projection type.
inverse	If TRUE, loc are map coordinates and coordinates in the mesh domain are calculated. If FALSE, loc are coordinates in the mesh domain and the forward map projection is calculated.

Value

For inla.mesh.map.lim, a list:

xlim	X axis limits in the map domain
ylim	Y axis limits in the map domain

No attempt is made to find minimal limits for partial spherical domains.

Functions

- `inla.mesh.map.lim`: Projection extent limit calculations

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.project\(\)](#)

`inla.mesh.project` *Methods for projecting to/from an inla.mesh*

Description

Calculate a lattice projection to/from an [inla.mesh\(\)](#)

Usage

```
inla.mesh.project(...)

## S3 method for class 'inla.mesh'
inla.mesh.project(mesh, loc = NULL, field = NULL, crs = NULL, ...)

## S3 method for class 'inla.mesh.1d'
inla.mesh.project(mesh, loc, field = NULL, ...)

## S3 method for class 'inla.mesh.projector'
inla.mesh.project(projector, field, ...)

inla.mesh.projector(...)

## S3 method for class 'inla.mesh'
inla.mesh.projector(
  mesh,
  loc = NULL,
  lattice = NULL,
  xlim = NULL,
  ylim = NULL,
  dims = c(100, 100),
  projection = NULL,
  crs = NULL,
  ...
)

## S3 method for class 'inla.mesh.1d'
inla.mesh.projector(mesh, loc = NULL, xlim = mesh$interval, dims = 100, ...)
```

Arguments

...	Additional arguments passed on to methods.
mesh	An inla.mesh() or inla.mesh.1d() object.
loc	Projection locations. Can be a matrix or a <code>SpatialPoints</code> or a <code>SpatialPointsDataFrame</code> object.
field	Basis function weights, one per mesh basis function, describing the function to be evaluated at the projection locations. <code>Function values for on the mesh</code>
crs	An optional CRS or <code>inla.CRS</code> object associated with <code>loc</code> and/or <code>lattice</code> .
projector	An <code>inla.mesh.projector</code> object.
lattice	An inla.mesh.lattice() object.
xlim	X-axis limits for a lattice. For R2 meshes, defaults to covering the domain.
ylim	Y-axis limits for a lattice. For R2 meshes, defaults to covering the domain.
dims	Lattice dimensions.
projection	One of <code>c("default", "longlat", "longsinlat", "mollweide")</code> .

Details

The call `inla.mesh.project(mesh, loc, field=..., ...)`, is a shortcut to `inla.mesh.project(inla.mesh.projector(mesh, loc), field)`.

Value

For `inla.mesh.project(mesh, ...)`, a list with projection information. For `inla.mesh.projector(mesh, ...)`, an `inla.mesh.projector` object. For `inla.mesh.project(projector, field, ...)`, a field projected from the mesh onto the locations given by the projector object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh\(\)](#), [inla.mesh.1d\(\)](#), [inla.mesh.lattice\(\)](#)

Examples

```
n <- 20
loc <- matrix(runif(n * 2), n, 2)
mesh <- inla.mesh.create(loc, refine = list(max.edge = 0.05))
proj <- inla.mesh.projector(mesh)
field <- cos(mesh$loc[, 1] * 2 * pi * 3) * sin(mesh$loc[, 2] * 2 * pi * 7)
image(proj$x, proj$y, inla.mesh.project(proj, field))

if (require(rgl)) {
  plot(mesh, rgl = TRUE, col = field, draw.edges = FALSE, draw.vertices = FALSE)
}
```

inla.mesh.query *High-quality triangulations*

Description

Query information about an inla.mesh object.

Usage

```
inla.mesh.query(mesh, ...)
```

Arguments

mesh	An inla.mesh object.
...	Query arguments. <ul style="list-style-type: none"> • tt.neighbours Compute neighbour triangles for triangles; list of vectors: list(triangles, orders) • vt.neighbours Compute neighbour triangles for vertices; list of vectors: list(vertices, orders)

Value

A list of query results.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.create\(\)](#), [inla.mesh.segment\(\)](#), [inla.mesh.lattice\(\)](#)

Examples

```
loc <- matrix(c(0.1, 0.15), 1, 2)
lattice <- inla.mesh.lattice(dims = c(10, 10))
mesh <- inla.mesh.create(loc = loc, lattice = lattice, extend = FALSE)

vt <- which(inla.mesh.query(mesh,
  vt.neighbours = list(
    mesh$idx$loc,
    4:6
  )
)$vt.neighbours)

mesh2 <- inla.mesh.create(mesh$loc,
  tv = mesh$graph$tv[vt, , drop = FALSE],
  refine = FALSE, extend = FALSE
)
```

<code>inla.mesh.segment</code>	<i>Constraint segments for inla.mesh</i> Constructs <code>inla.mesh.segment</code> objects that can be used to specify boundary and interior constraint edges in calls to <code>inla.mesh()</code> .
--------------------------------	--

Description

Constraint segments for `inla.mesh`

Constructs `inla.mesh.segment` objects that can be used to specify boundary and interior constraint edges in calls to `inla.mesh()`.

Usage

```
inla.mesh.segment(...)
```

```
## Default S3 method:
```

```
inla.mesh.segment(
  loc = NULL,
  idx = NULL,
  grp = NULL,
  is.bnd = TRUE,
  crs = NULL,
  ...
)
```

```
## S3 method for class 'inla.mesh.segment'
```

```
inla.mesh.segment(..., grp.default = 0)
```

```
inla.contour.segment(
  x = seq(0, 1, length.out = nrow(z)),
  y = seq(0, 1, length.out = ncol(z)),
  z,
  nlevels = 10,
  levels = pretty(range(z, na.rm = TRUE), nlevels),
  groups = seq_len(length(levels)),
  positive = TRUE,
  eps = NULL,
  crs = NULL
)
```

Arguments

<code>...</code>	Additional parameters. When joining segments, a list of <code>inla.mesh.segment</code> objects.
<code>loc</code>	Matrix of point locations.
<code>idx</code>	Segment index sequence vector or index pair matrix. The indices refer to the rows of <code>loc</code> . If <code>loc==NULL</code> , the indices will be interpreted as indices into the point specification supplied to <code>inla.mesh.create()</code> . If <code>is.bnd==TRUE</code> , defaults to linking all the points in <code>loc</code> , as <code>c(1:nrow(loc), 1L)</code> , otherwise <code>1:nrow(loc)</code> .

grp	Vector of group labels for each segment. Set to NULL to let the labels be chosen automatically in a call to <code>inla.mesh.create()</code> .
is.bnd	TRUE if the segments are boundary segments, otherwise FALSE.
crs	An optional CRS or <code>inla.CRS</code> object
grp.default	When joining segments, use this group label for segments that have <code>grp=NULL</code> .
x, y, z, nlevels, levels	Parameters specifying a set of surface contours, with syntax described in <code>contour()</code> .
groups	Vector of group ID:s, one for each contour level.
positive	TRUE if the contours should encircle positive level excursions in a counter clockwise direction.
eps	Tolerance for <code>inla.simplify.curve()</code> .

Value

An `inla.mesh.segment` object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

`inla.mesh.create()`, `inla.mesh.2d()`

Examples

```
## Create a square boundary and a diagonal interior segment
loc.bnd <- matrix(c(0, 0, 1, 0, 1, 1, 0, 1), 4, 2, byrow = TRUE)
loc.int <- matrix(c(0.9, 0.1, 0.1, 0.6), 2, 2, byrow = TRUE)
segm.bnd <- inla.mesh.segment(loc.bnd)
segm.int <- inla.mesh.segment(loc.int, is.bnd = FALSE)

## Points to be meshed
loc <- matrix(runif(10 * 2), 10, 2) * 0.9 + 0.05
mesh <- inla.mesh.create(loc,
  boundary = segm.bnd,
  interior = segm.int,
  refine = list()
)
plot(mesh)
## Not run:
mesh <- inla.mesh.create(loc, interior = list(segm.bnd, segm.int))
plot(mesh)

## End(Not run)
```

inla.models

Valid models in INLA

Description

This page describe the models implemented in inla, divided into sections: latent, group, mix, link, predictor, hazard, likelihood, prior, wrapper .

Usage

```
inla.models()
```

Value

Valid sections are: latent, group, mix, link, predictor, hazard, likelihood, prior, wrapper

Section ‘latent’. Valid models in this section are:

Model ‘linear’. Number of hyperparameters are 0.

Model ‘iid’. Number of hyperparameters are 1.

Hyperparameter ‘theta’ hyperid = ‘1001’

name = ‘log precision’

short.name = ‘prec’

prior = ‘loggamma’

param = ‘1.5e-05’

initial = ‘4’

fixed = ‘FALSE’

to.theta = ‘function(x) log(x)’

from.theta = ‘function(x) exp(x)’

Properties: doc = ‘Gaussian random effects in dim=1’

constr = ‘FALSE’

nrow.ncol = ‘FALSE’

augmented = ‘FALSE’

aug.factor = ‘1’

aug.constr = ‘NULL’

n.div.by = ‘NULL’

n.required = ‘FALSE’

set.default.values = ‘FALSE’

pdf = ‘indep’

Model ‘mec’. Number of hyperparameters are 4.

Hyperparameter ‘theta1’ hyperid = ‘2001’

name = ‘beta’

short.name = ‘b’

prior = ‘gaussian’

param = ‘1 0.001’

initial = ‘1’

fixed = ‘FALSE’

```

to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '2002'
  name = 'prec.u'
  short.name = 'prec'
  prior = 'loggamma'
  param = '1 1e-04'
  initial = '9.21034037197618'
  fixed = 'TRUE'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Hyperparameter 'theta3' hyperid = '2003'
  name = 'mean.x'
  short.name = 'mu.x'
  prior = 'gaussian'
  param = '0 1e-04'
  initial = '0'
  fixed = 'TRUE'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta4' hyperid = '2004'
  name = 'prec.x'
  short.name = 'prec.x'
  prior = 'loggamma'
  param = '1 10000'
  initial = '-9.21034037197618'
  fixed = 'TRUE'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'Classical measurement error model'
  constr = 'FALSE'
  nrow.ncol = 'FALSE'
  augmented = 'FALSE'
  aug.factor = '1'
  aug.constr = 'NULL'
  n.div.by = 'NULL'
  n.required = 'FALSE'
  set.default.values = 'FALSE'
  pdf = 'mec'
Model 'mcb'. Number of hyperparameters are 2.
Hyperparameter 'theta1' hyperid = '3001'
  name = 'beta'
  short.name = 'b'
  prior = 'gaussian'
  param = '1 0.001'

```

```

initial = '1'
fixed = 'FALSE'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '3002'
name = 'prec.u'
short.name = 'prec'
prior = 'loggamma'
param = '1 1e-04'
initial = '6.90775527898214'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'Berkson measurement error model'
constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
pdf = 'meb'

```

Model 'rgeneric'. Number of hyperparameters are 0.

Model 'rw1'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '4001'
name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'Random walk of order 1'
constr = 'TRUE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
min.diff = '1e-05'
pdf = 'rw1'

```

Model 'rw2'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '5001'

name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

Properties: doc = 'Random walk of order 2'

constr = 'TRUE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
min.diff = '0.001'
pdf = 'rw2'

Model 'crw2'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '6001'

name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

Properties: doc = 'Exact solution to the random walk of order 2'

constr = 'TRUE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '2'
aug.constr = '1'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
min.diff = '0.001'
pdf = 'crw2'

Model 'seasonal'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '7001'

name = 'log precision'

```

short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'Seasonal model for time series'
constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
pdf = 'seasonal'

```

Model 'besag'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '8001'
name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'The Besag area model (CAR-model)'
constr = 'TRUE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'besag'

```

Model 'besag2'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '9001'
name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'
fixed = 'FALSE'

```

```

to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '9002'
  name = 'scaling parameter'
  short.name = 'a'
  prior = 'loggamma'
  param = '10 10'
  initial = '0'
  fixed = 'FALSE'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'The shared Besag model'
  constr = 'TRUE'
  nrow.ncol = 'FALSE'
  augmented = 'FALSE'
  aug.factor = '1'
  aug.constr = '1 2'
  n.div.by = '2'
  n.required = 'TRUE'
  set.default.values = 'TRUE'
  pdf = 'besag2'

```

Model 'bym'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '10001'
  name = 'log unstructured precision'
  short.name = 'prec.unstruct'
  prior = 'loggamma'
  param = '1 5e-04'
  initial = '4'
  fixed = 'FALSE'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '10002'
  name = 'log spatial precision'
  short.name = 'prec.spatial'
  prior = 'loggamma'
  param = '1 5e-04'
  initial = '4'
  fixed = 'FALSE'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'The BYM-model (Besag-York-Mollier model)'
  constr = 'TRUE'
  nrow.ncol = 'FALSE'
  augmented = 'TRUE'
  aug.factor = '2'

```

```

aug.constr = '2'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'bym'

```

Model 'bym2'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '11001'

```

name = 'log precision'
short.name = 'prec'
prior = 'pc.prec'
param = '1 0.01'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '11002'

```

name = 'logit phi'
short.name = 'phi'
prior = 'pc'
param = '0.5 0.5'
initial = '-3'
fixed = 'FALSE'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

Properties: doc = 'The BYM-model with the PC priors'

```

constr = 'TRUE'
nrow.ncol = 'FALSE'
augmented = 'TRUE'
aug.factor = '2'
aug.constr = '2'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
status = 'experimental'
pdf = 'bym2'

```

Model 'besagproper'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '12001'

```

name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-04'
initial = '2'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter ‘theta2’ hyperid = ‘12002’
 name = ‘log diagonal’
 short.name = ‘diag’
 prior = ‘loggamma’
 param = ‘1 1’
 initial = ‘1’
 fixed = ‘FALSE’
 to.theta = ‘function(x) log(x)’
 from.theta = ‘function(x) exp(x)’
Properties: doc = ‘A proper version of the Besag model’
 constr = ‘FALSE’
 nrow.ncol = ‘FALSE’
 augmented = ‘FALSE’
 aug.factor = ‘1’
 aug.constr = ‘NULL’
 n.div.by = ‘NULL’
 n.required = ‘TRUE’
 set.default.values = ‘TRUE’
 status = ‘experimental’
 pdf = ‘besagproper’

Model ‘besagproper2’. Number of hyperparameters are 2.

Hyperparameter ‘theta1’ hyperid = ‘13001’
 name = ‘log precision’
 short.name = ‘prec’
 prior = ‘loggamma’
 param = ‘1 5e-04’
 initial = ‘2’
 fixed = ‘FALSE’
 to.theta = ‘function(x) log(x)’
 from.theta = ‘function(x) exp(x)’
Hyperparameter ‘theta2’ hyperid = ‘13002’
 name = ‘logit lambda’
 short.name = ‘lambda’
 prior = ‘gaussian’
 param = ‘0 0.45’
 initial = ‘3’
 fixed = ‘FALSE’
 to.theta = ‘function(x) log(x / (1 -x))’
 from.theta = ‘function(x) exp(x) / (1 + exp(x))’
Properties: doc = ‘An alternative proper version of the Besag model’
 constr = ‘FALSE’
 nrow.ncol = ‘FALSE’
 augmented = ‘FALSE’
 aug.factor = ‘1’
 aug.constr = ‘NULL’

```

n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
status = 'experimental'
pdf = 'besagproper2'

```

Model 'fgn'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '13101'

```

name = 'log precision'
short.name = 'prec'
prior = 'pc.prec'
param = '3 0.01'
initial = '1'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '13102'

```

name = 'logit H'
short.name = 'H'
prior = 'pcfgnh'
param = '0.9 0.1'
initial = '2'
fixed = 'FALSE'
to.theta = 'function(x) log((2 * x - 1) / (2 * (1 - x)))'
from.theta = 'function(x) 0.5 + 0.5 * exp(x) / (1 + exp(x))'

```

Properties: doc = 'Fractional Gaussian noise model'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'TRUE'
aug.factor = '5'
aug.constr = '1'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'TRUE'
order.default = '4'
order.defined = '3 4'
pdf = 'fgn'

```

Model 'fgn2'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '13111'

```

name = 'log precision'
short.name = 'prec'
prior = 'pc.prec'
param = '3 0.01'
initial = '1'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'

```

```

from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '13112'
  name = 'logit H'
  short.name = 'H'
  prior = 'pcfgnh'
  param = '0.9 0.1'
  initial = '2'
  fixed = 'FALSE'
  to.theta = 'function(x) log((2 * x - 1) / (2 * (1 - x)))'
  from.theta = 'function(x) 0.5 + 0.5 * exp(x) / (1 + exp(x))'
Properties: doc = 'Fractional Gaussian noise model (alt 2)'
  constr = 'FALSE'
  nrow.ncol = 'FALSE'
  augmented = 'TRUE'
  aug.factor = '4'
  aug.constr = '1'
  n.div.by = 'NULL'
  n.required = 'FALSE'
  set.default.values = 'TRUE'
  order.default = '4'
  order.defined = '3 4'
  pdf = 'fgn'

```

Model 'ar1'. Number of hyperparameters are 3.

```

Hyperparameter 'theta1' hyperid = '14001'
  name = 'log precision'
  short.name = 'prec'
  prior = 'loggamma'
  param = '1 5e-05'
  initial = '4'
  fixed = 'FALSE'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '14002'
  name = 'logit lag one correlation'
  short.name = 'rho'
  prior = 'normal'
  param = '0 0.15'
  initial = '2'
  fixed = 'FALSE'
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta3' hyperid = '14003'
  name = 'mean'
  short.name = 'mean'
  prior = 'normal'

```

```

param = '0 1'
initial = '0'
fixed = 'TRUE'
to.theta = 'function(x) x'
from.theta = 'function(x) x'

```

Properties: doc = 'Auto-regressive model of order 1 (AR(1))'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
pdf = 'ar1'

```

Model 'ar1c'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '14101'

```

name = 'log precision'
short.name = 'prec'
prior = 'pc.prec'
param = '1 0.01'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '14102'

```

name = 'logit lag one correlation'
short.name = 'rho'
prior = 'pc.cor0'
param = '0.5 0.5'
initial = '2'
fixed = 'FALSE'
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'

```

Properties: doc = 'Auto-regressive model of order 1 w/covariates'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'TRUE'
status = 'experimental'
pdf = 'ar1c'

```

Model 'ar'. Number of hyperparameters are 11.

Hyperparameter 'theta1' hyperid = '15001'

name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'pc.prec'
param = '3 0.01'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

Hyperparameter 'theta2' hyperid = '15002'

name = 'pacf1'
short.name = 'pacf1'
initial = '1'
fixed = 'FALSE'
prior = 'pc.cor0'
param = '0.5 0.5'
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'

Hyperparameter 'theta3' hyperid = '15003'

name = 'pacf2'
short.name = 'pacf2'
initial = '0'
fixed = 'FALSE'
prior = 'pc.cor0'
param = '0.5 0.4'
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'

Hyperparameter 'theta4' hyperid = '15004'

name = 'pacf3'
short.name = 'pacf3'
initial = '0'
fixed = 'FALSE'
prior = 'pc.cor0'
param = '0.5 0.3'
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'

Hyperparameter 'theta5' hyperid = '15005'

name = 'pacf4'
short.name = 'pacf4'
initial = '0'
fixed = 'FALSE'
prior = 'pc.cor0'
param = '0.5 0.2'
to.theta = 'function(x) log((1 + x) / (1 - x))'

```

from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'
Hyperparameter 'theta6' hyperid = '15006'
  name = 'pacf5'
  short.name = 'pacf5'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 -x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'
Hyperparameter 'theta7' hyperid = '15007'
  name = 'pacf6'
  short.name = 'pacf6'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 -x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'
Hyperparameter 'theta8' hyperid = '15008'
  name = 'pacf7'
  short.name = 'pacf7'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 -x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'
Hyperparameter 'theta9' hyperid = '15009'
  name = 'pacf8'
  short.name = 'pacf8'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 -x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'
Hyperparameter 'theta10' hyperid = '15010'
  name = 'pacf9'
  short.name = 'pacf9'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 -x))'

```

```

from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'
Hyperparameter 'theta11' hyperid = '15011'
  name = 'pacf10'
  short.name = 'pacf10'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'
Properties: doc = 'Auto-regressive model of order p (AR(p))'
  constr = 'FALSE'
  nrow.ncol = 'FALSE'
  augmented = 'FALSE'
  aug.factor = '1'
  aug.constr = 'NULL'
  n.div.by = 'NULL'
  n.required = 'FALSE'
  set.default.values = 'FALSE'
  pdf = 'ar'

```

Model 'ou'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '16001'
  name = 'log precision'
  short.name = 'prec'
  prior = 'loggamma'
  param = '1 5e-05'
  initial = '4'
  fixed = 'FALSE'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '16002'
  name = 'log phi'
  short.name = 'phi'
  prior = 'normal'
  param = '0 0.2'
  initial = '-1'
  fixed = 'FALSE'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'The Ornstein-Uhlenbeck process'
  constr = 'FALSE'
  nrow.ncol = 'FALSE'
  augmented = 'FALSE'
  aug.factor = '1'
  aug.constr = 'NULL'

```

```

n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
pdf = 'ou'

```

Model 'intslope'. Number of hyperparameters are 13.

Hyperparameter 'theta1' hyperid = '16101'

```

name = 'log precision1'
short.name = 'prec1'
initial = '4'
fixed = 'FALSE'
prior = 'wishart2d'
param = '4 1 1 0'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '16102'

```

name = 'log precision2'
short.name = 'prec2'
initial = '4'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta3' hyperid = '16103'

```

name = 'logit correlation'
short.name = 'cor'
initial = '4'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'

```

Hyperparameter 'theta4' hyperid = '16104'

```

name = 'gamma1'
short.name = 'g1'
initial = '1'
fixed = 'TRUE'
prior = 'normal'
param = '1 36'
to.theta = 'function(x) x'
from.theta = 'function(x) x'

```

Hyperparameter 'theta5' hyperid = '16105'

```

name = 'gamma2'
short.name = 'g2'
initial = '1'

```

```
fixed = 'TRUE'
prior = 'normal'
param = '1 36'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta6' hyperid = '16106'
  name = 'gamma3'
  short.name = 'g3'
  initial = '1'
  fixed = 'TRUE'
  prior = 'normal'
  param = '1 36'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta7' hyperid = '16107'
  name = 'gamma4'
  short.name = 'g4'
  initial = '1'
  fixed = 'TRUE'
  prior = 'normal'
  param = '1 36'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta8' hyperid = '16108'
  name = 'gamma5'
  short.name = 'g5'
  initial = '1'
  fixed = 'TRUE'
  prior = 'normal'
  param = '1 36'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta9' hyperid = '16109'
  name = 'gamma6'
  short.name = 'g6'
  initial = '1'
  fixed = 'TRUE'
  prior = 'normal'
  param = '1 36'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta10' hyperid = '16110'
  name = 'gamma7'
  short.name = 'g7'
  initial = '1'
```

```

    fixed = 'TRUE'
    prior = 'normal'
    param = '1 36'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta11' hyperid = '16111'
    name = 'gamma8'
    short.name = 'g8'
    initial = '1'
    fixed = 'TRUE'
    prior = 'normal'
    param = '1 36'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta12' hyperid = '16112'
    name = 'gamma9'
    short.name = 'g9'
    initial = '1'
    fixed = 'TRUE'
    prior = 'normal'
    param = '1 36'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta13' hyperid = '16113'
    name = 'gamma10'
    short.name = 'g10'
    initial = '1'
    fixed = 'TRUE'
    prior = 'normal'
    param = '1 36'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Properties: doc = 'Intecept-slope model with Wishart-prior'
    constr = 'FALSE'
    nrow.ncol = 'FALSE'
    augmented = 'FALSE'
    aug.factor = '1'
    aug.constr = 'NULL'
    n.div.by = 'NULL'
    n.required = 'FALSE'
    set.default.values = 'TRUE'
    status = 'experimental'
    pdf = 'intslope'
Model 'generic'. Number of hyperparameters are 1.
Hyperparameter 'theta' hyperid = '17001'

```

```

name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'A generic model'
constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'generic0'

```

Model 'generic0'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '18001'
name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'A generic model (type 0)'
constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'generic0'

```

Model 'generic1'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '19001'
name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '4'

```

```

    fixed = 'FALSE'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '19002'
    name = 'beta'
    short.name = 'beta'
    initial = '2'
    fixed = 'FALSE'
    prior = 'gaussian'
    param = '0 0.1'
    to.theta = 'function(x) log(x / (1 -x))'
    from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'A generic model (type 1)'
    constr = 'FALSE'
    nrow.ncol = 'FALSE'
    augmented = 'FALSE'
    aug.factor = '1'
    aug.constr = 'NULL'
    n.div.by = 'NULL'
    n.required = 'TRUE'
    set.default.values = 'TRUE'
    pdf = 'generic1'

```

Model 'generic2'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '20001'
    name = 'log precision cmatrix'
    short.name = 'prec'
    initial = '4'
    fixed = 'FALSE'
    prior = 'loggamma'
    param = '1 5e-05'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '20002'
    name = 'log precision random'
    short.name = 'prec.random'
    initial = '4'
    fixed = 'FALSE'
    prior = 'loggamma'
    param = '1 0.001'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Properties: doc = 'A generic model (type 2)'
    constr = 'FALSE'
    nrow.ncol = 'FALSE'
    augmented = 'FALSE'

```

```

aug.factor = '2'
aug.constr = '2'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'generic2'

```

Model 'generic3'. Number of hyperparameters are 11.

Hyperparameter 'theta1' hyperid = '21001'

```

name = 'log precision1'
short.name = 'prec1'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '21002'

```

name = 'log precision2'
short.name = 'prec2'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta3' hyperid = '21003'

```

name = 'log precision3'
short.name = 'prec3'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta4' hyperid = '21004'

```

name = 'log precision4'
short.name = 'prec4'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta5' hyperid = '21005'

```

name = 'log precision5'

```

```

short.name = 'prec5'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta6' hyperid = '21006'
name = 'log precision6'
short.name = 'prec6'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta7' hyperid = '21007'
name = 'log precision7'
short.name = 'prec7'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta8' hyperid = '21008'
name = 'log precision8'
short.name = 'prec8'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta9' hyperid = '21009'
name = 'log precision9'
short.name = 'prec9'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta10' hyperid = '21010'
name = 'log precision10'

```

```

short.name = 'prec10'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1.5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta11' hyperid = '21011'
name = 'log precision common'
short.name = 'prec.common'
initial = '0'
fixed = 'TRUE'
prior = 'loggamma'
param = '1.5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'A generic model (type 3)'
constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
status = 'experimental'
pdf = 'generic3'

```

Model 'spde'. Number of hyperparameters are 4.

```

Hyperparameter 'theta1' hyperid = '22001'
name = 'theta.T'
short.name = 'T'
initial = '2'
fixed = 'FALSE'
prior = 'normal'
param = '0.1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '22002'
name = 'theta.K'
short.name = 'K'
initial = '-2'
fixed = 'FALSE'
prior = 'normal'
param = '0.1'
to.theta = 'function(x) x'

```

```

from.theta = 'function(x) x'
Hyperparameter 'theta3' hyperid = '22003'
  name = 'theta.KT'
  short.name = 'KT'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta4' hyperid = '22004'
  name = 'theta.OC'
  short.name = 'OC'
  initial = '-20'
  fixed = 'TRUE'
  prior = 'normal'
  param = '0 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'A SPDE model'
  constr = 'FALSE'
  nrow.ncol = 'FALSE'
  augmented = 'FALSE'
  aug.factor = '1'
  aug.constr = 'NULL'
  n.div.by = 'NULL'
  n.required = 'TRUE'
  set.default.values = 'TRUE'
  pdf = 'spde'
Model 'spde2'. Number of hyperparameters are 100.
Hyperparameter 'theta1' hyperid = '23001'
  name = 'theta1'
  short.name = 't1'
  initial = '0'
  fixed = 'FALSE'
  prior = 'mvnorm'
  param = '1 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '23002'
  name = 'theta2'
  short.name = 't2'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'

```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta3' hyperid = '23003'
name = 'theta3'
short.name = 't3'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta4' hyperid = '23004'
name = 'theta4'
short.name = 't4'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta5' hyperid = '23005'
name = 'theta5'
short.name = 't5'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta6' hyperid = '23006'
name = 'theta6'
short.name = 't6'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta7' hyperid = '23007'
name = 'theta7'
short.name = 't7'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta8' hyperid = '23008'
name = 'theta8'
short.name = 't8'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta9' hyperid = '23009'
name = 'theta9'
short.name = 't9'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta10' hyperid = '23010'
name = 'theta10'
short.name = 't10'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta11' hyperid = '23011'
name = 'theta11'
short.name = 't11'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta12' hyperid = '23012'
name = 'theta12'
short.name = 't12'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta13' hyperid = '23013'
    name = 'theta13'
    short.name = 't13'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta14' hyperid = '23014'
    name = 'theta14'
    short.name = 't14'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta15' hyperid = '23015'
    name = 'theta15'
    short.name = 't15'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta16' hyperid = '23016'
    name = 'theta16'
    short.name = 't16'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta17' hyperid = '23017'
    name = 'theta17'
    short.name = 't17'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
```

```

    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta18' hyperid = '23018'
    name = 'theta18'
    short.name = 't18'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta19' hyperid = '23019'
    name = 'theta19'
    short.name = 't19'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta20' hyperid = '23020'
    name = 'theta20'
    short.name = 't20'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta21' hyperid = '23021'
    name = 'theta21'
    short.name = 't21'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta22' hyperid = '23022'
    name = 'theta22'
    short.name = 't22'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'

```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta23' hyperid = '23023'
name = 'theta23'
short.name = 't23'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta24' hyperid = '23024'
name = 'theta24'
short.name = 't24'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta25' hyperid = '23025'
name = 'theta25'
short.name = 't25'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta26' hyperid = '23026'
name = 'theta26'
short.name = 't26'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta27' hyperid = '23027'
name = 'theta27'
short.name = 't27'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta28' hyperid = '23028'
    name = 'theta28'
    short.name = 't28'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta29' hyperid = '23029'
    name = 'theta29'
    short.name = 't29'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta30' hyperid = '23030'
    name = 'theta30'
    short.name = 't30'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta31' hyperid = '23031'
    name = 'theta31'
    short.name = 't31'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta32' hyperid = '23032'
    name = 'theta32'
    short.name = 't32'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta33' hyperid = '23033'
name = 'theta33'
short.name = 't33'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta34' hyperid = '23034'
name = 'theta34'
short.name = 't34'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta35' hyperid = '23035'
name = 'theta35'
short.name = 't35'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta36' hyperid = '23036'
name = 'theta36'
short.name = 't36'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta37' hyperid = '23037'
name = 'theta37'
short.name = 't37'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta38' hyperid = '23038'
    name = 'theta38'
    short.name = 't38'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta39' hyperid = '23039'
    name = 'theta39'
    short.name = 't39'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta40' hyperid = '23040'
    name = 'theta40'
    short.name = 't40'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta41' hyperid = '23041'
    name = 'theta41'
    short.name = 't41'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta42' hyperid = '23042'
    name = 'theta42'
    short.name = 't42'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta43' hyperid = '23043'
name = 'theta43'
short.name = 't43'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta44' hyperid = '23044'
name = 'theta44'
short.name = 't44'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta45' hyperid = '23045'
name = 'theta45'
short.name = 't45'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta46' hyperid = '23046'
name = 'theta46'
short.name = 't46'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta47' hyperid = '23047'
name = 'theta47'
short.name = 't47'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta48' hyperid = '23048'
    name = 'theta48'
    short.name = 't48'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta49' hyperid = '23049'
    name = 'theta49'
    short.name = 't49'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta50' hyperid = '23050'
    name = 'theta50'
    short.name = 't50'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta51' hyperid = '23051'
    name = 'theta51'
    short.name = 't51'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta52' hyperid = '23052'
    name = 'theta52'
    short.name = 't52'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta53' hyperid = '23053'
name = 'theta53'
short.name = 't53'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta54' hyperid = '23054'
name = 'theta54'
short.name = 't54'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta55' hyperid = '23055'
name = 'theta55'
short.name = 't55'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta56' hyperid = '23056'
name = 'theta56'
short.name = 't56'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta57' hyperid = '23057'
name = 'theta57'
short.name = 't57'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta58' hyperid = '23058'
name = 'theta58'
short.name = 't58'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta59' hyperid = '23059'
name = 'theta59'
short.name = 't59'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta60' hyperid = '23060'
name = 'theta60'
short.name = 't60'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta61' hyperid = '23061'
name = 'theta61'
short.name = 't61'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta62' hyperid = '23062'
name = 'theta62'
short.name = 't62'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta63' hyperid = '23063'
name = 'theta63'
short.name = 't63'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta64' hyperid = '23064'
name = 'theta64'
short.name = 't64'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta65' hyperid = '23065'
name = 'theta65'
short.name = 't65'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta66' hyperid = '23066'
name = 'theta66'
short.name = 't66'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta67' hyperid = '23067'
name = 'theta67'
short.name = 't67'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta68' hyperid = '23068'
name = 'theta68'
short.name = 't68'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta69' hyperid = '23069'
name = 'theta69'
short.name = 't69'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta70' hyperid = '23070'
name = 'theta70'
short.name = 't70'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta71' hyperid = '23071'
name = 'theta71'
short.name = 't71'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta72' hyperid = '23072'
name = 'theta72'
short.name = 't72'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta73' hyperid = '23073'
name = 'theta73'
short.name = 't73'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta74' hyperid = '23074'
name = 'theta74'
short.name = 't74'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta75' hyperid = '23075'
name = 'theta75'
short.name = 't75'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta76' hyperid = '23076'
name = 'theta76'
short.name = 't76'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta77' hyperid = '23077'
name = 'theta77'
short.name = 't77'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta78' hyperid = '23078'
    name = 'theta78'
    short.name = 't78'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta79' hyperid = '23079'
    name = 'theta79'
    short.name = 't79'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta80' hyperid = '23080'
    name = 'theta80'
    short.name = 't80'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta81' hyperid = '23081'
    name = 'theta81'
    short.name = 't81'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta82' hyperid = '23082'
    name = 'theta82'
    short.name = 't82'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta83' hyperid = '23083'
name = 'theta83'
short.name = 't83'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta84' hyperid = '23084'
name = 'theta84'
short.name = 't84'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta85' hyperid = '23085'
name = 'theta85'
short.name = 't85'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta86' hyperid = '23086'
name = 'theta86'
short.name = 't86'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta87' hyperid = '23087'
name = 'theta87'
short.name = 't87'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta88' hyperid = '23088'
name = 'theta88'
short.name = 't88'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta89' hyperid = '23089'
name = 'theta89'
short.name = 't89'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta90' hyperid = '23090'
name = 'theta90'
short.name = 't90'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta91' hyperid = '23091'
name = 'theta91'
short.name = 't91'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta92' hyperid = '23092'
name = 'theta92'
short.name = 't92'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta93' hyperid = '23093'
name = 'theta93'
short.name = 't93'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta94' hyperid = '23094'
name = 'theta94'
short.name = 't94'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta95' hyperid = '23095'
name = 'theta95'
short.name = 't95'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta96' hyperid = '23096'
name = 'theta96'
short.name = 't96'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta97' hyperid = '23097'
name = 'theta97'
short.name = 't97'
initial = '0'
fixed = 'FALSE'
prior = 'none'
```

```

    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta98' hyperid = '23098'
    name = 'theta98'
    short.name = 't98'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta99' hyperid = '23099'
    name = 'theta99'
    short.name = 't99'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta100' hyperid = '23100'
    name = 'theta100'
    short.name = 't100'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Properties: doc = 'A SPDE2 model'
    constr = 'FALSE'
    nrow.ncol = 'FALSE'
    augmented = 'FALSE'
    aug.factor = '1'
    aug.constr = 'NULL'
    n.div.by = 'NULL'
    n.required = 'TRUE'
    set.default.values = 'TRUE'
    pdf = 'spde2'
Model 'spde3'. Number of hyperparameters are 100.
Hyperparameter 'theta1' hyperid = '24001'
    name = 'theta1'
    short.name = 't1'
    initial = '0'

```

```
fixed = 'FALSE'
prior = 'mvnorm'
param = '1 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '24002'
  name = 'theta2'
  short.name = 't2'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta3' hyperid = '24003'
  name = 'theta3'
  short.name = 't3'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta4' hyperid = '24004'
  name = 'theta4'
  short.name = 't4'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta5' hyperid = '24005'
  name = 'theta5'
  short.name = 't5'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta6' hyperid = '24006'
  name = 'theta6'
  short.name = 't6'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta7' hyperid = '24007'
  name = 'theta7'
  short.name = 't7'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta8' hyperid = '24008'
  name = 'theta8'
  short.name = 't8'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta9' hyperid = '24009'
  name = 'theta9'
  short.name = 't9'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta10' hyperid = '24010'
  name = 'theta10'
  short.name = 't10'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta11' hyperid = '24011'
  name = 'theta11'
  short.name = 't11'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta12' hyperid = '24012'
name = 'theta12'
short.name = 't12'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta13' hyperid = '24013'
name = 'theta13'
short.name = 't13'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta14' hyperid = '24014'
name = 'theta14'
short.name = 't14'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta15' hyperid = '24015'
name = 'theta15'
short.name = 't15'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta16' hyperid = '24016'
name = 'theta16'
short.name = 't16'
initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta17' hyperid = '24017'
name = 'theta17'
short.name = 't17'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta18' hyperid = '24018'
name = 'theta18'
short.name = 't18'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta19' hyperid = '24019'
name = 'theta19'
short.name = 't19'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta20' hyperid = '24020'
name = 'theta20'
short.name = 't20'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta21' hyperid = '24021'
name = 'theta21'
short.name = 't21'
initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta22' hyperid = '24022'
  name = 'theta22'
  short.name = 't22'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta23' hyperid = '24023'
  name = 'theta23'
  short.name = 't23'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta24' hyperid = '24024'
  name = 'theta24'
  short.name = 't24'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta25' hyperid = '24025'
  name = 'theta25'
  short.name = 't25'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta26' hyperid = '24026'
  name = 'theta26'
  short.name = 't26'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta27' hyperid = '24027'
  name = 'theta27'
  short.name = 't27'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta28' hyperid = '24028'
  name = 'theta28'
  short.name = 't28'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta29' hyperid = '24029'
  name = 'theta29'
  short.name = 't29'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta30' hyperid = '24030'
  name = 'theta30'
  short.name = 't30'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta31' hyperid = '24031'
  name = 'theta31'
  short.name = 't31'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta32' hyperid = '24032'
  name = 'theta32'
  short.name = 't32'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta33' hyperid = '24033'
  name = 'theta33'
  short.name = 't33'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta34' hyperid = '24034'
  name = 'theta34'
  short.name = 't34'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta35' hyperid = '24035'
  name = 'theta35'
  short.name = 't35'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta36' hyperid = '24036'
  name = 'theta36'
  short.name = 't36'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta37' hyperid = '24037'
  name = 'theta37'
  short.name = 't37'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta38' hyperid = '24038'
  name = 'theta38'
  short.name = 't38'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta39' hyperid = '24039'
  name = 'theta39'
  short.name = 't39'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta40' hyperid = '24040'
  name = 'theta40'
  short.name = 't40'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta41' hyperid = '24041'
  name = 'theta41'
  short.name = 't41'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta42' hyperid = '24042'
  name = 'theta42'
  short.name = 't42'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta43' hyperid = '24043'
  name = 'theta43'
  short.name = 't43'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta44' hyperid = '24044'
  name = 'theta44'
  short.name = 't44'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta45' hyperid = '24045'
  name = 'theta45'
  short.name = 't45'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta46' hyperid = '24046'
  name = 'theta46'
  short.name = 't46'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta47' hyperid = '24047'
  name = 'theta47'
  short.name = 't47'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta48' hyperid = '24048'
  name = 'theta48'
  short.name = 't48'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta49' hyperid = '24049'
  name = 'theta49'
  short.name = 't49'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta50' hyperid = '24050'
  name = 'theta50'
  short.name = 't50'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta51' hyperid = '24051'
  name = 'theta51'
  short.name = 't51'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta52' hyperid = '24052'
name = 'theta52'
short.name = 't52'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta53' hyperid = '24053'
name = 'theta53'
short.name = 't53'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta54' hyperid = '24054'
name = 'theta54'
short.name = 't54'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta55' hyperid = '24055'
name = 'theta55'
short.name = 't55'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta56' hyperid = '24056'
name = 'theta56'
short.name = 't56'
initial = '0'
```

```

fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta57' hyperid = '24057'
  name = 'theta57'
  short.name = 't57'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta58' hyperid = '24058'
  name = 'theta58'
  short.name = 't58'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta59' hyperid = '24059'
  name = 'theta59'
  short.name = 't59'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta60' hyperid = '24060'
  name = 'theta60'
  short.name = 't60'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta61' hyperid = '24061'
  name = 'theta61'
  short.name = 't61'
  initial = '0'

```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta62' hyperid = '24062'
  name = 'theta62'
  short.name = 't62'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta63' hyperid = '24063'
  name = 'theta63'
  short.name = 't63'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta64' hyperid = '24064'
  name = 'theta64'
  short.name = 't64'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta65' hyperid = '24065'
  name = 'theta65'
  short.name = 't65'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta66' hyperid = '24066'
  name = 'theta66'
  short.name = 't66'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta67' hyperid = '24067'
  name = 'theta67'
  short.name = 't67'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta68' hyperid = '24068'
  name = 'theta68'
  short.name = 't68'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta69' hyperid = '24069'
  name = 'theta69'
  short.name = 't69'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta70' hyperid = '24070'
  name = 'theta70'
  short.name = 't70'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta71' hyperid = '24071'
  name = 'theta71'
  short.name = 't71'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta72' hyperid = '24072'
  name = 'theta72'
  short.name = 't72'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta73' hyperid = '24073'
  name = 'theta73'
  short.name = 't73'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta74' hyperid = '24074'
  name = 'theta74'
  short.name = 't74'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta75' hyperid = '24075'
  name = 'theta75'
  short.name = 't75'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta76' hyperid = '24076'
  name = 'theta76'
  short.name = 't76'
  initial = '0'
```

```

    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta77' hyperid = '24077'
    name = 'theta77'
    short.name = 't77'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta78' hyperid = '24078'
    name = 'theta78'
    short.name = 't78'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta79' hyperid = '24079'
    name = 'theta79'
    short.name = 't79'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta80' hyperid = '24080'
    name = 'theta80'
    short.name = 't80'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta81' hyperid = '24081'
    name = 'theta81'
    short.name = 't81'
    initial = '0'

```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta82' hyperid = '24082'
  name = 'theta82'
  short.name = 't82'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta83' hyperid = '24083'
  name = 'theta83'
  short.name = 't83'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta84' hyperid = '24084'
  name = 'theta84'
  short.name = 't84'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta85' hyperid = '24085'
  name = 'theta85'
  short.name = 't85'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta86' hyperid = '24086'
  name = 'theta86'
  short.name = 't86'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta87' hyperid = '24087'
  name = 'theta87'
  short.name = 't87'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta88' hyperid = '24088'
  name = 'theta88'
  short.name = 't88'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta89' hyperid = '24089'
  name = 'theta89'
  short.name = 't89'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta90' hyperid = '24090'
  name = 'theta90'
  short.name = 't90'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta91' hyperid = '24091'
  name = 'theta91'
  short.name = 't91'
  initial = '0'
```

```
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta92' hyperid = '24092'
  name = 'theta92'
  short.name = 't92'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta93' hyperid = '24093'
  name = 'theta93'
  short.name = 't93'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta94' hyperid = '24094'
  name = 'theta94'
  short.name = 't94'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta95' hyperid = '24095'
  name = 'theta95'
  short.name = 't95'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta96' hyperid = '24096'
  name = 'theta96'
  short.name = 't96'
  initial = '0'
```

```

fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta97' hyperid = '24097'
  name = 'theta97'
  short.name = 't97'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta98' hyperid = '24098'
  name = 'theta98'
  short.name = 't98'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta99' hyperid = '24099'
  name = 'theta99'
  short.name = 't99'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta100' hyperid = '24100'
  name = 'theta100'
  short.name = 't100'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Properties: doc = 'A SPDE3 model'
constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'

```

```

aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'spde3'

```

Model 'iid1d'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '25001'

```

name = 'precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'wishart1d'
param = '2 1e-04'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'Gaussian random effect in dim=1 with Wishart prior'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'TRUE'
pdf = 'iid123d'

```

Model 'iid2d'. Number of hyperparameters are 3.

Hyperparameter 'theta1' hyperid = '26001'

```

name = 'log precision1'
short.name = 'prec1'
initial = '4'
fixed = 'FALSE'
prior = 'wishart2d'
param = '4 1 1 0'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '26002'

```

name = 'log precision2'
short.name = 'prec2'
initial = '4'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

```

Hyperparameter ‘theta3’ hyperid = ‘26003’
  name = ‘logit correlation’
  short.name = ‘cor’
  initial = ‘4’
  fixed = ‘FALSE’
  prior = ‘none’
  param = ‘’
  to.theta = ‘function(x) log((1 + x) / (1 - x))’
  from.theta = ‘function(x) 2 * exp(x) / (1 + exp(x)) - 1’
Properties: doc = ‘Gaussian random effect in dim=2 with Wishart prior’
  constr = ‘FALSE’
  nrow.ncol = ‘FALSE’
  augmented = ‘TRUE’
  aug.factor = ‘1’
  aug.constr = ‘1 2’
  n.div.by = ‘2’
  n.required = ‘TRUE’
  set.default.values = ‘TRUE’
  pdf = ‘iid123d’

```

Model ‘iid3d’. Number of hyperparameters are 6.

```

Hyperparameter ‘theta1’ hyperid = ‘27001’
  name = ‘log precision1’
  short.name = ‘prec1’
  initial = ‘4’
  fixed = ‘FALSE’
  prior = ‘wishart3d’
  param = ‘7 1 1 1 0 0 0’
  to.theta = ‘function(x) log(x)’
  from.theta = ‘function(x) exp(x)’
Hyperparameter ‘theta2’ hyperid = ‘27002’
  name = ‘log precision2’
  short.name = ‘prec2’
  initial = ‘4’
  fixed = ‘FALSE’
  prior = ‘none’
  param = ‘’
  to.theta = ‘function(x) log(x)’
  from.theta = ‘function(x) exp(x)’
Hyperparameter ‘theta3’ hyperid = ‘27003’
  name = ‘log precision3’
  short.name = ‘prec3’
  initial = ‘4’
  fixed = ‘FALSE’
  prior = ‘none’
  param = ‘’

```

```

to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta4' hyperid = '27004'
  name = 'logit correlation12'
  short.name = 'cor12'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta5' hyperid = '27005'
  name = 'logit correlation13'
  short.name = 'cor13'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta6' hyperid = '27006'
  name = 'logit correlation23'
  short.name = 'cor23'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Properties: doc = 'Gaussian random effect in dim=3 with Wishart prior'
  constr = 'FALSE'
  nrow.ncol = 'FALSE'
  augmented = 'TRUE'
  aug.factor = '1'
  aug.constr = '1 2 3'
  n.div.by = '3'
  n.required = 'TRUE'
  set.default.values = 'TRUE'
  pdf = 'iid123d'

```

Model 'iid4d'. Number of hyperparameters are 10.

```

Hyperparameter 'theta1' hyperid = '28001'
  name = 'log precision1'
  short.name = 'prec1'
  initial = '4'
  fixed = 'FALSE'

```

```

    prior = 'wishart4d'
    param = '11 1 1 1 1 0 0 0 0 0'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '28002'
    name = 'log precision2'
    short.name = 'prec2'
    initial = '4'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta3' hyperid = '28003'
    name = 'log precision3'
    short.name = 'prec3'
    initial = '4'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta4' hyperid = '28004'
    name = 'log precision4'
    short.name = 'prec4'
    initial = '4'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta5' hyperid = '28005'
    name = 'logit correlation12'
    short.name = 'cor12'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log((1 + x) / (1 - x))'
    from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta6' hyperid = '28006'
    name = 'logit correlation13'
    short.name = 'cor13'
    initial = '0'
    fixed = 'FALSE'

```

```

prior = 'none'
param = ''
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta7' hyperid = '28007'
name = 'logit correlation14'
short.name = 'cor14'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta8' hyperid = '28008'
name = 'logit correlation23'
short.name = 'cor23'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta9' hyperid = '28009'
name = 'logit correlation24'
short.name = 'cor24'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta10' hyperid = '28010'
name = 'logit correlation34'
short.name = 'cor34'
initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Properties: doc = 'Gaussian random effect in dim=4 with Wishart prior'
constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'TRUE'
aug.factor = '1'

```

```

aug.constr = '1 2 3 4'
n.div.by = '4'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'iid123d'

```

Model 'iid5d'. Number of hyperparameters are 15.

Hyperparameter 'theta1' hyperid = '29001'

```

name = 'log precision1'
short.name = 'prec1'
initial = '4'
fixed = 'FALSE'
prior = 'wishart5d'
param = '16 1 1 1 1 1 0 0 0 0 0 0 0 0 0'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '29002'

```

name = 'log precision2'
short.name = 'prec2'
initial = '4'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta3' hyperid = '29003'

```

name = 'log precision3'
short.name = 'prec3'
initial = '4'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta4' hyperid = '29004'

```

name = 'log precision4'
short.name = 'prec4'
initial = '4'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta5' hyperid = '29005'

```

name = 'log precision5'
short.name = 'prec5'

```

```

initial = '4'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta6' hyperid = '29006'
  name = 'logit correlation12'
  short.name = 'cor12'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta7' hyperid = '29007'
  name = 'logit correlation13'
  short.name = 'cor13'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta8' hyperid = '29008'
  name = 'logit correlation14'
  short.name = 'cor14'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta9' hyperid = '29009'
  name = 'logit correlation15'
  short.name = 'cor15'
  initial = '0'
  fixed = 'FALSE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta10' hyperid = '29010'
  name = 'logit correlation23'
  short.name = 'cor23'

```

```

    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log((1 + x) / (1 - x))'
    from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta11' hyperid = '29011'
    name = 'logit correlation24'
    short.name = 'cor24'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log((1 + x) / (1 - x))'
    from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta12' hyperid = '29012'
    name = 'logit correlation25'
    short.name = 'cor25'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log((1 + x) / (1 - x))'
    from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta13' hyperid = '29013'
    name = 'logit correlation34'
    short.name = 'cor34'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log((1 + x) / (1 - x))'
    from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta14' hyperid = '29014'
    name = 'logit correlation35'
    short.name = 'cor35'
    initial = '0'
    fixed = 'FALSE'
    prior = 'none'
    param = ''
    to.theta = 'function(x) log((1 + x) / (1 - x))'
    from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta15' hyperid = '29015'
    name = 'logit correlation45'
    short.name = 'cor45'

```

```

initial = '0'
fixed = 'FALSE'
prior = 'none'
param = ''
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Properties: doc = 'Gaussian random effect in dim=5 with Wishart prior'
constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'TRUE'
aug.factor = '1'
aug.constr = '1 2 3 4 5'
n.div.by = '5'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'iid123d'

```

Model '2diid'. Number of hyperparameters are 3.

```

Hyperparameter 'theta1' hyperid = '30001'
name = 'log precision1'
short.name = 'prec1'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

```

Hyperparameter 'theta2' hyperid = '30002'
name = 'log precision2'
short.name = 'prec2'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

```

Hyperparameter 'theta3' hyperid = '30003'
name = 'correlation'
short.name = 'cor'
initial = '4'
fixed = 'FALSE'
prior = 'normal'
param = '0 0.15'
to.theta = 'function(x) log((1 + x) / (1 - x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'

```

```

Properties: doc = '(This model is obsolete)'

```

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = '1 2'
n.div.by = '2'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'iid123d'

```

Model 'z'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '31001'
name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'The z-model in a classical mixed model formulation'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
pdf = 'z'
status = 'experimental'

```

Model 'rw2d'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '32001'
name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'Thin-plate spline model'

```

constr = 'TRUE'
nrow.ncol = 'TRUE'
augmented = 'FALSE'
aug.factor = '1'

```

```

aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'TRUE'
pdf = 'rw2d'

```

Model 'rw2diid'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '33001'

```

name = 'log precision'
short.name = 'prec'
prior = 'pc.prec'
param = '1 0.01'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '33002'

```

name = 'logit phi'
short.name = 'phi'
prior = 'pc'
param = '0.5 0.5'
initial = '3'
fixed = 'FALSE'
to.theta = 'function(x) log(x / (1 - x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

Properties: doc = 'Thin-plate spline with iid noise'

```

constr = 'TRUE'
nrow.ncol = 'TRUE'
augmented = 'TRUE'
aug.factor = '2'
aug.constr = '2'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'TRUE'
status = 'experimental'
pdf = 'rw2diid'

```

Model 'slm'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '34001'

```

name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter ‘theta2’ hyperid = ‘34002’
 name = ‘rho’
 short.name = ‘rho’
 initial = ‘0’
 fixed = ‘FALSE’
 prior = ‘normal’
 param = ‘0 10’
 to.theta = ‘function(x) log(x / (1 -x))’
 from.theta = ‘function(x) 1 / (1 + exp(-x))’
Properties: doc = ‘Spatial lag model’
 constr = ‘FALSE’
 nrow.ncol = ‘FALSE’
 augmented = ‘FALSE’
 aug.factor = ‘1’
 aug.constr = ‘NULL’
 n.div.by = ‘NULL’
 n.required = ‘TRUE’
 set.default.values = ‘TRUE’
 pdf = ‘slm’
 status = ‘experimental’

Model ‘matern2d’. Number of hyperparameters are 2.

Hyperparameter ‘theta1’ hyperid = ‘35001’
 name = ‘log precision’
 short.name = ‘prec’
 initial = ‘4’
 fixed = ‘FALSE’
 prior = ‘loggamma’
 param = ‘1 5e-05’
 to.theta = ‘function(x) log(x)’
 from.theta = ‘function(x) exp(x)’

Hyperparameter ‘theta2’ hyperid = ‘35002’
 name = ‘log range’
 short.name = ‘range’
 initial = ‘2’
 fixed = ‘FALSE’
 prior = ‘loggamma’
 param = ‘1 0.01’
 to.theta = ‘function(x) log(x)’
 from.theta = ‘function(x) exp(x)’

Properties: doc = ‘Matern covariance function on a regular grid’
 constr = ‘FALSE’
 nrow.ncol = ‘TRUE’
 augmented = ‘FALSE’
 aug.factor = ‘1’
 aug.constr = ‘NULL’

```

n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'TRUE'
pdf = 'matern2d'

```

Model 'dmatern'. Number of hyperparameters are 3.

Hyperparameter 'theta1' hyperid = '35101'

```

name = 'log precision'
short.name = 'prec'
initial = '3'
fixed = 'FALSE'
prior = 'pc.prec'
param = '1 0.01'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '35102'

```

name = 'log range'
short.name = 'range'
initial = '0'
fixed = 'FALSE'
prior = 'pc.range'
param = '1 0.5'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta3' hyperid = '35103'

```

name = 'log nu'
short.name = 'nu'
initial = '-0.693147180559945'
fixed = 'TRUE'
prior = 'loggamma'
param = '0.5 1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'Dense Matern field'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'TRUE'
set.default.values = 'TRUE'
status = 'experimental'
pdf = 'dmatern'

```

Model 'copy'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '36001'

```

name = 'beta'
short.name = 'b'
initial = '1'
fixed = 'TRUE'
prior = 'normal'
param = '1 10'
to.theta = 'function(x,REPLACE.ME.low,REPLACE.ME.high) {} if (all(is.infinite(c(low,high)
  || low == high) {}) return(x) else if (all(is.finite(c(low,high)))) {}
  stopifnot(low < high) return(log(-(low-x) / (high-x))) else if (is.finite(low)
  && is.infinite(high) && high > low) {} return(log(x -low)) else {} stop("Condition
  not yet implemented") '
from.theta = 'function(x,REPLACE.ME.low,REPLACE.ME.high) {} if (all(is.infinite(c(low,high)
  || low == high) {}) return(x) else if (all(is.finite(c(low,high)))) {}
  stopifnot(low < high) return(low + exp(x) / (1 + exp(x)) * (high -low)) else
  if (is.finite(low) && is.infinite(high) && high > low) {} return(low + exp(x))
  else {} stop("Condition not yet implemented") '

```

Properties: **doc** = 'Create a copy of a model component'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
pdf = 'NA'

```

Model 'clinear'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '37001'

```

name = 'beta'
short.name = 'b'
initial = '1'
fixed = 'FALSE'
prior = 'normal'
param = '1 10'
to.theta = 'function(x,REPLACE.ME.low,REPLACE.ME.high) {} if (all(is.infinite(c(low,high)
  || low == high) {}) stopifnot(low < high) return(x) else if (all(is.finite(c(low,high)
  {} stopifnot(low < high) return(log(-(low-x) / (high-x))) else if (is.finite(low)
  && is.infinite(high) && high > low) {} return(log(x -low)) else {} stop("Condition
  not yet implemented") '
from.theta = 'function(x,REPLACE.ME.low,REPLACE.ME.high) {} if (all(is.infinite(c(low,high)
  || low == high) {}) stopifnot(low < high) return(x) else if (all(is.finite(c(low,high)
  {} stopifnot(low < high) return(low + exp(x) / (1 + exp(x)) * (high -low))
  else if (is.finite(low) && is.infinite(high) && high > low) {} return(low
  + exp(x)) else {} stop("Condition not yet implemented") '

```

Properties: **doc** = 'Constrained linear effect'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'

```

```

aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
pdf = 'clinear'

```

Model 'sigm'. Number of hyperparameters are 3.

Hyperparameter 'theta1' hyperid = '38001'

```

name = 'beta'
short.name = 'b'
initial = '1'
fixed = 'FALSE'
prior = 'normal'
param = '1 10'
to.theta = 'function(x) x'
from.theta = 'function(x) x'

```

Hyperparameter 'theta2' hyperid = '38002'

```

name = 'loghalflife'
short.name = 'halflife'
initial = '3'
fixed = 'FALSE'
prior = 'loggamma'
param = '3 1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta3' hyperid = '38003'

```

name = 'logshape'
short.name = 'shape'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '10 10'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'Sigmoidal effect of a covariate'

```

constr = 'FALSE'
nrow.ncol = 'FALSE'
augmented = 'FALSE'
aug.factor = '1'
aug.constr = 'NULL'
n.div.by = 'NULL'
n.required = 'FALSE'
set.default.values = 'FALSE'
status = 'experimental'
pdf = 'sigm'

```

Model ‘revsigm’. Number of hyperparameters are 3.

Hyperparameter ‘theta1’ hyperid = ‘39001’

name = ‘beta’
short.name = ‘b’
initial = ‘1’
fixed = ‘FALSE’
prior = ‘normal’
param = ‘1 10’
to.theta = ‘function(x) x’
from.theta = ‘function(x) x’

Hyperparameter ‘theta2’ hyperid = ‘39002’

name = ‘loghalflife’
short.name = ‘halflife’
initial = ‘3’
fixed = ‘FALSE’
prior = ‘loggamma’
param = ‘3 1’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

Hyperparameter ‘theta3’ hyperid = ‘39003’

name = ‘logshape’
short.name = ‘shape’
initial = ‘0’
fixed = ‘FALSE’
prior = ‘loggamma’
param = ‘10 10’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

Properties: doc = ‘Reverse sigmoidal effect of a covariate’

constr = ‘FALSE’
nrow.ncol = ‘FALSE’
augmented = ‘FALSE’
aug.factor = ‘1’
aug.constr = ‘NULL’
n.div.by = ‘NULL’
n.required = ‘FALSE’
set.default.values = ‘FALSE’
status = ‘experimental’
pdf = ‘sigm’

Model ‘log1exp’. Number of hyperparameters are 3.

Hyperparameter ‘theta1’ hyperid = ‘39011’

name = ‘beta’
short.name = ‘b’
initial = ‘1’
fixed = ‘FALSE’

```

    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '39012'
    name = 'alpha'
    short.name = 'a'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta3' hyperid = '39013'
    name = 'gamma'
    short.name = 'g'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Properties: doc = 'A nonlinear model of a covariate'
    constr = 'FALSE'
    nrow.ncol = 'FALSE'
    augmented = 'FALSE'
    aug.factor = '1'
    aug.constr = 'NULL'
    n.div.by = 'NULL'
    n.required = 'FALSE'
    set.default.values = 'FALSE'
    status = 'experimental'
    pdf = 'log1exp'

```

Model 'logdist'. Number of hyperparameters are 3.

```

Hyperparameter 'theta1' hyperid = '39021'
    name = 'beta'
    short.name = 'b'
    initial = '1'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '39022'
    name = 'alpha1'

```

```

short.name = 'a1'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '0.1 1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta3' hyperid = '39023'
  name = 'alpha2'
  short.name = 'a2'
  initial = '0'
  fixed = 'FALSE'
  prior = 'loggamma'
  param = '0.1 1'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'A nonlinear model of a covariate'
  constr = 'FALSE'
  nrow.ncol = 'FALSE'
  augmented = 'FALSE'
  aug.factor = '1'
  aug.constr = 'NULL'
  n.div.by = 'NULL'
  n.required = 'FALSE'
  set.default.values = 'FALSE'
  status = 'experimental'
  pdf = 'logdist'

```

Section 'group'. Valid models in this section are:

Model 'exchangeable'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '40001'
  name = 'logit correlation'
  short.name = 'rho'
  initial = '1'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 0.2'
  to.theta = 'function(x,REPLACE.ME.ngroup) log((1 + x * (ngroup - 1)) / (1 - x))'
  from.theta = 'function(x,REPLACE.ME.ngroup) (exp(x) - 1) / (exp(x) + ngroup - 1)'

```

Properties: doc = 'Exchangeable correlations'

Model 'exchangeablepos'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '40101'
  name = 'logit correlation'
  short.name = 'rho'
  initial = '1'

```

```

fixed = 'FALSE'
prior = 'pc.cor0'
param = '0.5 0.5'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

Properties: **doc** = 'Exchangeable positive correlations'

Model 'ar1'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '41001'

```

name = 'logit correlation'
short.name = 'rho'
initial = '2'
fixed = 'FALSE'
prior = 'normal'
param = '0 0.15'
to.theta = 'function(x) log((1 + x) / (1 -x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'

```

Properties: **doc** = 'AR(1) correlations'

Model 'ar'. Number of hyperparameters are 11.

Hyperparameter 'theta1' hyperid = '42001'

```

name = 'log precision'
short.name = 'prec'
initial = '0'
fixed = 'TRUE'
prior = 'pc.prec'
param = '3 0.01'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '42002'

```

name = 'pacf1'
short.name = 'pacf1'
initial = '2'
fixed = 'FALSE'
prior = 'pc.cor0'
param = '0.5 0.5'
to.theta = 'function(x) log((1 + x) / (1 -x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'

```

Hyperparameter 'theta3' hyperid = '42003'

```

name = 'pacf2'
short.name = 'pacf2'
initial = '0'
fixed = 'FALSE'
prior = 'pc.cor0'
param = '0.5 0.4'
to.theta = 'function(x) log((1 + x) / (1 -x))'
from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) -1'

```

```

Hyperparameter 'theta4' hyperid = '42004'
  name = 'pacf3'
  short.name = 'pacf3'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.3'
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta5' hyperid = '42005'
  name = 'pacf4'
  short.name = 'pacf4'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.2'
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta6' hyperid = '42006'
  name = 'pacf5'
  short.name = 'pacf5'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta7' hyperid = '42007'
  name = 'pacf6'
  short.name = 'pacf6'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'
Hyperparameter 'theta8' hyperid = '42008'
  name = 'pacf7'
  short.name = 'pacf7'
  initial = '0'
  fixed = 'FALSE'
  prior = 'pc.cor0'
  param = '0.5 0.1'
  to.theta = 'function(x) log((1 + x) / (1 - x))'
  from.theta = 'function(x) 2 * exp(x) / (1 + exp(x)) - 1'

```

```

Hyperparameter ‘theta9’ hyperid = ‘42009’
  name = ‘pacf8’
  short.name = ‘pacf8’
  initial = ‘0’
  fixed = ‘FALSE’
  prior = ‘pc.cor0’
  param = ‘0.5 0.1’
  to.theta = ‘function(x) log((1 + x) / (1 - x))’
  from.theta = ‘function(x) 2 * exp(x) / (1 + exp(x)) - 1’
Hyperparameter ‘theta10’ hyperid = ‘42010’
  name = ‘pacf9’
  short.name = ‘pacf9’
  initial = ‘0’
  fixed = ‘FALSE’
  prior = ‘pc.cor0’
  param = ‘0.5 0.1’
  to.theta = ‘function(x) log((1 + x) / (1 - x))’
  from.theta = ‘function(x) 2 * exp(x) / (1 + exp(x)) - 1’
Hyperparameter ‘theta11’ hyperid = ‘42011’
  name = ‘pacf10’
  short.name = ‘pacf10’
  initial = ‘0’
  fixed = ‘FALSE’
  prior = ‘pc.cor0’
  param = ‘0.5 0.1’
  to.theta = ‘function(x) log((1 + x) / (1 - x))’
  from.theta = ‘function(x) 2 * exp(x) / (1 + exp(x)) - 1’

```

Properties: doc = ‘AR(p) correlations’

Model ‘rw1’. Number of hyperparameters are 1.

```

Hyperparameter ‘theta’ hyperid = ‘43001’
  name = ‘log precision’
  short.name = ‘prec’
  prior = ‘loggamma’
  param = ‘1 5e-05’
  initial = ‘0’
  fixed = ‘TRUE’
  to.theta = ‘function(x) log(x)’
  from.theta = ‘function(x) exp(x)’

```

Properties: doc = ‘Random walk of order 1’

Model ‘rw2’. Number of hyperparameters are 1.

```

Hyperparameter ‘theta’ hyperid = ‘44001’
  name = ‘log precision’
  short.name = ‘prec’
  prior = ‘loggamma’
  param = ‘1 5e-05’

```

```

initial = '0'
fixed = 'TRUE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: **doc** = 'Random walk of order 2'

Model 'besag'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '45001'

```

name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '0'
fixed = 'TRUE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: **doc** = 'Besag model'

Model 'iid'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '46001'

```

name = 'log precision'
short.name = 'prec'
prior = 'loggamma'
param = '1 5e-05'
initial = '0'
fixed = 'TRUE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: **doc** = 'Independent model'

Section 'mix'. Valid models in this section are:

Model 'gaussian'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '47001'

```

name = 'log precision'
short.name = 'prec'
prior = 'pc.prec'
param = '1 0.01'
initial = '0'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: **doc** = 'Gaussian mixture'

Model 'loggamma'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '47101'

```

name = 'log precision'
short.name = 'prec'
prior = 'pc.mgamma'
param = '4.8'

```

```

initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: **doc** = 'LogGamma mixture'

Model 'mloggamma'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '47201'

```

name = 'log precision'
short.name = 'prec'
prior = 'pc.mgamma'
param = '4.8'
initial = '4'
fixed = 'FALSE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: **doc** = 'Minus-LogGamma mixture'

Section 'link'. Valid models in this section are:

Model 'default'. Number of hyperparameters are 0.

Model 'cloglog'. Number of hyperparameters are 0.

Model 'loglog'. Number of hyperparameters are 0.

Model 'identity'. Number of hyperparameters are 0.

Model 'inverse'. Number of hyperparameters are 0.

Model 'log'. Number of hyperparameters are 0.

Model 'loga'. Number of hyperparameters are 0.

Model 'neglog'. Number of hyperparameters are 0.

Model 'logit'. Number of hyperparameters are 0.

Model 'probit'. Number of hyperparameters are 0.

Model 'cauchit'. Number of hyperparameters are 0.

Model 'tan'. Number of hyperparameters are 0.

Model 'quantile'. Number of hyperparameters are 0.

Model 'pquantile'. Number of hyperparameters are 0.

Model 'sslogit'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '48001'

```

name = 'sensitivity'
short.name = 'sens'
prior = 'logitbeta'
param = '10 5'
initial = '1'
fixed = 'FALSE'
to.theta = 'function(x) log(x / (1 - x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

Hyperparameter 'theta2' hyperid = '48002'

```

name = 'specificity'
short.name = 'spec'
prior = 'logitbeta'

```

```

param = '10 5'
initial = '1'
fixed = 'FALSE'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Logit link with sensitivity and specificity'
status = 'disabled'
pdf = 'NA'

```

Model 'logoffset'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '49001'
name = 'beta'
short.name = 'b'
prior = 'normal'
param = '0 100'
initial = '0'
fixed = 'TRUE'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'Log-link with an offset'
pdf = 'logoffset'

```

Model 'logitoffset'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '49011'
name = 'prob'
short.name = 'p'
prior = 'normal'
param = '-1 100'
initial = '-1'
fixed = 'FALSE'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Logit-link with an offset'
status = 'experimental'
pdf = 'logitoffset'

```

Model 'robit'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '49021'
name = 'log degrees of freedom'
short.name = 'dof'
initial = '1.6094379124341'
fixed = 'TRUE'
prior = 'pc.dof'
param = '50 0.5'
to.theta = 'function(x) log(x -2)'
from.theta = 'function(x) 2 + exp(x)'
Properties: doc = 'Robit link'
status = 'experimental'

```

pdf = 'robit'

Model 'sn'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '49031'

name = 'skewness'

short.name = 'skew'

initial = '0.00123456789'

fixed = 'FALSE'

prior = 'pc.sn'

param = '10'

to.theta = 'function(x, skew.max = 0.988) log((1 + x / skew.max) / (1 - x / skew.max))'

from.theta = 'function(x, skew.max = 0.988) skew.max * (2 * exp(x) / (1 + exp(x)) - 1)'

Hyperparameter 'theta2' hyperid = '49032'

name = 'intercept'

short.name = 'intercept'

initial = '0'

fixed = 'FALSE'

prior = 'linksnintercept'

param = '0 0'

to.theta = 'function(x) log(x / (1 - x))'

from.theta = 'function(x) exp(x) / (1 + exp(x))'

Properties: doc = 'Skew-normal link'

pdf = 'linksn'

Model 'test1'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '50001'

name = 'beta'

short.name = 'b'

prior = 'normal'

param = '0 100'

initial = '0'

fixed = 'FALSE'

to.theta = 'function(x) x'

from.theta = 'function(x) x'

Properties: doc = 'A test1-link function (experimental)'

pdf = 'NA'

Model 'special1'. Number of hyperparameters are 11.

Hyperparameter 'theta1' hyperid = '51001'

name = 'log precision'

short.name = 'prec'

initial = '0'

fixed = 'FALSE'

prior = 'loggamma'

param = '1 1'

to.theta = 'function(x) x'

from.theta = 'function(x) x'

```
Hyperparameter 'theta2' hyperid = '51002'  
  name = 'beta1'  
  short.name = 'beta1'  
  initial = '0'  
  fixed = 'FALSE'  
  prior = 'mvnorm'  
  param = '0 100'  
  to.theta = 'function(x) x'  
  from.theta = 'function(x) x'  
Hyperparameter 'theta3' hyperid = '51003'  
  name = 'beta2'  
  short.name = 'beta2'  
  initial = '0'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) x'  
  from.theta = 'function(x) x'  
Hyperparameter 'theta4' hyperid = '51004'  
  name = 'beta3'  
  short.name = 'beta3'  
  initial = '0'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) x'  
  from.theta = 'function(x) x'  
Hyperparameter 'theta5' hyperid = '51005'  
  name = 'beta4'  
  short.name = 'beta4'  
  initial = '0'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) x'  
  from.theta = 'function(x) x'  
Hyperparameter 'theta6' hyperid = '51006'  
  name = 'beta5'  
  short.name = 'beta5'  
  initial = '0'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) x'  
  from.theta = 'function(x) x'
```

Hyperparameter 'theta7' hyperid = '51007'

```
name = 'beta6'  
short.name = 'beta6'  
initial = '0'  
fixed = 'FALSE'  
prior = 'none'  
param = ''  
to.theta = 'function(x) x'  
from.theta = 'function(x) x'
```

Hyperparameter 'theta8' hyperid = '51008'

```
name = 'beta7'  
short.name = 'beta7'  
initial = '0'  
fixed = 'FALSE'  
prior = 'none'  
param = ''  
to.theta = 'function(x) x'  
from.theta = 'function(x) x'
```

Hyperparameter 'theta9' hyperid = '51009'

```
name = 'beta8'  
short.name = 'beta8'  
initial = '0'  
fixed = 'FALSE'  
prior = 'none'  
param = ''  
to.theta = 'function(x) x'  
from.theta = 'function(x) x'
```

Hyperparameter 'theta10' hyperid = '51010'

```
name = 'beta9'  
short.name = 'beta9'  
initial = '0'  
fixed = 'FALSE'  
prior = 'none'  
param = ''  
to.theta = 'function(x) x'  
from.theta = 'function(x) x'
```

Hyperparameter 'theta11' hyperid = '51011'

```
name = 'beta10'  
short.name = 'beta10'  
initial = '0'  
fixed = 'FALSE'  
prior = 'none'  
param = ''  
to.theta = 'function(x) x'  
from.theta = 'function(x) x'
```

Properties: **doc** = 'A special1-link function (experimental)'
pdf = 'NA'

Model 'special2'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '52001'
name = 'beta'
short.name = 'b'
prior = 'normal'
param = '0 10'
initial = '0'
fixed = 'FALSE'
to.theta = 'function(x) x'
from.theta = 'function(x) x'

Properties: **doc** = 'A special2-link function (experimental)'
pdf = 'NA'

Section 'predictor'. Valid models in this section are:

Model 'predictor'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '53001'
name = 'log precision'
short.name = 'prec'
initial = '13.8155105579643'
fixed = 'TRUE'
prior = 'loggamma'
param = '1 1e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

Properties: **doc** = '(do not use)'

Section 'hazard'. Valid models in this section are:

Model 'rw1'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '54001'
name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

Properties: **doc** = 'A random walk of order 1 for the log-hazard'

Model 'rw2'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '55001'
name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'

```

param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'A random walk of order 2 for the log-hazard'

Model 'iid'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '55501'
name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'An iid model for the log-hazard'

Section 'likelihood'. Valid models in this section are:

Model 'poisson'. Number of hyperparameters are 0.

Model 'xpoisson'. Number of hyperparameters are 0.

Model 'cenpoisson'. Number of hyperparameters are 0.

Model 'cenpoisson2'. Number of hyperparameters are 0.

Model 'gpoisson'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '56001'
name = 'overdispersion'
short.name = 'phi'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

```

Hyperparameter 'theta2' hyperid = '56002'

```

```

name = 'p'
short.name = 'p'
initial = '1'
fixed = 'TRUE'
prior = 'normal'
param = '1 100'
to.theta = 'function(x) x'
from.theta = 'function(x) x'

```

Properties: doc = 'The generalized Poisson likelihood'

```

survival = 'FALSE'
discrete = 'TRUE'
link = 'default log logoffset'
pdf = 'gpoisson'
status = 'experimental'

```

Model ‘poisson.special1’. Number of hyperparameters are 1.

Hyperparameter ‘theta’ hyperid = ‘56100’
 name = ‘logit probability’
 short.name = ‘prob’
 initial = ‘-1’
 fixed = ‘FALSE’
 prior = ‘gaussian’
 param = ‘-1 0.2’
 to.theta = ‘function(x) log(x / (1 -x))’
 from.theta = ‘function(x) exp(x) / (1 + exp(x))’
Properties: doc = ‘The Poisson.special1 likelihood’
 survival = ‘FALSE’
 discrete = ‘TRUE’
 link = ‘default log’
 pdf = ‘poisson-special’

Model ‘binomial’. Number of hyperparameters are 0.

Model ‘xbinomial’. Number of hyperparameters are 0.

Model ‘pom’. Number of hyperparameters are 10.

Hyperparameter ‘theta1’ hyperid = ‘57101’
 name = ‘theta1’
 short.name = ‘theta1’
 initial = ‘NA’
 fixed = ‘FALSE’
 prior = ‘dirichlet’
 param = ‘3’
 to.theta = ‘function(x) x’
 from.theta = ‘function(x) x’
Hyperparameter ‘theta2’ hyperid = ‘57102’
 name = ‘theta2’
 short.name = ‘theta2’
 initial = ‘NA’
 fixed = ‘FALSE’
 prior = ‘none’
 param = ‘’
 to.theta = ‘function(x) log(x)’
 from.theta = ‘function(x) exp(x)’
Hyperparameter ‘theta3’ hyperid = ‘57103’
 name = ‘theta3’
 short.name = ‘theta3’
 initial = ‘NA’
 fixed = ‘FALSE’
 prior = ‘none’
 param = ‘’
 to.theta = ‘function(x) log(x)’
 from.theta = ‘function(x) exp(x)’

```
Hyperparameter 'theta4' hyperid = '57104'  
  name = 'theta4'  
  short.name = 'theta4'  
  initial = 'NA'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) log(x)'  
  from.theta = 'function(x) exp(x)'  
Hyperparameter 'theta5' hyperid = '57105'  
  name = 'theta5'  
  short.name = 'theta5'  
  initial = 'NA'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) log(x)'  
  from.theta = 'function(x) exp(x)'  
Hyperparameter 'theta6' hyperid = '57106'  
  name = 'theta6'  
  short.name = 'theta6'  
  initial = 'NA'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) log(x)'  
  from.theta = 'function(x) exp(x)'  
Hyperparameter 'theta7' hyperid = '57107'  
  name = 'theta7'  
  short.name = 'theta7'  
  initial = 'NA'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) log(x)'  
  from.theta = 'function(x) exp(x)'  
Hyperparameter 'theta8' hyperid = '57108'  
  name = 'theta8'  
  short.name = 'theta8'  
  initial = 'NA'  
  fixed = 'FALSE'  
  prior = 'none'  
  param = ''  
  to.theta = 'function(x) log(x)'  
  from.theta = 'function(x) exp(x)'
```

Hyperparameter ‘theta9’ hyperid = ‘57109’

```

name = ‘theta9’
short.name = ‘theta9’
initial = ‘NA’
fixed = ‘FALSE’
prior = ‘none’
param = ‘’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

```

Hyperparameter ‘theta10’ hyperid = ‘57110’

```

name = ‘theta10’
short.name = ‘theta10’
initial = ‘NA’
fixed = ‘FALSE’
prior = ‘none’
param = ‘’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

```

Properties: doc = ‘Likelihood for the proportional odds model’

```

status = ‘experimental’
survival = ‘FALSE’
discrete = ‘TRUE’
link = ‘default identity’
pdf = ‘pom’

```

Model ‘bgev’. Number of hyperparameters are 12.

Hyperparameter ‘theta1’ hyperid = ‘57201’

```

name = ‘spread’
short.name = ‘sd’
initial = ‘0’
fixed = ‘FALSE’
prior = ‘loggamma’
param = ‘1 3’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

```

Hyperparameter ‘theta2’ hyperid = ‘57202’

```

name = ‘tail’
short.name = ‘xi’
initial = ‘-4’
fixed = ‘FALSE’
prior = ‘pc.gevtail’
param = ‘7 0 0.5’
to.theta = ‘function(x, interval = c(REPLACE.ME.low, REPLACE.ME.high)) log(-(interval[1]
-x) / (interval[2] -x))’
from.theta = ‘function(x, interval = c(REPLACE.ME.low, REPLACE.ME.high))
interval[1] + (interval[2] -interval[1]) * exp(x) / (1.0 + exp(x))’

```

Hyperparameter ‘theta3’ hyperid = ‘57203’

```
name = 'beta1'
short.name = 'beta1'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta4' hyperid = '57204'
name = 'beta2'
short.name = 'beta2'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta5' hyperid = '57205'
name = 'beta3'
short.name = 'beta3'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta6' hyperid = '57206'
name = 'beta4'
short.name = 'beta4'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta7' hyperid = '57207'
name = 'beta5'
short.name = 'beta5'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta8' hyperid = '57208'
```

```

name = 'beta6'
short.name = 'beta6'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta9' hyperid = '57209'
name = 'beta7'
short.name = 'beta7'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta10' hyperid = '57210'
name = 'beta8'
short.name = 'beta8'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta11' hyperid = '57211'
name = 'beta9'
short.name = 'beta9'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta12' hyperid = '57212'
name = 'beta10'
short.name = 'beta'
initial = 'NA'
fixed = 'FALSE'
prior = 'normal'
param = '0.300'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Properties: doc = 'The blended Generalized Extreme Value likelihood'

```

```

status = 'experimental'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default identity log'
pdf = 'bgev'

```

Model 'gamma'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '58001'
name = 'precision parameter'
short.name = 'prec'
initial = '4.60517018598809'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 0.01'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

```

Properties: doc = 'The Gamma likelihood'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log quantile'
pdf = 'gamma'

```

Model 'gammasurv'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '58101'
name = 'precision parameter'
short.name = 'prec'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 0.01'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

```

Properties: doc = 'The Gamma likelihood (survival)'
survival = 'TRUE'
discrete = 'FALSE'
status = 'experimental'
link = 'default log'
pdf = 'gammasurv'

```

Model 'gammajw'. Number of hyperparameters are 0.

Model 'gammajwsurv'. Number of hyperparameters are 0.

Model 'gammacount'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '59001'
name = 'log alpha'
short.name = 'alpha'
initial = '0'
fixed = 'FALSE'
prior = 'pc.gammacount'

```

```

param = '3'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'A Gamma generalisation of the Poisson likelihood'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
status = 'experimental'
pdf = 'gammacount'

```

Model 'qkumar'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '60001'

```

name = 'precision parameter'
short.name = 'prec'
initial = '1'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 0.1'
to.theta = 'function(x,sc = 0.1) log(x) / sc'
from.theta = 'function(x,sc = 0.1) exp(sc * x)'

```

Properties: doc = 'A quantile version of the Kumar likelihood'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga cauchit'
pdf = 'qkumar'

```

Model 'qloglogistic'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '60011'

```

name = 'log alpha'
short.name = 'alpha'
initial = '1'
fixed = 'FALSE'
prior = 'loggamma'
param = '25 25'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'A quantile loglogistic likelihood'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default log neglog'
pdf = 'qloglogistic'

```

Model 'qloglogisticsurv'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '60021'

```

name = 'log alpha'
short.name = 'alpha'
initial = '1'
fixed = 'FALSE'

```

```

prior = 'loggamma'
param = '25 25'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'A quantile loglogistic likelihood (survival)'
survival = 'TRUE'
discrete = 'FALSE'
link = 'default log neglog'
pdf = 'qloglogistic'

```

Model 'beta'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '61001'
name = 'precision parameter'
short.name = 'phi'
initial = '2.30258509299405'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 0.1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'The Beta likelihood'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga cauchit probit cloglog loglog'
pdf = 'beta'

```

Model 'betabinomial'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '62001'
name = 'overdispersion'
short.name = 'rho'
initial = '0'
fixed = 'FALSE'
prior = 'gaussian'
param = '0 0.4'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'The Beta-Binomial likelihood'
survival = 'FALSE'
discrete = 'TRUE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'betabinomial'

```

Model 'betabinomialna'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '62101'
name = 'overdispersion'
short.name = 'rho'
initial = '0'
fixed = 'FALSE'

```

```

prior = 'gaussian'
param = '0 0.4'
to.theta = 'function(x) log(x / (1 - x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'The Beta-Binomial Normal approximation likelihood'
survival = 'FALSE'
discrete = 'TRUE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'betabinomialna'

```

Model 'cbinomial'. Number of hyperparameters are 0.

Model 'nbinomial'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '63001'

```

name = 'size'
short.name = 'size'
initial = '2.30258509299405'
fixed = 'FALSE'
prior = 'pc.mgamma'
param = '7'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'The negBinomial likelihood'

```

survival = 'FALSE'
discrete = 'TRUE'
link = 'default log logoffset quantile'
pdf = 'nbinomial'

```

Model 'nbinomial2'. Number of hyperparameters are 0.

Model 'simplex'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '64001'

```

name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'The simplex likelihood'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga cauchit probit cloglog loglog'
pdf = 'simplex'

```

Model 'gaussian'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '65001'

```

name = 'log precision'
short.name = 'prec'

```

```

initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '65002'
  name = 'log precision offset'
  short.name = 'precoffset'
  initial = '72.0873067782343'
  fixed = 'TRUE'
  prior = 'none'
  param = ''
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'The Gaussian likelihood'
  survival = 'FALSE'
  discrete = 'FALSE'
  link = 'default identity logit loga cauchit log logoffset'
  pdf = 'gaussian'

```

Model 'agaussian'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '66001'
  name = 'log precision'
  short.name = 'prec'
  initial = '4'
  fixed = 'FALSE'
  prior = 'loggamma'
  param = '1 5e-05'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'The aggregated Gaussian likelihood'
  status = 'experimental'
  survival = 'FALSE'
  discrete = 'FALSE'
  link = 'default identity logit loga cauchit log logoffset'
  pdf = 'agaussian'

```

Model 'circularnormal'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '67001'
  name = 'log precision parameter'
  short.name = 'prec'
  initial = '2'
  fixed = 'FALSE'
  prior = 'loggamma'
  param = '1 0.01'
  to.theta = 'function(x) log(x)'

```

```

from.theta = 'function(x) exp(x)'
Properties: doc = 'The circular Gaussian likelihood'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default tan'
pdf = 'circular-normal'
status = 'experimental'
Model 'wrappedcauchy'. Number of hyperparameters are 1.
Hyperparameter 'theta' hyperid = '68001'
name = 'log precision parameter'
short.name = 'prec'
initial = '2'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 0.005'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'The wrapped Cauchy likelihood'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default tan'
pdf = 'wrapped-cauchy'
status = 'disabled'
Model 'iidgamma'. Number of hyperparameters are 2.
Hyperparameter 'theta1' hyperid = '69001'
name = 'logshape'
short.name = 'shape'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '100 100'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '69002'
name = 'lograte'
short.name = 'rate'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '100 100'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = '(experimental)'
survival = 'FALSE'
discrete = 'FALSE'

```

```

link = 'default identity'
pdf = 'iidgamma'
status = 'experimental'

```

Model 'iidlogitbeta'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '70001'

```

name = 'log.a'
short.name = 'a'
initial = '1'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '70002'

```

name = 'log.b'
short.name = 'b'
initial = '1'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = '(experimental)'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga'
pdf = 'iidlogitbeta'
status = 'experimental'

```

Model 'loggammafrailty'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '71001'

```

name = 'log precision'
short.name = 'prec'
initial = '4'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = '(experimental)'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default identity'
pdf = 'loggammafrailty'
status = 'experimental'

```

Model 'logistic'. Number of hyperparameters are 1.

Hyperparameter ‘theta’ hyperid = ‘72001’

name = ‘log precision’
short.name = ‘prec’
initial = ‘1’
fixed = ‘FALSE’
prior = ‘loggamma’
param = ‘1 5e-05’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

Properties: doc = ‘The Logistic likelihood’

survival = ‘FALSE’
discrete = ‘FALSE’
link = ‘default identity’
pdf = ‘logistic’

Model ‘sn’. Number of hyperparameters are 2.

Hyperparameter ‘theta1’ hyperid = ‘74001’

name = ‘log precision’
short.name = ‘prec’
initial = ‘4’
fixed = ‘FALSE’
prior = ‘loggamma’
param = ‘1 5e-05’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

Hyperparameter ‘theta2’ hyperid = ‘74002’

name = ‘logit skew’
short.name = ‘skew’
initial = ‘0.00123456789’
fixed = ‘FALSE’
prior = ‘pc.sn’
param = ‘10’
to.theta = ‘function(x,skew.max = 0.988) log((1 + x / skew.max) / (1 -x / skew.max))’
from.theta = ‘function(x,skew.max = 0.988) skew.max * (2 * exp(x) / (1 + exp(x)) -1)’

Properties: doc = ‘The Skew-Normal likelihood’

status = ‘experimental’
survival = ‘FALSE’
discrete = ‘FALSE’
link = ‘default identity’
pdf = ‘sn’

Model ‘gev’. Number of hyperparameters are 2.

Hyperparameter ‘theta1’ hyperid = ‘76001’

name = ‘log precision’
short.name = ‘prec’
initial = ‘4’

```

fixed = 'FALSE'
prior = 'loggamma'
param = '1.5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '76002'
  name = 'tail parameter'
  short.name = 'tail'
  initial = '0'
  fixed = 'FALSE'
  prior = 'gaussian'
  param = '0.25'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Properties: doc = 'The Generalized Extreme Value likelihood'
  survival = 'FALSE'
  discrete = 'FALSE'
  link = 'default identity'
  status = 'experimental'
  pdf = 'gev'

```

Model 'lognormal'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '77101'
  name = 'log precision'
  short.name = 'prec'
  initial = '0'
  fixed = 'FALSE'
  prior = 'loggamma'
  param = '1.5e-05'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'The log-Normal likelihood'
  survival = 'FALSE'
  discrete = 'FALSE'
  link = 'default identity'
  pdf = 'lognormal'

```

Model 'lognormalsurv'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '78001'
  name = 'log precision'
  short.name = 'prec'
  initial = '0'
  fixed = 'FALSE'
  prior = 'loggamma'
  param = '1.5e-05'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'

```

Properties: **doc** = ‘The log-Normal likelihood (survival)’
survival = ‘TRUE’
discrete = ‘FALSE’
link = ‘default identity’
pdf = ‘lognormal’

Model ‘exponential’. Number of hyperparameters are 0.

Model ‘exponentialsurv’. Number of hyperparameters are 0.

Model ‘coxph’. Number of hyperparameters are 0.

Model ‘weibull’. Number of hyperparameters are 1.

Hyperparameter ‘theta’ hyperid = ‘79001’
name = ‘log alpha’
short.name = ‘alpha’
initial = ‘-2’
fixed = ‘FALSE’
prior = ‘pc.alphaw’
param = ‘5’
to.theta = ‘function(x,sc = 0.1) log(x) / sc’
from.theta = ‘function(x,sc = 0.1) exp(sc * x)’

Properties: **doc** = ‘The Weibull likelihood’
survival = ‘FALSE’
discrete = ‘FALSE’
link = ‘default log neglog quantile’
pdf = ‘weibull’

Model ‘weibullsurv’. Number of hyperparameters are 1.

Hyperparameter ‘theta’ hyperid = ‘79101’
name = ‘log alpha’
short.name = ‘alpha’
initial = ‘-2’
fixed = ‘FALSE’
prior = ‘pc.alphaw’
param = ‘5’
to.theta = ‘function(x,sc = 0.1) log(x) / sc’
from.theta = ‘function(x,sc = 0.1) exp(sc * x)’

Properties: **doc** = ‘The Weibull likelihood (survival)’
survival = ‘TRUE’
discrete = ‘FALSE’
link = ‘default log neglog quantile’
pdf = ‘weibull’

Model ‘loglogistic’. Number of hyperparameters are 1.

Hyperparameter ‘theta’ hyperid = ‘80001’
name = ‘log alpha’
short.name = ‘alpha’
initial = ‘1’
fixed = ‘FALSE’
prior = ‘loggamma’

```

param = '25 25'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'The loglogistic likelihood'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default log neglog'
pdf = 'loglogistic'

```

Model 'loglogisticsurv'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '80011'

```

name = 'log alpha'
short.name = 'alpha'
initial = '1'
fixed = 'FALSE'
prior = 'loggamma'
param = '25 25'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Properties: doc = 'The loglogistic likelihood (survival)'

```

survival = 'TRUE'
discrete = 'FALSE'
link = 'default log neglog'
pdf = 'loglogistic'

```

Model 'weibullcure'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '81001'

```

name = 'log alpha'
short.name = 'a'
initial = '0.1'
fixed = 'FALSE'
prior = 'pc.alphaw'
param = '5'
to.theta = 'function(x,sc = 0.1) log(x) / sc'
from.theta = 'function(x,sc = 0.1) exp(sc * x)'

```

Hyperparameter 'theta2' hyperid = '81002'

```

name = 'logit probability'
short.name = 'prob'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 - x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

Properties: doc = 'The Weibull-cure likelihood (survival)'

```

survival = 'TRUE'
discrete = 'FALSE'

```

```
link = 'default log neglog'
pdf = 'weibullcure'
```

Model 'stochvol'. Number of hyperparameters are 1.

```
Hyperparameter 'theta' hyperid = '82001'
name = 'log precision'
short.name = 'prec'
initial = '500'
fixed = 'TRUE'
prior = 'loggamma'
param = '1 0.005'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
```

```
Properties: doc = 'The Gaussian stochvol likelihood'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'stochvolgaussian'
```

Model 'stochvolns'. Number of hyperparameters are 2.

```
Hyperparameter 'theta1' hyperid = '82101'
name = 'logit skew'
short.name = 'skew'
initial = '0.00123456789'
fixed = 'FALSE'
prior = 'pc.sn'
param = '10'
to.theta = 'function(x, skew.max = 0.988) log((1 + x / skew.max) / (1 - x / skew.max))'
from.theta = 'function(x, skew.max = 0.988) skew.max * (2 * exp(x) / (1 + exp(x)) - 1)'
```

```
Hyperparameter 'theta2' hyperid = '82102'
name = 'log precision'
short.name = 'prec'
initial = '500'
fixed = 'TRUE'
prior = 'loggamma'
param = '1 0.005'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
```

```
Properties: doc = 'The SkewNormal stochvol likelihood'
status = 'experimental'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'stochvolns'
```

Model 'stochvolt'. Number of hyperparameters are 1.

```
Hyperparameter 'theta' hyperid = '83001'
```

```

name = 'log degrees of freedom'
short.name = 'dof'
initial = '4'
fixed = 'FALSE'
prior = 'pc.dof'
param = '15 0.5'
to.theta = 'function(x) log(x - 2)'
from.theta = 'function(x) 2 + exp(x)'

```

Properties: doc = 'The Student-t stochvol likelihood'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'stochvolt'

```

Model 'stochvolnig'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '84001'

```

name = 'skewness'
short.name = 'skew'
initial = '0'
fixed = 'FALSE'
prior = 'gaussian'
param = '0 10'
to.theta = 'function(x) x'
from.theta = 'function(x) x'

```

Hyperparameter 'theta2' hyperid = '84002'

```

name = 'shape'
short.name = 'shape'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 0.5'
to.theta = 'function(x) log(x - 1)'
from.theta = 'function(x) 1 + exp(x)'

```

Properties: doc = 'The Normal inverse Gaussian stochvol likelihood'

```

survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'stochvolnig'

```

Model 'zeroinflatedpoisson0'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '85001'

```

name = 'logit probability'
short.name = 'prob'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'

```

```

to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Zero-inflated Poisson, type 0'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'zeroinflated'

```

Model 'zeroinflatedpoisson1'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '86001'
name = 'logit probability'
short.name = 'prob'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Zero-inflated Poisson, type 1'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'zeroinflated'

```

Model 'zeroinflatedpoisson2'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '87001'
name = 'log alpha'
short.name = 'a'
initial = '0.693147180559945'
fixed = 'FALSE'
prior = 'gaussian'
param = '0.693147180559945 1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'Zero-inflated Poisson, type 2'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'zeroinflated'

```

Model 'zeroinflatedcnpoisson0'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '87101'
name = 'logit probability'
short.name = 'prob'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'

```

```

to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Zero-inflated censored Poisson, type 0'
status = 'experimental'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'zeroinflated'

```

Model 'zeroinflatedcenpoisson1'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '87201'
name = 'logit probability'
short.name = 'prob'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Zero-inflated censored Poisson, type 1'
status = 'experimental'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'zeroinflated'

```

Model 'zeroinflatedbetabinomial0'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '88001'
name = 'overdispersion'
short.name = 'rho'
initial = '0'
fixed = 'FALSE'
prior = 'gaussian'
param = '0 0.4'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta2' hyperid = '88002'
name = 'logit probability'
short.name = 'prob'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Zero-inflated Beta-Binomial, type 0'
survival = 'FALSE'

```

```

discrete = 'TRUE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'zeroinflated'

```

Model 'zeroinflatedbetabinomial1'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '89001'
  name = 'overdispersion'
  short.name = 'rho'
  initial = '0'
  fixed = 'FALSE'
  prior = 'gaussian'
  param = '0 0.4'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

```

Hyperparameter 'theta2' hyperid = '89002'
  name = 'logit probability'
  short.name = 'prob'
  initial = '-1'
  fixed = 'FALSE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

```

Properties: doc = 'Zero-inflated Beta-Binomial, type 1'
survival = 'FALSE'
discrete = 'TRUE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'zeroinflated'

```

Model 'zeroinflatedbinomial0'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '90001'
  name = 'logit probability'
  short.name = 'prob'
  initial = '-1'
  fixed = 'FALSE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

```

Properties: doc = 'Zero-inflated Binomial, type 0'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'zeroinflated'

```

Model 'zeroinflatedbinomial1'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '91001'
  name = 'logit probability'

```

```

short.name = 'prob'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 - x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Zero-inflated Binomial, type 1'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'zeroinflated'

```

Model 'zeroinflatedbinomial2'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '92001'
name = 'alpha'
short.name = 'alpha'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'Zero-inflated Binomial, type 2'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'zeroinflated'

```

Model 'zeroinflatedbinomial2'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '93001'
name = 'alpha1'
short.name = 'alpha1'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '93002'
name = 'alpha2'
short.name = 'alpha2'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x)'

```

```

from.theta = 'function(x) exp(x)'
Properties: doc = 'Zero and N inflated binomial, type 2'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'NA'
Model 'zeroninflatedbinomial3'. Number of hyperparameters are 2.
Hyperparameter 'theta1' hyperid = '93101'
  name = 'alpha0'
  short.name = 'alpha0'
  initial = '1'
  fixed = 'FALSE'
  prior = 'loggamma'
  param = '1 1'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '93102'
  name = 'alphaN'
  short.name = 'alphaN'
  initial = '1'
  fixed = 'FALSE'
  prior = 'loggamma'
  param = '1 1'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Properties: doc = 'Zero and N inflated binomial, type 3'
  status = 'experimental'
  survival = 'FALSE'
  discrete = 'FALSE'
  link = 'default logit loga cauchit probit cloglog loglog robit sn'
  pdf = 'zeroinflated'
Model 'zeroinflatedbetabinomial2'. Number of hyperparameters are 2.
Hyperparameter 'theta1' hyperid = '94001'
  name = 'log alpha'
  short.name = 'a'
  initial = '0.693147180559945'
  fixed = 'FALSE'
  prior = 'gaussian'
  param = '0.693147180559945 1'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '94002'
  name = 'beta'
  short.name = 'b'
  initial = '0'

```

```

fixed = 'FALSE'
prior = 'gaussian'
param = '0 1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'Zero inflated Beta-Binomial, type 2'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default logit loga cauchit probit cloglog loglog robit sn'
pdf = 'zeroinflated'

```

Model 'zeroinflatednbinomial0'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '95001'
  name = 'log size'
  short.name = 'size'
  initial = '2.30258509299405'
  fixed = 'FALSE'
  prior = 'pc.mgamma'
  param = '7'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '95002'
  name = 'logit probability'
  short.name = 'prob'
  initial = '-1'
  fixed = 'FALSE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Zero inflated negBinomial, type 0'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'zeroinflated'

```

Model 'zeroinflatednbinomial1'. Number of hyperparameters are 2.

```

Hyperparameter 'theta1' hyperid = '96001'
  name = 'log size'
  short.name = 'size'
  initial = '2.30258509299405'
  fixed = 'FALSE'
  prior = 'pc.mgamma'
  param = '7'
  to.theta = 'function(x) log(x)'
  from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '96002'

```

```

name = 'logit probability'
short.name = 'prob'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Properties: doc = 'Zero inflated negBinomial, type 1'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'zeroinflated'

```

Model 'zeroinflatednbinomial1strata2'. Number of hyperparameters are 11.

```

Hyperparameter 'theta1' hyperid = '97001'
name = 'log size'
short.name = 'size'
initial = '2.30258509299405'
fixed = 'FALSE'
prior = 'pc.mgamma'
param = '7'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '97002'
name = 'logit probability 1'
short.name = 'prob1'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta3' hyperid = '97003'
name = 'logit probability 2'
short.name = 'prob2'
initial = '-1'
fixed = 'FALSE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta4' hyperid = '97004'
name = 'logit probability 3'
short.name = 'prob3'
initial = '-1'

```

```

fixed = 'TRUE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta5' hyperid = '97005'
  name = 'logit probability 4'
  short.name = 'prob4'
  initial = '-1'
  fixed = 'TRUE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta6' hyperid = '97006'
  name = 'logit probability 5'
  short.name = 'prob5'
  initial = '-1'
  fixed = 'TRUE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta7' hyperid = '97007'
  name = 'logit probability 6'
  short.name = 'prob6'
  initial = '-1'
  fixed = 'TRUE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta8' hyperid = '97008'
  name = 'logit probability 7'
  short.name = 'prob7'
  initial = '-1'
  fixed = 'TRUE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta9' hyperid = '97009'
  name = 'logit probability 8'
  short.name = 'prob8'
  initial = '-1'

```

```

fixed = 'TRUE'
prior = 'gaussian'
param = '-1 0.2'
to.theta = 'function(x) log(x / (1 -x))'
from.theta = 'function(x) exp(x) / (1 + exp(x))'
Hyperparameter 'theta10' hyperid = '97010'
  name = 'logit probability 9'
  short.name = 'prob9'
  initial = '-1'
  fixed = 'TRUE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

```

Hyperparameter 'theta11' hyperid = '97011'
  name = 'logit probability 10'
  short.name = 'prob10'
  initial = '-1'
  fixed = 'TRUE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

```

Properties: doc = 'Zero inflated negBinomial, type 1, strata 2'
  status = 'experimental'
  survival = 'FALSE'
  discrete = 'FALSE'
  link = 'default log'
  pdf = 'zeroinflated'

```

Model 'zeroinflatednbinomial1strata3'. Number of hyperparameters are 11.

```

Hyperparameter 'theta1' hyperid = '98001'
  name = 'logit probability'
  short.name = 'prob'
  initial = '-1'
  fixed = 'FALSE'
  prior = 'gaussian'
  param = '-1 0.2'
  to.theta = 'function(x) log(x / (1 -x))'
  from.theta = 'function(x) exp(x) / (1 + exp(x))'

```

```

Hyperparameter 'theta2' hyperid = '98002'
  name = 'log size l'
  short.name = 'size1'
  initial = '2.30258509299405'
  fixed = 'FALSE'
  prior = 'pc.mgamma'

```

```
param = '7'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta3' hyperid = '98003'
name = 'log size 2'
short.name = 'size2'
initial = '2.30258509299405'
fixed = 'FALSE'
prior = 'pc.mgamma'
param = '7'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta4' hyperid = '98004'
name = 'log size 3'
short.name = 'size3'
initial = '2.30258509299405'
fixed = 'TRUE'
prior = 'pc.mgamma'
param = '7'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta5' hyperid = '98005'
name = 'log size 4'
short.name = 'size4'
initial = '2.30258509299405'
fixed = 'TRUE'
prior = 'pc.mgamma'
param = '7'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta6' hyperid = '98006'
name = 'log size 5'
short.name = 'size5'
initial = '2.30258509299405'
fixed = 'TRUE'
prior = 'pc.mgamma'
param = '7'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta7' hyperid = '98007'
name = 'log size 6'
short.name = 'size6'
initial = '2.30258509299405'
fixed = 'TRUE'
prior = 'pc.mgamma'
```

```

    param = '7'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta8' hyperid = '98008'
    name = 'log size 7'
    short.name = 'size7'
    initial = '2.30258509299405'
    fixed = 'TRUE'
    prior = 'pc.mgamma'
    param = '7'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta9' hyperid = '98009'
    name = 'log size 8'
    short.name = 'size8'
    initial = '2.30258509299405'
    fixed = 'TRUE'
    prior = 'pc.mgamma'
    param = '7'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta10' hyperid = '98010'
    name = 'log size 9'
    short.name = 'size9'
    initial = '2.30258509299405'
    fixed = 'TRUE'
    prior = 'pc.mgamma'
    param = '7'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta11' hyperid = '98011'
    name = 'log size 10'
    short.name = 'size10'
    initial = '2.30258509299405'
    fixed = 'TRUE'
    prior = 'pc.mgamma'
    param = '7'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Properties: doc = 'Zero inflated negBinomial, type 1, strata 3'
    status = 'experimental'
    survival = 'FALSE'
    discrete = 'FALSE'
    link = 'default log'
    pdf = 'zeroinflated'

```

Model ‘zeroinflatednbinomial2’. Number of hyperparameters are 2.

Hyperparameter ‘theta1’ hyperid = ‘99001’

name = ‘log size’
short.name = ‘size’
initial = ‘2.30258509299405’
fixed = ‘FALSE’
prior = ‘pc.mgamma’
param = ‘7’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

Hyperparameter ‘theta2’ hyperid = ‘99002’

name = ‘log alpha’
short.name = ‘a’
initial = ‘0.693147180559945’
fixed = ‘FALSE’
prior = ‘gaussian’
param = ‘2 1’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

Properties: doc = ‘Zero inflated negBinomial, type 2’

survival = ‘FALSE’
discrete = ‘FALSE’
link = ‘default log’
pdf = ‘zeroinflated’

Model ‘t’. Number of hyperparameters are 2.

Hyperparameter ‘theta1’ hyperid = ‘100001’

name = ‘log precision’
short.name = ‘prec’
initial = ‘0’
fixed = ‘FALSE’
prior = ‘loggamma’
param = ‘1 5e-05’
to.theta = ‘function(x) log(x)’
from.theta = ‘function(x) exp(x)’

Hyperparameter ‘theta2’ hyperid = ‘100002’

name = ‘log degrees of freedom’
short.name = ‘dof’
initial = ‘5’
fixed = ‘FALSE’
prior = ‘pc.dof’
param = ‘15 0.5’
to.theta = ‘function(x) log(x - 2)’
from.theta = ‘function(x) 2 + exp(x)’

Properties: doc = ‘Student-t likelihood’

survival = ‘FALSE’

```

discrete = 'FALSE'
link = 'default identity'
pdf = 'student-t'

```

Model 'tstrata'. Number of hyperparameters are 11.

Hyperparameter 'theta1' hyperid = '101001'

```

name = 'log degrees of freedom'
short.name = 'dof'
initial = '4'
fixed = 'FALSE'
prior = 'pc.dof'
param = '15 0.5'
to.theta = 'function(x) log(x - 5)'
from.theta = 'function(x) 5 + exp(x)'

```

Hyperparameter 'theta2' hyperid = '101002'

```

name = 'log precision1'
short.name = 'prec1'
initial = '2'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta3' hyperid = '101003'

```

name = 'log precision2'
short.name = 'prec2'
initial = '2'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta4' hyperid = '101004'

```

name = 'log precision3'
short.name = 'prec3'
initial = '2'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta5' hyperid = '101005'

```

name = 'log precision4'
short.name = 'prec4'
initial = '2'
fixed = 'FALSE'

```

```
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta6' hyperid = '101006'
name = 'log precision5'
short.name = 'prec5'
initial = '2'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta7' hyperid = '101007'
name = 'log precision6'
short.name = 'prec6'
initial = '2'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta8' hyperid = '101008'
name = 'log precision7'
short.name = 'prec7'
initial = '2'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta9' hyperid = '101009'
name = 'log precision8'
short.name = 'prec8'
initial = '2'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 5e-05'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta10' hyperid = '101010'
name = 'log precision9'
short.name = 'prec9'
initial = '2'
fixed = 'FALSE'
```

```

    prior = 'loggamma'
    param = '1 5e-05'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Hyperparameter 'theta11' hyperid = '101011'
    name = 'log precision10'
    short.name = 'prec10'
    initial = '2'
    fixed = 'FALSE'
    prior = 'loggamma'
    param = '1 5e-05'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Properties: doc = 'A stratified version of the Student-t likelihood'
    survival = 'FALSE'
    discrete = 'FALSE'
    link = 'default identity'
    pdf = 'tstrata'

```

Model 'nmix'. Number of hyperparameters are 15.

```

Hyperparameter 'theta1' hyperid = '101101'
    name = 'beta1'
    short.name = 'beta1'
    initial = '2.30258509299405'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 0.5'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '101102'
    name = 'beta2'
    short.name = 'beta2'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta3' hyperid = '101103'
    name = 'beta3'
    short.name = 'beta3'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'

```

```
    from.theta = 'function(x) x'
Hyperparameter 'theta4' hyperid = '101104'
  name = 'beta4'
  short.name = 'beta4'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta5' hyperid = '101105'
  name = 'beta5'
  short.name = 'beta5'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta6' hyperid = '101106'
  name = 'beta6'
  short.name = 'beta6'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta7' hyperid = '101107'
  name = 'beta7'
  short.name = 'beta7'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta8' hyperid = '101108'
  name = 'beta8'
  short.name = 'beta8'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
```

```
    from.theta = 'function(x) x'
Hyperparameter 'theta9' hyperid = '101109'
    name = 'beta9'
    short.name = 'beta9'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta10' hyperid = '101110'
    name = 'beta10'
    short.name = 'beta10'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta11' hyperid = '101111'
    name = 'beta11'
    short.name = 'beta11'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta12' hyperid = '101112'
    name = 'beta12'
    short.name = 'beta12'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta13' hyperid = '101113'
    name = 'beta13'
    short.name = 'beta13'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
```

```

from.theta = 'function(x) x'
Hyperparameter 'theta14' hyperid = '101114'
  name = 'beta14'
  short.name = 'beta14'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta15' hyperid = '101115'
  name = 'beta15'
  short.name = 'beta15'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Properties: doc = 'Binomial-Poisson mixture'
  status = 'experimental'
  survival = 'FALSE'
  discrete = 'TRUE'
  link = 'default logit loga probit'
  pdf = 'nmix'
Model 'nmixnb'. Number of hyperparameters are 16.
Hyperparameter 'theta1' hyperid = '101121'
  name = 'beta1'
  short.name = 'beta1'
  initial = '2.30258509299405'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 0.5'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta2' hyperid = '101122'
  name = 'beta2'
  short.name = 'beta2'
  initial = '0'
  fixed = 'FALSE'
  prior = 'normal'
  param = '0 1'
  to.theta = 'function(x) x'
  from.theta = 'function(x) x'
Hyperparameter 'theta3' hyperid = '101123'

```

```
name = 'beta3'
short.name = 'beta3'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta4' hyperid = '101124'
name = 'beta4'
short.name = 'beta4'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta5' hyperid = '101125'
name = 'beta5'
short.name = 'beta5'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta6' hyperid = '101126'
name = 'beta6'
short.name = 'beta6'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta7' hyperid = '101127'
name = 'beta7'
short.name = 'beta7'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta8' hyperid = '101128'
```

```
name = 'beta8'
short.name = 'beta8'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta9' hyperid = '101129'
name = 'beta9'
short.name = 'beta9'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta10' hyperid = '101130'
name = 'beta10'
short.name = 'beta10'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta11' hyperid = '101131'
name = 'beta11'
short.name = 'beta11'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta12' hyperid = '101132'
name = 'beta12'
short.name = 'beta12'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Hyperparameter 'theta13' hyperid = '101133'
```

```

    name = 'beta13'
    short.name = 'beta13'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta14' hyperid = '101134'
    name = 'beta14'
    short.name = 'beta14'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta15' hyperid = '101135'
    name = 'beta15'
    short.name = 'beta15'
    initial = '0'
    fixed = 'FALSE'
    prior = 'normal'
    param = '0 1'
    to.theta = 'function(x) x'
    from.theta = 'function(x) x'
Hyperparameter 'theta16' hyperid = '101136'
    name = 'overdispersion'
    short.name = 'overdispersion'
    initial = '0'
    fixed = 'FALSE'
    prior = 'pc.gamma'
    param = '7'
    to.theta = 'function(x) log(x)'
    from.theta = 'function(x) exp(x)'
Properties: doc = 'NegBinomial-Poisson mixture'
    status = 'experimental'
    survival = 'FALSE'
    discrete = 'TRUE'
    link = 'default logit loga probit'
    pdf = 'nmixnb'
Model 'gp'. Number of hyperparameters are 1.
Hyperparameter 'theta' hyperid = '101201'
    name = 'tail'
    short.name = 'xi'

```

```

initial = '-4'
fixed = 'FALSE'
prior = 'pc.gevtail'
param = '7 0 0.5'
to.theta = 'function(x,interval = c(REPLACE.ME.low,REPLACE.ME.high)) log(-(interval[1]
  -x) / (interval[2] -x))'
from.theta = 'function(x,interval = c(REPLACE.ME.low,REPLACE.ME.high))
  interval[1] + (interval[2] -interval[1]) * exp(x) / (1.0 + exp(x))'

```

Properties: **doc** = 'Generalized Pareto likelihood'

```

status = 'experimental'
survival = 'FALSE'
discrete = 'TRUE'
link = 'default quantile'
pdf = 'genPareto'

```

Model 'dgp'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '101201'

```

name = 'tail'
short.name = 'xi'
initial = '2'
fixed = 'FALSE'
prior = 'pc.gevtail'
param = '7 0 0.5'
to.theta = 'function(x,interval = c(REPLACE.ME.low,REPLACE.ME.high)) log(-(interval[1]
  -x) / (interval[2] -x))'
from.theta = 'function(x,interval = c(REPLACE.ME.low,REPLACE.ME.high))
  interval[1] + (interval[2] -interval[1]) * exp(x) / (1.0 + exp(x))'

```

Properties: **doc** = 'Discrete generalized Pareto likelihood'

```

status = 'experimental'
survival = 'FALSE'
discrete = 'TRUE'
link = 'default quantile'
pdf = 'dgp'

```

Model 'logperiodogram'. Number of hyperparameters are 0.

Model 'tweedie'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '102101'

```

name = 'p'
short.name = 'p'
initial = '0'
fixed = 'FALSE'
prior = 'normal'
param = '0 10'
to.theta = 'function(x,interval = c(1.0,2.0)) log(-(interval[1] -x) / (interval[2]
  -x))'
from.theta = 'function(x,interval = c(1.0,2.0)) interval[1] + (interval[2]
  -interval[1]) * exp(x) / (1.0 + exp(x))'

```

Hyperparameter 'theta2' hyperid = '102201'

```

name = 'dispersion'
short.name = 'phi'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '1 0.1'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Properties: doc = 'Tweedie distribution'
status = 'experimental'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'tweedie'

```

Model 'fmri'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '103101'

```

name = 'precision'
short.name = 'prec'
initial = '0'
fixed = 'FALSE'
prior = 'loggamma'
param = '10 10'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'

```

Hyperparameter 'theta2' hyperid = '103202'

```

name = 'dof'
short.name = 'df'
initial = '4'
fixed = 'TRUE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'

```

Properties: doc = 'fmri distribution (special nc-chi)'

```

status = 'experimental'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log'
pdf = 'fmri'

```

Model 'fmrisurv'. Number of hyperparameters are 2.

Hyperparameter 'theta1' hyperid = '104101'

```

name = 'precision'
short.name = 'prec'
initial = '0'
fixed = 'FALSE'

```

```

prior = 'loggamma'
param = '10 10'
to.theta = 'function(x) log(x)'
from.theta = 'function(x) exp(x)'
Hyperparameter 'theta2' hyperid = '104201'
name = 'dof'
short.name = 'df'
initial = '4'
fixed = 'TRUE'
prior = 'normal'
param = '0 1'
to.theta = 'function(x) x'
from.theta = 'function(x) x'
Properties: doc = 'fmri distribution (special nc-chi)
status = 'experimental'
survival = 'TRUE'
discrete = 'FALSE'
link = 'default log'
pdf = 'fmri'

```

Model 'gompertz'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '105101'
name = 'shape'
short.name = 'alpha'
initial = '-20'
fixed = 'FALSE'
prior = 'normal'
param = '-20 0.01'
to.theta = 'function(x, sc = 0.1) log(x) / sc'
from.theta = 'function(x, sc = 0.1) exp(sc * x)'
Properties: doc = 'gompertz distribution'
status = 'experimental'
survival = 'FALSE'
discrete = 'FALSE'
link = 'default log neglog'
pdf = 'gompertz'

```

Model 'gompertzsurv'. Number of hyperparameters are 1.

```

Hyperparameter 'theta' hyperid = '106101'
name = 'shape'
short.name = 'alpha'
initial = '-20'
fixed = 'FALSE'
prior = 'normal'
param = '-20 0.01'
to.theta = 'function(x, sc = 0.1) log(x) / sc'
from.theta = 'function(x, sc = 0.1) exp(sc * x)'

```

Properties: `doc` = 'gompertz distribution'
`status` = 'experimental'
`survival` = 'TRUE'
`discrete` = 'FALSE'
`link` = 'default log neglog'
`pdf` = 'gompertz'

Section 'prior'. Valid models in this section are:

Model 'normal'. Number of parameters in the prior = 2
Model 'gaussian'. Number of parameters in the prior = 2
Model 'linksnintercept'. Number of parameters in the prior = 2
Model 'wishart1d'. Number of parameters in the prior = 2
Model 'wishart2d'. Number of parameters in the prior = 4
Model 'wishart3d'. Number of parameters in the prior = 7
Model 'wishart4d'. Number of parameters in the prior = 11
Model 'wishart5d'. Number of parameters in the prior = 16
Model 'loggamma'. Number of parameters in the prior = 2
Model 'gamma'. Number of parameters in the prior = 2
Model 'minuslogsqrtruncnormal'. Number of parameters in the prior = 2
Model 'logtnormal'. Number of parameters in the prior = 2
Model 'logtgaussian'. Number of parameters in the prior = 2
Model 'flat'. Number of parameters in the prior = 0
Model 'logflat'. Number of parameters in the prior = 0
Model 'logiflat'. Number of parameters in the prior = 0
Model 'mvnorm'. Number of parameters in the prior = -1
Model 'pc.alpha'. Number of parameters in the prior = 1
Model 'pc.ar'. Number of parameters in the prior = 1
Model 'dirichlet'. Number of parameters in the prior = 1
Model 'none'. Number of parameters in the prior = 0
Model 'invalid'. Number of parameters in the prior = 0
Model 'betacorrelation'. Number of parameters in the prior = 2
Model 'logitbeta'. Number of parameters in the prior = 2
Model 'pc.prec'. Number of parameters in the prior = 2
Model 'pc.dof'. Number of parameters in the prior = 2
Model 'pc.cor0'. Number of parameters in the prior = 2
Model 'pc.cor1'. Number of parameters in the prior = 2
Model 'pc.fgnh'. Number of parameters in the prior = 2
Model 'pc.spde.GA'. Number of parameters in the prior = 4
Model 'pc.matern'. Number of parameters in the prior = 3
Model 'pc.range'. Number of parameters in the prior = 2
Model 'pc.sn'. Number of parameters in the prior = 1
Model 'pc.gamma'. Number of parameters in the prior = 1
Model 'pc.mgamma'. Number of parameters in the prior = 1
Model 'pc.gammacount'. Number of parameters in the prior = 1
Model 'pc.gevtail'. Number of parameters in the prior = 3
Model 'pc'. Number of parameters in the prior = 2

Model 'ref.ar'. Number of parameters in the prior = 0

Model 'pom'. Number of parameters in the prior = 0

Model 'jeffreystdf'. Number of parameters in the prior = 0

Model 'expression:'. Number of parameters in the prior = -1

Model 'table:'. Number of parameters in the prior = -1

Section 'wrapper'. Valid models in this section are:

Model 'joint'. Number of hyperparameters are 1.

Hyperparameter 'theta' hyperid = '102001'

name = 'log precision'

short.name = 'prec'

initial = '0'

fixed = 'TRUE'

prior = 'loggamma'

param = '1 5e-05'

to.theta = 'function(x) log(x)'

from.theta = 'function(x) exp(x)'

Properties: doc = '(experimental)'

constr = 'FALSE'

nrow.ncol = 'FALSE'

augmented = 'FALSE'

aug.factor = '1'

aug.constr = 'NULL'

n.div.by = 'NULL'

n.required = 'FALSE'

set.default.values = 'FALSE'

pdf = 'NA'

Examples

```
## How to set hyperparameters to pass as the argument 'hyper'. This
## format is compatible with the old style (using 'initial', 'fixed',
## 'prior', 'param'), but the new style using 'hyper' take precedence
## over the old style. The two styles can also be mixed. The old style
## might be removed from the code in the future...

## Only a subset need to be given
hyper = list(theta = list(initial = 2))
## The `name` can be used instead of 'theta', or 'theta1', 'theta2',...
hyper = list(precision = list(initial = 2))
hyper = list(precision = list(prior = "flat", param = numeric(0)))
hyper = list(theta2 = list(initial=3), theta1 = list(prior = "gaussian"))
## The 'short.name' can be used instead of 'name'
hyper = list(rho = list(param = c(0,1)))
```

inla.nmix.lambda.fitted

Estimate posterior distributions of fitted lambda values

Description

For use with 'nmix' and 'nmixnb' models. This function takes the information contained in an object returned by `inla()` and uses the contents to create fitted lambda values using the linear predictor for $\log(\lambda)$, the input covariate values, and samples from the posteriors of the model hyperparameters. Fitted values from the linear predictor are exponentiated, by default, before being returned.

Usage

```
inla.nmix.lambda.fitted(result, sample.size = 1000,
                        return.posterior = FALSE, scale = "exp")
```

Arguments

<code>result</code>	The output object from a call to <code>inla()</code> , where the family argument has been set to 'nmix' or 'nmixnb'. For the function to work, the call to <code>inla()</code> should also include the argument <code>control.compute=list(config = TRUE)</code> .
<code>sample.size</code>	The size of the sample from the posteriors of the model hyperparameters. This sample size ends up being the size of the estimated posterior for a fitted lambda value. Default is 1000. Larger values are recommended.
<code>return.posterior</code>	A logical value for whether or not to return the full estimated posteriors for each fitted value (TRUE), or just a summary of the posteriors (FALSE). Default is FALSE.
<code>scale</code>	A character string, where the default string, "exp", causes values from the linear predictor to be exponentiated before being returned. The string, "log", causes values to be returned on the $\log(\lambda)$ scale.

Value

<code>fitted.summary</code>	A data frame with summaries of estimated posteriors of fitted lambda values. The number of rows equals the number of rows in the data used to create the 'nmix' or 'nmixnb' model. There are six columns of summary statistics for each estimated posterior. Columns include an index, <code>mean.lambda</code> , <code>sd.lambda</code> , <code>quant025.lambda</code> , <code>median.lambda</code> , <code>quant975.lambda</code> , and <code>mode.lambda</code> .
<code>fitted.posterior</code>	A data frame containing samples that comprise the full estimated posteriors of fitted values. The number of rows equals the number of rows in the data used to create the 'nmix' or 'nmixnb' model. The number of columns equals one plus the number of samples specified by the <code>sample.size</code> argument.

Note

This function is experimental.

Author(s)

Tim Meehan <tmeehan@audubon.org>

References

See documentation for families "nmix" and "nmixmb": `inla.doc("nmix")`

Examples

```
## an example analysis of an N-mixture model using simulated data
## set parameters
n <- 75                # number of study sites
nrep.max <- 5          # number of surveys per site
b0 <- 0.5              # lambda intercept, expected abundance
b1 <- 2.0              # effect of x1 on lambda
a0 <- 1.0              # p intercept, detection probability
a2 <- 0.5              # effect of x2 on p
size <- 3.0           # size of theta
overdispersion <- 1 / size # for negative binomial distribution

## make empty vectors and matrix
x1 <- c(); x2 <- c()
lambdas <- c(); Ns <- c()
y <- matrix(NA, n, nrep.max)

## fill vectors and matrix
for(i in 1:n) {
  x1.i <- runif(1) - 0.5
  lambda <- exp(b0 + b1 * x1.i)
  N <- rnbinom(1, mu = lambda, size = size)
  x2.i <- runif(1) - 0.5
  eta <- a0 + a2 * x2.i
  p <- exp(eta) / (exp(eta) + 1)
  nr <- sample(1:nrep.max, 1)
  y[i, 1:nr] <- rbinom(nr, size = N, prob = p)
  x1 <- c(x1, x1.i); x2 <- c(x2, x2.i)
  lambdas <- c(lambdas, lambda); Ns <- c(Ns, N)
}

## bundle counts, lambda intercept, and lambda covariates
Y <- inla.mdata(y, 1, x1)

## run inla and summarize output
result <- inla(Y ~ 1 + x2,
  data = list(Y=Y, x2=x2),
  family = "nmixnb",
  control.fixed = list(mean = 0, mean.intercept = 0, prec = 0.01,
    prec.intercept = 0.01),
  control.family = list(hyper = list(theta1 = list(param = c(0, 0.01)),
    theta2 = list(param = c(0, 0.01)),
    theta3 = list(prior = "flat",
      param = numeric()))),
  control.compute=list(config = TRUE)) # important argument
summary(result)

## get and evaluate fitted values
```

```
lam.fits <- inla.nmix.lambda.fitted(result, 5000)$fitted.summary
plot(lam.fits$median.lambda, lambdas)
round(sum(lam.fits$median.lambda), 0); sum(Ns)
```

inla.nonconvex.hull *Nonconvex set extensions.*

Description

Constructs a nonconvex boundary for a point set using morphological operations.

Usage

```
inla.nonconvex.hull(
  points,
  convex = -0.15,
  concave = convex,
  resolution = 40,
  eps = NULL,
  crs = NULL
)

inla.nonconvex.hull.basic(
  points,
  convex = -0.15,
  resolution = 40,
  eps = NULL,
  crs = NULL
)
```

Arguments

points	2D point coordinates (2-column matrix). Can alternatively be a <code>SpatialPoints</code> or <code>SpatialPointsDataFrame</code> object.
convex	The desired extension radius. Also determines the smallest allowed convex curvature radius. Negative values are interpreted as fractions of the approximate initial set diameter.
concave	The desired minimal concave curvature radius. Default is <code>concave=convex</code> .
resolution	The internal computation resolution. A warning will be issued when this needs to be increased for higher accuracy, with the required resolution stated.
eps	The polygonal curve simplification tolerance used for simplifying the resulting boundary curve. See inla.simplify.curve() for details.
crs	An optional CRS or <code>inla.CRS</code> object

Details

Morphological dilation by `convex`, followed by closing by `concave`, with minimum concave curvature radius `concave`. If the dilated set has no gaps of width between

$$2convex(\sqrt{1 + 2concave/convex} - 1)$$

and 2concave , then the minimum convex curvature radius is convex. Special case `concave=0` delegates to `inla.nonconvex.hull.basic`

The implementation is based on the identity

$$\text{dilation}(a) \& \text{closing}(b) = \text{dilation}(a + b) \& \text{erosion}(b)$$

where all operations are with respect to disks with the specified radii.

Value

An `inla.mesh.segment()` object.

Note

Requires `nndistF` from the `splancs` package.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

Examples

```
if (require(splancs)) {
  loc <- matrix(runif(20), 10, 2)
  boundary <- inla.nonconvex.hull(loc, convex = 0.2)
  lines(boundary, add = FALSE)
  points(loc)
}
```

inla.option

Set and get global options for INLA

Description

Set and get global options for INLA

Usage

```
inla.setOption(...)
inla.getOption(option)
```

Arguments

... Option and value, like `option=value` or `option, value`; see the Examples

option The option to get. If `option = NULL` then `inla.getOption` then `inla.getOption` will return a named list of current values, otherwise, `option` must be one of

`inla.call`: The path to the `inla`-program.

`inla.arg`: Additional arguments to `inla.call`

`fmesher.call`: The path to the `fmesher`-program

`fmesher.arg`: Additional arguments to `fmesher.call`

`num.threads`: Character string with the number of threads to use as A:B

blas.num.threads: Number of threads to use for openblas and mklblas (see inla for details)

smtplib: Sparse matrix library to use, one of band, taucs (default) or pardiso

mkl: Use binaries built with Intel MKL? (If possible)

vecLib: This option applies to Mac only. If TRUE and mkl=FALSE, link with vecLib BLAS and LAPACK libs (if available)

vecLibPath: This option applies to Mac only. Path to vecLib-libraries. If empty, use default.

pardiso.license: The full path to the PARDISO license file

keep: Keep temporary files?

working.directory: The name of the working directory.

silent: Run the inla-program in a silent mode?

debug : Run the inla-program in a debug mode?

cygwin : The home of the Cygwin installation (default "C:/cygwin") [Remote computing for Windows only]

ssh.auth.sock: The ssh bind-address (value of \$SSH_AUTH_SOCK in the Cygwin-shell). [Remote computing for Windows only]

show.warning.graph.file : Give a warning for using the obsolete argument graph.file instead of graph

scale.model.default : The default value of argument scale.model which optionally scale intrinsic models to have generalized unit average variance

short.summary : Use a less verbose output for summary. Useful for Markdown documents.

inla.timeout : The timeout limit, in whole seconds, for calls to the inla binary. Default is 0, meaning no timeout limit. Set to a positive integer to terminate inla calls if they run to long. Fractional seconds are rounded up to the nearest integer. This feature is EXPERIMENTAL and might change at a later stage.

fmesh.timeout : The timeout limit, in whole seconds, for calls to the fmesher binary. Default is 0, meaning no timeout limit. Set to a positive integer to terminate fmesher calls that may enter infinite loops due to special geometry regularity. Fractional seconds are rounded up to the nearest integer.

inla.mode : Which mode to use in INLA? Default is "classic". Other options are (USE AT YOUR OWN RISK, THIS IS WORK IN PROGRESS!!!) options are "twostage" and "experimental".

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
## set number of threads
inla.setOption("num.threads", "4:1")
## alternative format
inla.setOption(num.threads="4:1")
## check it
inla.getOption("num.threads")
```

inla.over_sp_mesh *Check which mesh triangles are inside a polygon*

Description

Wrapper for the `sp::over()` method to find triangle centroids or vertices inside sp polygon objects

Usage

```
inla.over_sp_mesh(x, y, type = c("centroid", "vertex"), ignore.CRS = FALSE)
```

Arguments

x	geometry (typically a <code>sp::SpatialPolygons()</code> object) for the queries
y	an <code>inla.mesh()</code> object
type	the query type; either 'centroid' (default, for triangle centroids), or 'vertex' (for mesh vertices)
ignore.CRS	logical; whether to ignore the coordinate system information in x and y (default FALSE)

Value

A vector of triangle indices (when type is 'centroid') or vertex indices (when type is 'vertex')

Author(s)

Haakon Bakka, <bakka@inla.org>, and Finn Lindgren <finn.lindgren@gmail.com>

Examples

```
# Create a polygon and a mesh
obj <- sp::SpatialPolygons(list(Polygons(list(Polygon(rbind(
  c(0, 0),
  c(50, 0),
  c(50, 50),
  c(0, 50)
))),
ID = 1
)),
proj4string = inla.CRS("longlat_globe")
)
mesh <- inla.mesh.create(globe = 2, crs = inla.CRS("sphere"))

## 3 vertices found in the polygon
inla.over_sp_mesh(obj, mesh, type = "vertex")

## 3 triangles found in the polygon
inla.over_sp_mesh(obj, mesh)

## Multiple transformations can lead to slightly different results due to edge cases
## 4 triangles found in the polygon
inla.over_sp_mesh(obj, inla.spTransform(mesh, CRSobj = inla.CRS("mollweide_norm")), ignore.CRS = FALSE)
```

```
## Ignoring mismatching coordinate systems is rarely useful
## 20 triangles "found in" the polygon
inla.over_sp_mesh(obj, inla.spTransform(mesh, CRSobj = inla.CRS("mollweide_norm")), ignore.CRS = TRUE)
```

inla.pardiso *PARDISO support in R-INLA*

Description

Describe and check the PARDISO support in R-INLA

Usage

```
inla.pardiso()
inla.pardiso.check()
```

Details

inla.pardiso() describes the PARDISO support in R-INLA, how to get the license key and enable it in the R-INLA package. inla.pardiso.check() check if the PARDISO support is working.

Author(s)

Havard Rue <hrue@r-inla.org>

inla.priors.used *Print priors used*

Description

Print the priors used for the hyperparameters

Usage

```
inla.priors.used(result, digits=6L)
```

Arguments

result	An inla-object, typically the output from an inla()-call
digits	The digits argument to the function format()

Details

This function provides a more human-friendly output of result\$all.hyper of all the priors used for the hyperparameters. Since not all information about the model is encoded in this object, more hyperparameters than actually used, may be printed. In particular, group.theta1 is printed even though the argument group in f() is not used. Similarly for spde-models, but the user should know that, for example, only the two first ones are actually used. Hopefully, this issue will be fixed in the future.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
r = inla(y ~ 1 + x, data = data.frame(y = 1:10, x = rep(1:5, 2)))
inla.priors.used(r)
```

inla.prune

Prune the INLA-package

Description

Prune the INLA-package by deleting binary files not supported by the running OS

Usage

```
inla.prune(ask = TRUE)
```

Arguments

ask Logical. If TRUE, then ask for user confirmation before deleting. If FALSE, then delete without user confirmation.

Value

No value is returned.

Author(s)

Havard Rue <hrue@r-inla.org>

inla.qstat

Control and view a remote inla-queue

Description

Control and view a remote inla-queue of submitted jobs

Usage

```
inla.qget(id, remove = TRUE)
inla.qdel(id)
inla.qstat(id)
inla.qlog(id)
inla.qnuke()
## S3 method for class 'inla.q'
summary(object,...)
## S3 method for class 'inla.q'
print(x,...)
```

Arguments

id	The job-id which is the output from inla when the job is submitted, the job-number or job-name. For inla.qstat, id is optional and if omitted all the jobs will be listed.
remove	Logical If FALSE, leave the job on the server after retrieval, otherwise remove it (default).
x	An inla.q-object which is the output from inla.qstat
object	An inla.q-object which is the output from inla.qstat
...	other arguments.

Details

inla.qstat show job(s) on the server, inla.qget fetch the results (and by default remove the files on the server), inla.qdel removes a job on the server and inla.qnuke remove all jobs on the server. inla.qlog fetches the logfile only.

The recommended procedure is to use `r=inla(..., inla.call="submit")` and then do `r=inla.qget(r)` at a later stage. If the job is not finished, then `r` will not be overwritten and this step can be repeated. The reason for this procedure, is that some information usually stored in the result object does not go through the remote server, hence have to be appended to the results that are retrieved from the server. Hence doing `r=inla(..., inla.call="submit")` and then later retrieve it using `r=inla.qget(1)`, say, then `r` does not contain all the usual information. All the main results are there, but administrative information which is required to call `inla.hyperpar` or `inla.rerun` are not there.

Value

inla.qstat returns an inla.q-object with information about current jobs.

Author(s)

Havard Rue

See Also

[inla](#)

Examples

```
## Not run:
r = inla(y~1, data = data.frame(y=rnorm(10)), inla.call="submit")
inla.qstat()
r = inla.qget(r, remove=FALSE)
inla.qdel(1)
inla.qnuke()

## End(Not run)
```

inla.reorderings	<i>Reorderings methods for sparse matrices</i>
------------------	--

Description

Provide the names of all implemented reordering schemes

Usage

```
inla.reorderings()
```

Arguments

None

Value

The names of all available reorderings

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
inla.reorderings()
```

inla.rerun	<i>Rerun an analysis</i>
------------	--------------------------

Description

Rerun [inla](#) on an inla-object (output from `link{inla}`)

Usage

```
inla.rerun(object, plain=FALSE)
```

Arguments

object	An inla-object, ie the output from an inla-call
plain	Logical. If FALSE (default), then make changes in object to improve the performance

Value

This function will take the result in object, and rerun inla again. If plain is FALSE, start the optimization from the mode in object so that we can obtain an improvement the mode for the hyperparameters. Otherwise, start from the same configuration as for object. The returned value is an inla-object.

See Also[inla](#)**Examples**

```
r = inla(y ~ 1, data = data.frame(y=1:10))
r = inla.rerun(r)
```

inla.row.kron	<i>Row-wise Kronecker products</i>
---------------	------------------------------------

Description

Takes two Matrices and computes the row-wise Kronecker product. Optionally applies row-wise weights and/or applies an additional 0/1 row-wise Kronecker matrix product, as needed by [inla.spde.make.A\(\)](#).

Usage

```
inla.row.kron(M1, M2, repl = NULL, n.repl = NULL, weights = NULL)
```

Arguments

M1	A matrix that can be transformed into a sparse Matrix.
M2	A matrix that can be transformed into a sparse Matrix.
repl	An optional index vector. For each entry, specifies which replicate the row belongs to, in the sense used in inla.spde.make.A() .
n.repl	The maximum replicate index, in the sense used in inla.spde.make.A() .
weights	Optional scaling weights to be applied row-wise to the resulting matrix.

Value

A sparseMatrix object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also[inla.spde.make.A\(\)](#)

inla.sample	<i>Generate samples, and functions thereof, from an approximated posterior of a fitted model</i>
-------------	--

Description

This function generate samples, and functions of those, from an approximated posterior of a fitted model (an inla-object)

Usage

```
inla.posterior.sample(n = 1L, result, selection = list(),
  intern = FALSE,
  use.improved.mean = TRUE, skew.corr = TRUE,
  add.names = TRUE, seed = 0L, num.threads = NULL,
  parallel.configs = TRUE, verbose=FALSE)
inla.posterior.sample.eval(fun, samples, return.matrix = TRUE, ...)
```

Arguments

n	Number of samples.
result	The inla-object, ie the output from an inla-call. The inla-object must be created with <code>control.compute=list(config=TRUE)</code> .
selection	Select what part of the sample to return. By default, the whole sample is returned. <code>selection</code> is a named list with the name of the components of the sample, and what indices of them to return. Names include APredictor, Predictor, (Intercept), and otherwise names in the formula. The values of the list, is interpreted as indices. If they are negative, they are interpreted as 'not', a zero is interpreted as 'all', and positive indices are interpreted as 'only'. The names of elements of each samples refer to the indices in the full sample.
intern	Logical. If TRUE then produce samples in the internal scale for the hyperparameter, if FALSE then produce samples in the user-scale. (For example log-precision (intern) and precision (user-scale))
use.improved.mean	Logical. If TRUE then use the marginal mean values when constructing samples. If FALSE then use the mean in the Gaussian approximations.
skew.corr	Logical. If TRUE then correct samples for skewness, if FALSE, do not correct samples for skewness (ie use the Gaussian).
add.names	Logical. If TRUE then add name for each elements of each sample. If FALSE, only add name for the first sample. (This save space.)
seed	See the same argument in <code>?inla.qsample</code> for further information. In order to produce reproducible results, you ALSO need to make sure the RNG in R is in the same state, see example below. When seed is non-zero, <code>num.threads</code> is forced to "1:1" and <code>parallel.configs</code> is set to FALSE, since parallel sampling would not produce a reproducible sequence of pseudo-random numbers.
num.threads	The number of threads to use in the format 'A:B' defining the number threads in the outer (A) and inner (B) layer for nested parallelism. A '0' will be replaced intelligently. <code>seed!=0</code> requires serial computations.

<code>parallel.configs</code>	Logical. If TRUE and not on Windows, then try to run each configuration in parallel (not Windows) using <code>A</code> threads (see <code>num.threads</code>), where each of them is using <code>B:0</code> threads.
<code>verbose</code>	Logical. Run in verbose mode or not.
<code>fun</code>	The function to evaluate for each sample. Upon entry, the variable names defined in the model are defined as the value of the sample. The list of names are defined in <code>result\$misc\$configs\$contents</code> where <code>result</code> is an <code>inla</code> -object. This includes predefined names for for the linear predictor (<code>Predictor</code> and <code>APredictor</code>), and the intercept (<code>(Intercept)</code> or <code>Intercept</code>). The hyperparameters are defined as <code>theta</code> , no matter if they are in the internal scale or not. The function <code>fun</code> can also return a vector. To simplify usage, <code>fun</code> can also be a vector character's. In this case <code>fun</code> it is interpreted as (strict) variable names, and a function is created that return these variables: if argument <code>fun</code> equals <code>c("Intercept", "a[1:2]")</code> , then this is equivalent to pass <code>function() return(c(get('Intercept'), get('a[1:2]')))</code> .
<code>samples</code>	<code>samples</code> is the output from <code>inla.posterior.sample()</code>
<code>return.matrix</code>	Logical. If TRUE, then return the samples of <code>fun</code> as matrix, otherwise, as a list.
<code>...</code>	Additional arguments to <code>fun</code>

Details

The hyperparameters are sampled from the configurations used to do the numerical integration, hence if you want a higher resolution, you need to to change the `int.strategy` variable and `friends`. The latent field is sampled from the Gaussian approximation conditioned on the hyperparameters, but with a correction for the mean (default), and optional (and by default) corrected for the estimated skewness.

The `log.density` report is only correct when there is no constraints. With constraints, it correct the Gaussian part of the sample for the constraints.

After the sample is (optional) skewness corrected, the `log.density` is is not exact for correcting for constraints, but the error is very small in most cases.

Value

`inla.posterior.sample` returns a list of the samples, where each sample is a list with names `hyperpar` and `latent`, and with their marginal densities in `logdens$hyperpar` and `logdens$latent` and the joint density is in `logdens$joint`. `inla.posterior.sample.eval` return a list or a matrix of `fun` applied to each sample.

Author(s)

Harvard Rue <hrue@r-inla.org> and Cristian Chiuchiolu <cristian.chiuchiolu@kaust.edu.sa>

Examples

```
r = inla(y ~ 1 ,data = data.frame(y=rnorm(1)), control.compute = list(config=TRUE))
samples = inla.posterior.sample(2,r)

## reproducible results:
inla.seed = as.integer(runif(1)*.Machine$integer.max)
set.seed(12345)
x = inla.posterior.sample(10, r, seed = inla.seed, num.threads="1:1")
```

```

set.seed(12345)
xx = inla.posterior.sample(10, r, seed = inla.seed, num.threads="1.1")
all.equal(x, xx)

set.seed(1234)
n = 25
xx = rnorm(n)
yy = rev(xx)
z = runif(n)
y = rnorm(n)
r = inla(y ~ 1 + z + f(xx) + f(yy, copy="xx"),
        data = data.frame(y, z, xx, yy),
        control.compute = list(config=TRUE),
        family = "gaussian")
r.samples = inla.posterior.sample(10, r)

fun = function(...) {
  mean(xx) - mean(yy)
}
f1 = inla.posterior.sample.eval(fun, r.samples)

fun = function(...) {
  c(exp(Intercept), exp(Intercept + z))
}
f2 = inla.posterior.sample.eval(fun, r.samples)

fun = function(...) {
  return (theta[1]/(theta[1] + theta[2]))
}
f3 = inla.posterior.sample.eval(fun, r.samples)

## Predicting nz new observations, and
## comparing the estimated one with the true one
set.seed(1234)
n = 100
alpha = beta = s = 1
z = rnorm(n)
y = alpha + beta * z + rnorm(n, sd = s)
r = inla(y ~ 1 + z,
        data = data.frame(y, z),
        control.compute = list(config=TRUE),
        family = "gaussian")
r.samples = inla.posterior.sample(10^3, r)

## just return samples of the intercept
intercepts = inla.posterior.sample.eval("Intercept", r.samples)

nz = 3
znew = rnorm(nz)
fun = function(zz = NA) {
  ## theta[1] is the precision
  return (Intercept + z * zz +
          rnorm(length(zz), sd = sqrt(1/theta[1])))
}
par(mfrow=c(1, nz))
f1 = inla.posterior.sample.eval(fun, r.samples, zz = znew)
for(i in 1:nz) {

```

```

    hist(f1[i, ], n = 100, prob = TRUE)
    m = alpha + beta * znew[i]
    xx = seq(m-4*s, m+4*s, by = s/100)
    lines(xx, dnorm(xx, mean=m, sd = s), lwd=2)
  }

##
## Be aware that using non-clean variable names might be a little tricky
##
n <- 100
X <- matrix(rnorm(n^2), n, 2)
x <- X[, 1]
xx <- X[, 2]
xxx <- x*xx

y <- 1 + 2*x + 3*xx + 4*xxx + rnorm(n, sd = 0.01)

r <- inla(y ~ X[, 1]*X[, 2],
          data = list(y = y, X = X),
          control.compute = list(config = TRUE))
print(round(dig = 4, r$summary.fixed[, "mean"]))

sam <- inla.posterior.sample(100, r)
sam.extract <- inla.posterior.sample.eval(
  (function(...) {
    beta.1 <- get("X[, 1]")
    beta.2 <- get("X[, 2]")
    beta.12 <- get("X[, 1]:X[, 2]")
    return(c(Intercept, beta.1, beta.2, beta.12))
  }), sam)
print(round(dig = 4, rowMeans(sam.extract)))

## a simpler form can also be used here, and in the examples below
sam.extract <- inla.posterior.sample.eval(
  c("Intercept", "X[, 1]", "X[, 2]", "X[, 1]:X[, 2]"), sam)
print(round(dig = 4, rowMeans(sam.extract)))

r <- inla(y ~ x + xx + xxx,
          data = list(y = y, x = x, xx = xx, xxx = xxx),
          control.compute = list(config = TRUE))

sam <- inla.posterior.sample(100, r)
sam.extract <- inla.posterior.sample.eval(
  (function(...) {
    return(c(Intercept, x, xx, xxx))
  }), sam)
print(round(dig = 4, rowMeans(sam.extract)))

sam.extract <- inla.posterior.sample.eval(c("Intercept", "x", "xx", "xxx"), sam)
print(round(dig = 4, rowMeans(sam.extract)))

r <- inla(y ~ x*xx,
          data = list(y = y, x = x, xx = xx),
          control.compute = list(config = TRUE))

sam <- inla.posterior.sample(100, r)
sam.extract <- inla.posterior.sample.eval(

```

```
(function(...) {
  return(c(Intercept, x, xx, get("x:xx")))
}), sam)
print(round(dig = 4, rowMeans(sam.extract)))

sam.extract <- inla.posterior.sample.eval(c("Intercept", "x", "xx", "x:xx"), sam)
print(round(dig = 4, rowMeans(sam.extract)))
```

inla.simplify.curve *Recursive curve simplification.*

Description

Attempts to simplify a polygonal curve by joining nearly colinear segments.

Usage

```
inla.simplify.curve(loc, idx, eps)
```

Arguments

loc	Coordinate matrix.
idx	Index vector into loc specifying a polygonal curve.
eps	Straightness tolerance.

Details

Uses a variation of the binary splitting Ramer-Douglas-Peucker algorithm, with a width eps ellipse instead of a rectangle, motivated by prediction ellipse for Brownian bridge.

Value

An index vector into loc specifying the simplified polygonal curve.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

Examples

```
theta <- seq(0, 2 * pi, length = 1000)
loc <- cbind(cos(theta), sin(theta))
idx <- inla.simplify.curve(loc = loc, idx = 1:nrow(loc), eps = 0.01)
print(c(nrow(loc), length(idx)))
plot(loc, type = "l")
lines(loc[idx, ], col = "red")
```

inla.spde.make.A *Observation/prediction matrices for mesh models.*

Description

Constructs observation/prediction weight matrices for models based on [inla.mesh\(\)](#) and [inla.mesh.1d\(\)](#) objects.

Usage

```
inla.spde.make.A(
  mesh = NULL,
  loc = NULL,
  index = NULL,
  group = NULL,
  repl = 1L,
  n.spde = NULL,
  n.group = NULL,
  n.repl = NULL,
  group.mesh = NULL,
  weights = NULL,
  A.loc = NULL,
  A.group = NULL,
  group.index = NULL,
  block = NULL,
  n.block = NULL,
  block.rescale = c("none", "count", "weights", "sum"),
  ...
)
```

Arguments

mesh	An inla.mesh() or inla.mesh.1d() object specifying a function basis on a mesh domain. Alternatively, an inla.spde object that includes a mesh (e.g. from inla.spde2.matern()).
loc	Observation/prediction coordinates. mesh and loc defines a matrix A.loc of mapping weights between basis function weights and field values. If loc is NULL, A.loc is defined as Diagonal(n.spde, 1).
index	For each observation/prediction value, an index into loc. Default is seq_len(nrow(A.loc)).
group	For each observation/prediction value, an index into the group model.
repl	For each observation/prediction value, the replicate index.
n.spde	The number of basis functions in the mesh model. (Note: may be different than the number of mesh vertices/nodes/knots.)
n.group	The size of the group model.
n.repl	The total number of replicates.
group.mesh	An optional inla.mesh.1d() object for the group model.
weights	Optional scaling weights to be applied row-wise to the resulting matrix.

<code>A.loc</code>	Optional precomputed observation/prediction matrix. <code>A.loc</code> can be specified instead of <code>mesh+loc</code> , optionally with <code>index</code> supplied.
<code>A.group</code>	Optional precomputed observation/prediction matrix for the group model. <code>A.group</code> can be specified instead of <code>group</code> and/or <code>group.mesh</code> , optionally with <code>group.index</code> supplied.
<code>group.index</code>	For each observation/prediction value, an index into the rows of <code>A.group</code> .
<code>block</code>	Optional indices specifying block groupings: Entries with the same <code>block</code> value are joined into a single row in the resulting matrix, and the <code>block</code> values are the row indices. This is intended for construction of approximate integration schemes for regional data problems. See inla.spde.make.block.A() for details.
<code>n.block</code>	The number of blocks.
<code>block.rescale</code>	Specifies what scaling method should be used when joining entries as grouped by a block specification. See inla.spde.make.block.A() for details.
<code>...</code>	Additional parameters. Currently unused.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde.make.index\(\)](#)

Examples

```
loc <- matrix(runif(10000 * 2) * 1000, 10000, 2)
mesh <- inla.mesh.2d(
  loc = loc,
  cutoff = 50,
  max.edge = c(50, 500)
)
A <- inla.spde.make.A(mesh, loc = loc)
```

`inla.spde.make.block.A`

Observation matrices for mesh models.

Description

Constructs observation/prediction weight matrices for numerical integration schemes for regional data problems. Primarily intended for internal use by [inla.spde.make.A\(\)](#).

Usage

```
inla.spde.make.block.A(
  A,
  block,
  n.block = max(block),
  weights = NULL,
  rescale = c("none", "count", "weights", "sum")
)
```

Arguments

A	A precomputed observation/prediction matrix for locations that are to be joined.
block	Indices specifying block groupings: Entries with the same block value are joined into a single row in the resulting matrix, and the block values are the row indices.
n.block	The number of blocks.
weights	Optional scaling weights to be applied row-wise to the input A matrix.
rescale	Specifies what scaling method should be used when joining the rows of the A matrix as grouped by the block specification. <ul style="list-style-type: none"> 'none': Straight sum, no rescaling. 'count': Divide by the number of entries in the block. 'weights': Divide by the sum of the weight values within each block. 'sum': Divide by the resulting row sums.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde.make.A\(\)](#)

inla.spde.make.index *SPDE model index vector generation*

Description

Generates a list of named index vectors for an SPDE model.

Usage

```
inla.spde.make.index(name, n.spde, n.group = 1, n.repl = 1, ...)
```

Arguments

name	A character string with the base name of the effect.
n.spde	The size of the model, typically from <code>spde\$n.spde</code> .
n.group	The size of the group model.
n.repl	The number of model replicates.
...	Additional parameters. Currently unused.

Value

A list of named index vectors.

name	Indices into the vector of latent variables
name.group	'group' indices
name.repl	Indices for replicates

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde.make.A\(\)](#), [inla.spde2.result\(\)](#)

Examples

```
loc <- matrix(runif(100 * 2), 100, 2)
mesh <- inla.mesh.create.helper(points.domain = loc, max.edge = c(0.1, 0.5))
spde <- inla.spde2.matern(mesh)
index <- inla.spde.make.index("spatial", spde$n.spde, n.repl = 2)
spatial.A <- inla.spde.make.A(mesh, loc,
  index = rep(1:nrow(loc), 2),
  repl = rep(1:2, each = nrow(loc))
)
y <- 10 + rnorm(100 * 2)
stack <- inla.stack(
  data = list(y = y),
  A = list(spatial.A),
  effects = list(c(index, list(intercept = 1))),
  tag = "tag"
)
data <- inla.stack.data(stack, spde = spde)
formula <- y ~ -1 + intercept + f(spatial,
  model = spde,
  replicate = spatial.repl
)
result <- inla(formula,
  family = "gaussian", data = data,
  control.predictor = list(A = inla.stack.A(stack))
)
spde.result <- inla.spde2.result(result, "spatial", spde)
```

inla.spde.models

List SPDE models supported by inla.spde objects

Description

List SPDE models supported by inla.spde objects

Usage

```
inla.spde.models(function.names = FALSE)
```

```
inla.spde1.models()
```

```
inla.spde2.models()
```

Arguments

`function.names` If FALSE, return list model name lists. If TRUE, return list of model object constructor function names.

Details

Returns a list of available SPDE model type name lists, one for each inla.spde model class (currently `inla.spde1()` and `inla.spde2()`).

Value

List of available SPDE model type name lists.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

Examples

```
## Display help for each supported inla.spde2 model:
for (model in inla.spde2.models()) {
  print(help(paste("inla.spde2.", model, sep = "")))
}

## Display help for each supported inla.spde* model:
models <- inla.spde.models()
for (type in names(models)) {
  for (model in models[[type]]) {
    print(help(paste("inla.", type, ".", model, sep = "")))
  }
}

## Display help for each supported inla.spde* model (equivalent to above):
for (model in inla.spde.models(function.names = TRUE)) {
  print(help(model))
}
```

<code>inla.spde.precision</code>	<i>Precision matrices for SPDE models Calculates the precision matrix for given parameter values based on an inla.spde model object.</i>
----------------------------------	--

Description

Precision matrices for SPDE models

Calculates the precision matrix for given parameter values based on an `inla.spde` model object.

Usage

```
inla.spde.precision(...)

inla.spde1.precision(spde, ...)

## S3 method for class 'inla.spde1'
inla.spde.precision(spde, ...)

inla.spde2.precision(
  spde,
```

```

    theta = NULL,
    phi0 = inla.spde2.theta2phi0(spde, theta),
    phi1 = inla.spde2.theta2phi1(spde, theta),
    phi2 = inla.spde2.theta2phi2(spde, theta),
    ...
  )

## S3 method for class 'inla.spde2'
inla.spde.precision(
  spde,
  theta = NULL,
  phi0 = inla.spde2.theta2phi0(spde, theta),
  phi1 = inla.spde2.theta2phi1(spde, theta),
  phi2 = inla.spde2.theta2phi2(spde, theta),
  ...
)

```

Arguments

...	Additional parameters passed on to other methods.
spde	An inla.spde object.
theta	The parameter vector.
phi0	Internal parameter for a generic model. Expert option only.
phi1	Internal parameter for a generic model. Expert option only.
phi2	Internal parameter for a generic model. Expert option only.

Value

A sparse precision matrix.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde.models\(\)](#), [inla.spde2.generic\(\)](#), [inla.spde2.theta2phi0\(\)](#), [inla.spde2.theta2phi1\(\)](#), [inla.spde2.theta2phi2\(\)](#)

inla.spde.result

SPDE result extraction from INLA estimation results

Description

Extract field and parameter values and distributions for an inla.spde SPDE effect from an inla result object.

Usage

```

inla.spde.result(...)

inla.spde1.result(inla, name, spde, do.transform = TRUE, ...)

## S3 method for class 'inla.spde1'
inla.spde.result(inla, name, spde, do.transform = TRUE, ...)

inla.spde2.result(inla, name, spde, do.transform = TRUE, ...)

## S3 method for class 'inla.spde2'
inla.spde.result(inla, name, spde, do.transform = TRUE, ...)

```

Arguments

...	Further arguments passed to and from other methods.
inla	An inla object obtained from a call to <code>inla()</code>
name	A character string with the name of the SPDE effect in the inla formula.
spde	The <code>inla.spde</code> object used for the effect in the inla formula. (Note: this could have been stored in the inla output, but isn't.) Usually the result of a call to <code>inla.spde2.matern()</code> .
do.transform	If TRUE, also calculate marginals transformed to user-scale. Setting to FALSE is useful for large non-stationary models, as transforming many marginal densities is time-consuming.

Value

For `inla.spde2` models, a list, where the nominal range and variance are defined as the values that would have been obtained with a stationary model and no boundary effects:

<code>marginals.kappa</code>	Marginal densities for kappa
<code>marginals.log.kappa</code>	Marginal densities for log(kappa)
<code>marginals.log.range.nominal</code>	Marginal densities for log(range)
<code>marginals.log.tau</code>	Marginal densities for log(tau)
<code>marginals.log.variance.nominal</code>	Marginal densities for log(variance)
<code>marginals.range.nominal</code>	Marginal densities for range
<code>marginals.tau</code>	Marginal densities for tau
<code>marginals.theta</code>	Marginal densities for the theta parameters
<code>marginals.values</code>	Marginal densities for the field values
<code>marginals.variance.nominal</code>	Marginal densities for variance

```

summary.hyperpar
    The SPDE related part of the inla hyperpar output summary
summary.log.kappa
    Summary statistics for log(kappa)
summary.log.range.nominal
    Summary statistics for log(range)
summary.log.tau
    Summary statistics for log(tau)
summary.log.variance.nominal
    Summary statistics for log(kappa)
summary.theta
    Summary statistics for the theta parameters
summary.values
    Summary statistics for the field values

```

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde.models\(\)](#), [inla.spde2.matern\(\)](#)

Examples

```

loc <- matrix(runif(100 * 2), 100, 2)
mesh <- inla.mesh.create.helper(points.domain = loc, max.edge = c(0.1, 0.5))
spde <- inla.spde2.matern(mesh)
index <- inla.spde.make.index("spatial", mesh$n, n.repl = 2)
spatial.A <- inla.spde.make.A(mesh, loc,
  index = rep(1:nrow(loc), 2),
  repl = rep(1:2, each = nrow(loc))
)
## Toy example with no spatial correlation (range=zero)
y <- 10 + rnorm(100 * 2)
stack <- inla.stack(
  data = list(y = y),
  A = list(spatial.A),
  effects = list(c(index, list(intercept = 1))),
  tag = "tag"
)
data <- inla.stack.data(stack, spde = spde)
formula <- y ~ -1 + intercept + f(spatial,
  model = spde,
  replicate = spatial.repl
)
result <- inla(formula,
  family = "gaussian", data = data,
  control.predictor = list(A = inla.stack.A(stack))
)
spde.result <- inla.spde.result(result, "spatial", spde)
plot(spde.result$marginals.range.nominal[[1]], type = "l")

```

inla.spde.sample *Sample from SPDE models*

Description

Old methods fo sampling from a SPDE model. For new code, use [inla.spde.precision\(\)](#) and [inla.qsample\(\)](#) instead.

Usage

```
inla.spde.sample(...)

## Default S3 method:
inla.spde.sample(precision, seed = NULL, ...)

## S3 method for class 'inla.spde'
inla.spde.sample(spde, seed = NULL, ...)
```

Arguments

...	Parameters passed on to other methods.
precision	A precision matrix.
seed	The seed for the pseudo-random generator.
spde	An inla.spde object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde.precision\(\)](#), [inla.qsample\(\)](#)

inla.spde1.create *Old SPDE model objects for INLA*

Description

Create an inla.spde1 model object.

Usage

```
inla.spde1.create(
  mesh,
  model = c("matern", "imatern", "matern.osc"),
  param = NULL,
  ...
)
```

```
inla.spde1.matern(mesh, ...)
```

```
inla.spde1.imatern(mesh, ...)
```

```
inla.spde1.matern.osc(mesh, ...)
```

Arguments

mesh	The mesh to build the model on, as an inla.mesh() object.
model	The name of the model.
param	Model specific parameters.
...	Additional parameters passed on to other methods.

Details

Note: This is an old spde object format retained for backwards compatibility. Please use [inla.spde2\(\)](#) models for new code.

This method constructs an object for SPDE models. Currently implemented:

model="matern"

$$(\kappa^2(u) - \Delta)^{\alpha/2}(\tau(u))$$

$$x(u) = W(u)$$

param:

- alpha = 1 or 2
- basis.T = Matrix of basis functions for $\log \tau(u)$
- basis.K = Matrix of basis functions for $\log \kappa^2(u)$

model="imatern"

$$(-\Delta)^{\alpha/2}(\tau(u))$$

$$x(u) = W(u)$$

param:

- alpha = 1 or 2
- basis.T = Matrix of basis functions for $\log \tau(u)$

Value

An `inla.spde1` object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde2.matern\(\)](#), [inla.mesh.2d\(\)](#), [inla.mesh.basis\(\)](#)

Examples

```

n <- 100
field.fcn <- function(loc) (10 * cos(2 * pi * 2 * (loc[, 1] + loc[, 2])))
loc <- matrix(runif(n * 2), n, 2)
## One field, 2 observations per location
idx.y <- rep(1:n, 2)
y <- field.fcn(loc[idx.y, ]) + rnorm(length(idx.y))

mesh <- inla.mesh.create(loc, refine = list(max.edge = 0.05))
spde <- inla.spde1.create(mesh, model = "matern")
data <- list(y = y, field = mesh$idx$loc[idx.y])
formula <- y ~ -1 + f(field, model = spde)
result <- inla(formula, data = data, family = "normal")

## Plot the mesh structure:
plot(mesh)

if (require(rgl)) {
  ## Plot the posterior mean:
  plot(mesh,
        rgl = TRUE,
        result$summary.random$field[, "mean"],
        color.palette = colorRampPalette(c("blue", "green", "red")))
  )
  ## Plot residual field:
  plot(mesh,
        rgl = TRUE,
        result$summary.random$field[, "mean"] - field.fcn(mesh$loc),
        color.palette = colorRampPalette(c("blue", "green", "red")))
  )
}

```

inla.spde2.generic *Generic spde2 model creation.*

Description

Creates an inla.spde2 object describing the internal structure of an 'spde2' model.

Usage

```

inla.spde2.generic(
  M0,
  M1,
  M2,
  B0,
  B1,
  B2,
  theta.mu,
  theta.Q,
  transform = c("logit", "log", "identity"),

```

```

theta.initial = theta.mu,
fixed = rep(FALSE, length(theta.mu)),
theta.fixed = theta.initial[fixed],
BLC = cbind(0, diag(nrow = length(theta.mu))),
...
)

```

Arguments

M0	The symmetric M0 matrix.
M1	The square M1 matrix.
M2	The symmetric M2 matrix.
B0	Basis definition matrix for ϕ_0 .
B1	Basis definition matrix for ϕ_2 .
B2	Basis definition matrix for ϕ_2 .
theta.mu	Prior expectation for the θ vector
theta.Q	Prior precision for the θ vector
transform	Transformation link for ϕ_2 . Valid settings are "logit", "log", and "identity"
theta.initial	Initial value for the θ vector. Default theta.mu
fixed	Logical vector. For every TRUE value, treat the corresponding theta value as known.
theta.fixed	Vector holding the values of fixed theta values. Default =theta.initial[fixed]
BLC	Basis definition matrix for linear combinations of theta.
...	Additional parameters, currently unused.
spde	An inla.spde2 object.
theta	parameter values to be mapped.

Value

For `inla.spde2.generic`, an `inla.spde2()` object.

For `inla.spde2.theta2phi0/1/2`, a vector of ϕ values.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

`inla.spde2.models()`, `inla.spde2.matern()`

inla.spde2.matern *Matern SPDE model object for INLA*

Description

Create an `inla.spde2` model object for a Matern model. Use `inla.spde2.pcmatern` instead for a PC prior for the parameters.

Usage

```
inla.spde2.matern(
  mesh,
  alpha = 2,
  param = NULL,
  constr = FALSE,
  extraconstr.int = NULL,
  extraconstr = NULL,
  fractional.method = c("parsimonious", "null"),
  B.tau = matrix(c(0, 1, 0), 1, 3),
  B.kappa = matrix(c(0, 0, 1), 1, 3),
  prior.variance.nominal = 1,
  prior.range.nominal = NULL,
  prior.tau = NULL,
  prior.kappa = NULL,
  theta.prior.mean = NULL,
  theta.prior.prec = 0.1,
  n.iid.group = 1,
  ...
)
```

```
inla.spde2.theta2phi0(spde, theta)
```

```
inla.spde2.theta2phi1(spde, theta)
```

```
inla.spde2.theta2phi2(spde, theta)
```

Arguments

<code>mesh</code>	The mesh to build the model on, as an <code>inla.mesh()</code> or <code>inla.mesh.1d()</code> object.
<code>alpha</code>	Fractional operator order, $0 < \alpha \leq 2$ supported. ($\nu = \alpha - d/2$)
<code>param</code>	Parameter, e.g. generated by <code>param2.matern.orig</code>
<code>constr</code>	If TRUE, apply an integrate-to-zero constraint. Default FALSE.
<code>extraconstr.int</code>	Field integral constraints.
<code>extraconstr</code>	Direct linear combination constraints on the basis weights.
<code>fractional.method</code>	Specifies the approximation method to use for fractional (non-integer) alpha values. 'parsimonious' gives an overall approximate minimal covariance error, 'null' uses approximates low-order properties.

B.tau	Matrix with specification of log-linear model for τ .
B.kappa	Matrix with specification of log-linear model for κ .
prior.variance.nominal	Nominal prior mean for the field variance
prior.range.nominal	Nominal prior mean for the spatial range
prior.tau	Prior mean for tau (overrides prior.variance.nominal)
prior.kappa	Prior mean for kappa (overrides prior.range.nominal)
theta.prior.mean	(overrides prior.*)
theta.prior.prec	Scalar, vector or matrix, specifying the joint prior precision for <i>theta</i> .
n.iid.group	If greater than 1, build an explicitly iid replicated model, to support constraints applied to the combined replicates, for example in a time-replicated spatial model. Constraints can either be specified for a single mesh, in which case it's applied to the average of the replicates (ncol(A) should be mesh\$n for 2D meshes, mesh\$m for 1D), or as general constraints on the collection of replicates (ncol(A) should be mesh\$n * n.iid.group for 2D meshes, mesh\$m * n.iid.group for 1D).
...	Additional parameters for special uses.

Details

This method constructs a Matern SPDE model, with spatial scale parameter $\kappa(u)$ and variance rescaling parameter $\tau(u)$.

$$(\kappa^2(u) - \Delta)^{\alpha/2}(\tau(u))$$

$$x(u) = W(u)$$

Stationary models are supported for $0 < \alpha \leq 2$, with spectral approximation methods used for non-integer α , with approximation method determined by `fractional.method`.

Non-stationary models are supported for $\alpha = 2$ only, with

- $\log \tau(u) = B_0^\tau(u) + \sum_{k=1}^p B_k^\tau(u)\theta_k$
- $\log \kappa(u) = B_0^\kappa(u) + \sum_{k=1}^p B_k^\kappa(u)\theta_k$

The same parameterisation is used in the stationary cases, but with B_0^τ , B_k^τ , B_0^κ , and B_k^κ constant across u .

Integration and other general linear constraints are supported via the `constr`, `extraconstr.int`, and `extraconstr` parameters, which also interact with `n.iid.group`.

Value

An `inla.spde2` object.

Functions

- `inla.spde2.theta2phi0`: Convert from theta vector to phi0 values in the internal spde2 model representation
- `inla.spde2.theta2phi1`: Convert from theta vector to phi1 values in the internal spde2 model representation
- `inla.spde2.theta2phi2`: Convert from theta vector to phi2 values in the internal spde2 model representation

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.2d\(\)](#), [inla.mesh.create\(\)](#), [inla.mesh.1d\(\)](#), [inla.mesh.basis\(\)](#), [inla.spde2.pcmatern\(\)](#), [inla.spde2.generic\(\)](#)

Examples

```
n = 100
field.fcn = function(loc) (10*cos(2*pi*2*(loc[,1]+loc[,2])))
loc = matrix(runif(n*2),n,2)
## One field, 2 observations per location
idx.y = rep(1:n,2)
y = field.fcn(loc[idx.y,]) + rnorm(length(idx.y))

mesh = inla.mesh.create(loc, refine=list(max.edge=0.05))
spde = inla.spde2.matern(mesh)
data = list(y=y, field=mesh$idx$loc[idx.y])
formula = y ~ -1 + f(field, model=spde)
result = inla(formula, data=data, family="normal")

## Plot the mesh structure:
plot(mesh)

if (require(rgl)) {
  col.pal = colorRampPalette(c("blue","cyan","green","yellow","red"))
  ## Plot the posterior mean:
  plot(mesh, rgl=TRUE,
        result$summary.random$field["mean"],
        color.palette = col.pal)
  ## Plot residual field:
  plot(mesh, rgl=TRUE,
        result$summary.random$field["mean"]-field.fcn(mesh$loc),
        color.palette = col.pal)
}

result.field = inla.spde.result(result, "field", spde)
plot(result.field$marginals.range.nominal[[1]])
```

`inla.spde2.matern.sd.basis`*Approximate variance-compensating basis functions*

Description

Calculates an approximate basis for tau and kappa for an `inla.spde2.matern` model where tau is a rescaling parameter.

Usage

```
inla.spde2.matern.sd.basis(  
  mesh,  
  B.sd,  
  B.range,  
  method = 1,  
  local.offset.compensation = FALSE,  
  alpha = 2,  
  ...  
)
```

Arguments

<code>mesh</code>	An inla.mesh() object.
<code>B.sd</code>	Desired basis for log-standard deviations.
<code>B.range</code>	Desired basis for spatial range.
<code>method</code>	Construction method selector. Expert option only.
<code>local.offset.compensation</code>	If FALSE, only compensate in the average for the tau offset.
<code>alpha</code>	The model alpha parameter.
<code>...</code>	Additional parameters passed on to internal <code>inla.spde2.matern</code> calls.

Value

List of basis specifications

<code>B.tau</code>	Basis for log(tau)
<code>B.kappa</code>	Basis for log(kappa)

Intended for passing on to [inla.spde2.matern\(\)](#).

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde2.matern\(\)](#)

inla.spde2.pcmatern *Matern SPDE model object with PC prior for INLA*

Description

Create an `inla.spde2` model object for a Matern model, using a PC prior for the parameters.

Usage

```
inla.spde2.pcmatern(
  mesh,
  alpha = 2,
  param = NULL,
  constr = FALSE,
  extraconstr.int = NULL,
  extraconstr = NULL,
  fractional.method = c("parsimonious", "null"),
  n.iid.group = 1,
  prior.range = NULL,
  prior.sigma = NULL
)
```

Arguments

<code>mesh</code>	The mesh to build the model on, as an <code>inla.mesh()</code> or <code>inla.mesh.1d()</code> object.
<code>alpha</code>	Fractional operator order, $0 < \alpha \leq 2$ supported, for $\nu = \alpha - d/2 > 0$.
<code>param</code>	Further model parameters. Not currently used.
<code>constr</code>	If TRUE, apply an integrate-to-zero constraint. Default FALSE.
<code>extraconstr.int</code>	Field integral constraints.
<code>extraconstr</code>	Direct linear combination constraints on the basis weights.
<code>fractional.method</code>	Specifies the approximation method to use for fractional (non-integer) alpha values. 'parsimonious' gives an overall approximate minimal covariance error, 'null' uses approximates low-order properties.
<code>n.iid.group</code>	If greater than 1, build an explicitly iid replicated model, to support constraints applied to the combined replicates, for example in a time-replicated spatial model. Constraints can either be specified for a single mesh, in which case it's applied to the average of the replicates (<code>ncol(A)</code> should be <code>mesh\$n</code> for 2D meshes, <code>mesh\$m</code> for 1D), or as general constraints on the collection of replicates (<code>ncol(A)</code> should be <code>mesh\$n * n.iid.group</code> for 2D meshes, <code>mesh\$m * n.iid.group</code> for 1D).
<code>prior.range</code>	A length 2 vector, with <code>(range0, Prange)</code> specifying that $P(\rho < \rho_0) = p_\rho$, where ρ is the spatial range of the random field. If <code>Prange</code> is NA, then <code>range0</code> is used as a fixed range value.
<code>prior.sigma</code>	A length 2 vector, with <code>(sigma0, Psigma)</code> specifying that $P(\sigma > \sigma_0) = p_\sigma$, where σ is the marginal standard deviation of the field. If <code>Psigma</code> is NA, then <code>sigma0</code> is used as a fixed range value.

Details

This method constructs a Matern SPDE model, with spatial range ρ and standard deviation parameter σ . In the parameterisation

$$\begin{aligned} & (\kappa^2 - \Delta)^{\alpha/2}(\tau \\ & x(u)) = W(u) \end{aligned}$$

the spatial scale parameter $\kappa = \sqrt{8\nu}/\rho$, where $\nu = \alpha - d/2$, and τ is proportional to $1/\sigma$.

Stationary models are supported for $0 < \alpha \leq 2$, with spectral approximation methods used for non-integer α , with approximation method determined by `fractional.method`.

Integration and other general linear constraints are supported via the `constr`, `extraconstr.int`, and `extraconstr` parameters, which also interact with `n.iid.group`.

The joint PC prior density for the spatial range, ρ , and the marginal standard deviation, σ , and is

$$\begin{aligned} \pi(\rho, \sigma) = & \\ & \frac{d\lambda_\rho}{2} \rho^{-1-d/2} \exp(-\lambda_\rho \rho^{-d/2}) \\ & \lambda_\sigma \exp(-\lambda_\sigma \sigma) \end{aligned}$$

where λ_ρ and λ_σ are hyperparameters that must be determined by the analyst. The practical approach for this in INLA is to require the user to indirectly specify these hyperparameters through

$$P(\rho < \rho_0) = p_\rho$$

and

$$P(\sigma > \sigma_0) = p_\sigma$$

where the user specifies the lower tail quantile and probability for the range (ρ_0 and p_ρ) and the upper tail quantile and probability for the standard deviation (σ_0 and p_σ).

This allows the user to control the priors of the parameters by supplying knowledge of the scale of the problem. What is a reasonable upper magnitude for the spatial effect and what is a reasonable lower scale at which the spatial effect can operate? The shape of the prior was derived through a construction that shrinks the spatial effect towards a base model of no spatial effect in the sense of distance measured by Kullback-Leibler divergence.

The prior is constructed in two steps, under the idea that having a spatial field is an extension of not having a spatial field. First, a spatially constant random effect ($\rho = \infty$) with finite variance is more complex than not having a random effect ($\sigma = 0$). Second, a spatial field with spatial variation ($\rho < \infty$) is more complex than the random effect with no spatial variation. Each of these extensions are shrunk towards the simpler model and, as a result, we shrink the spatial field towards the base model of no spatial variation and zero variance ($\rho = \infty$ and $\sigma = 0$).

The details behind the construction of the prior is presented in Fuglstad, et al. (2016) and is based on the PC prior framework (Simpson, et al., 2015).

Value

An `inla.spde2` object.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

References

Fuglstad, G.-A., Simpson, D., Lindgren, F., and Rue, H. (2016) Constructing Priors that Penalize the Complexity of Gaussian Random Fields. arXiv:1503.00256

Simpson, D., Rue, H., Martins, T., Riebler, A., and Sørbye, S. (2015) Penalising model component complexity: A principled, practical approach to constructing priors. arXiv:1403.4630

See Also

[inla.mesh.2d\(\)](#), [inla.mesh.create\(\)](#), [inla.mesh.1d\(\)](#), [inla.mesh.basis\(\)](#), [inla.spde2.matern\(\)](#), [inla.spde2.generic\(\)](#)

Examples

```
## Spatial interpolation
n = 100
field.fcn = function(loc) (10*cos(2*pi*2*(loc[,1]+loc[,2])))
loc = matrix(runif(n*2),n,2)
## One field, 2 observations per location
idx.y = rep(1:n,2)
y = field.fcn(loc[idx.y,]) + rnorm(length(idx.y))

mesh = inla.mesh.2d(loc, max.edge=0.05, cutoff=0.01)
spde = inla.spde2.pcmatern(mesh,
  prior.range=c(0.01,0.1), prior.sigma=c(100,0.1))
data = list(y=y, field=mesh$idx$loc[idx.y])
formula = y ~ -1 + f(field, model=spde)
result = inla(formula, data=data, family="normal")

## Plot the mesh structure:
plot(mesh)

if (require(rgl)) {
  col.pal = colorRampPalette(c("blue","cyan","green","yellow","red"))
  ## Plot the posterior mean:
  plot(mesh, rgl=TRUE,
    result$summary.random$field["mean"],
    color.palette = col.pal)
  ## Plot residual field:
  plot(mesh, rgl=TRUE,
    result$summary.random$field["mean"]-field.fcn(mesh$loc),
    color.palette = col.pal)
}

result.field = inla.spde.result(result, "field", spde)
par(mfrow=c(2,1))
plot(result.field$marginals.range.nominal[[1]],
  type="l", main="Posterior density for range")
plot(inla.tmarginal(sqrt, result.field$marginals.variance.nominal[[1]]),
  type="l", main="Posterior density for std.dev.")
par(mfrow=c(1,1))

## Spatial model
set.seed(1234234)

## Generate spatial locations
```

```

nObs = 200
loc = matrix(runif(nObs*2), nrow = nObs, ncol = 2)

## Generate observation of spatial field
nu = 1.0
rhoT = 0.2
kappaT = sqrt(8*nu)/rhoT
sigT = 1.0
Sig = sigT^2*inla.matern.cov(nu = nu,
                             kappa = kappaT,
                             x = as.matrix(dist(loc)),
                             d = 2,
                             corr = TRUE)

L = t(chol(Sig))
u = L %%% rnorm(nObs)

## Construct observation with nugget
sigN = 0.1
y = u + sigN*rnorm(nObs)

## Create the mesh and spde object
mesh = inla.mesh.2d(loc,
                    max.edge = 0.05,
                    cutoff = 0.01)
spde = inla.spde2.pcmatern(mesh,
                           prior.range = c(0.01, 0.05),
                           prior.sigma = c(10, 0.05))

## Create projection matrix for observations
A = inla.spde.make.A(mesh = mesh,
                     loc = loc)

## Run model without any covariates
idx = 1:spde$n.spde
res = inla(y ~ f(idx, model = spde) - 1,
           data = list(y = y, idx = idx, spde = spde),
           control.predictor = list(A = A))

## Re-run model with fixed range
spde.fixed = inla.spde2.pcmatern(mesh,
                                 prior.range = c(0.2, NA),
                                 prior.sigma = c(10, 0.05))

res.fixed = inla(y ~ f(idx, model = spde) - 1,
                 data = list(y = y, idx = idx, spde = spde.fixed),
                 control.predictor = list(A = A))

```

inla.spTransform

Wrapper method for sp::spTransform

Description

Handles transformation of various inla objects according to coordinate reference systems of sp::CRS or inla.CRS class.

Usage

```

inla.spTransform(x, ...)

## Default S3 method:
inla.spTransform(x, crs0, crs1, passthrough = FALSE, ...)

## S3 method for class 'SpatialPoints'
inla.spTransform(x, CRSobj, passthrough = FALSE, ...)

## S3 method for class 'SpatialPointsDataFrame'
inla.spTransform(x, CRSobj, passthrough = FALSE, ...)

## S3 method for class 'inla.mesh.lattice'
inla.spTransform(x, CRSobj, passthrough = FALSE, ...)

## S3 method for class 'inla.mesh.segment'
inla.spTransform(x, CRSobj, passthrough = FALSE, ...)

## S3 method for class 'inla.mesh'
inla.spTransform(x, CRSobj, passthrough = FALSE, ...)

```

Arguments

x	The object that should be transformed from its current CRS to a new CRS
...	Potential additional arguments
crs0	The source sp::CRS or inla.CRS object
crs1	The target sp::CRS or inla.CRS object
passthrough	default FALSE. Setting to TRUE allows objects with no CRS information to be passed through without transformation.
CRSobj	The target sp::CRS or inla.CRS object

Details

`inla.spTransform.default` Low level transformation of raw coordinates.

Value

The object is returned with its coordinates transformed

Author(s)

Finn Lindgren finn.lindgren@gmail.com

See Also

[inla.CRS\(\)](#)

Examples

```
if (require(rgdal)) {
  latt <- inla.mesh.lattice(-10:10, 40:60)
  mesh1 <- inla.mesh.create(
    lattice = latt, extend = FALSE, refine = FALSE,
    crs = inla.CRS("longlat")
  )
  mesh2 <- inla.spTransform(mesh1, inla.CRS("lambert"))
  summary(mesh1)
  summary(mesh2)
}
```

inla.sp_get_crs	<i>Extract CRS information</i>
-----------------	--------------------------------

Description

Wrapper for CRS(proj4args) (PROJ4) and CRS(wkt) for sp::Spatial objects.

This function is a convenience method to workaround PROJ4/PROJ6 differences, and the lack of a crs extraction method for Spatial objects.

Usage

```
inla.sp_get_crs(x)
```

Arguments

x A sp::Spatial object

Value

A CRS object, or NULL if no valid CRS identified

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

Examples

```
## Not run:
if (interactive()) {
  s <- sp::SpatialPoints(matrix(1:6, 3, 2), proj4string = inla.CRS("sphere"))
  inla.sp_get_crs(s)
}

## End(Not run)
```

inla.ssh.copy.id	<i>Setup remote computing</i>
------------------	-------------------------------

Description

Initialize the definition file and print the path to the internal script to transfer ssh-keys

Usage

```
inla.remote()
inla.ssh.copy.id()
```

Arguments

None

Value

inla.remote is used once to setup the remote host information file (definition file) in the users home directory; see the FAQ entry on this issue for more information. inla.ssh.copy.id will return the path to the internal script to transfer ssh-keys.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
##See the FAQ entry on this issue on r-inla.org.
```

inla.stack.remove.unused	<i>Data stacking for advanced INLA models</i>
--------------------------	---

Description

Functions for combining data, effects and observation matrices into inla.stack objects, and extracting information from such objects.

Usage

```
inla.stack.remove.unused(stack)

inla.stack.compress(stack, remove.unused = TRUE)

inla.stack(..., compress = TRUE, remove.unused = TRUE)

inla.stack.sum(
  data,
  A,
```

```

    effects,
    tag = "",
    compress = TRUE,
    remove.unused = TRUE
)

inla.stack.join(..., compress = TRUE, remove.unused = TRUE)

inla.stack.index(stack, tag)

inla.stack.LHS(stack)

inla.stack.RHS(stack)

inla.stack.data(stack, ...)

inla.stack.A(stack)

```

Arguments

stack	A <code>inla.data.stack</code> object, created by a call to <code>inla.stack</code> , <code>inla.stack.sum</code> , or <code>inla.stack.join</code> .
remove.unused	If TRUE, compress the model by removing rows of effects corresponding to all-zero columns in the A matrix (and removing those columns).
...	For <code>inla.stack.join</code> , two or more data stacks of class <code>inla.data.stack</code> , created by a call to <code>inla.stack</code> , <code>inla.stack.sum</code> , or <code>inla.stack.join</code> . For <code>inla.stack.data</code> , a list of variables to be joined with the data list.
compress	If TRUE, compress the model by removing duplicated rows of effects, replacing the corresponding A-matrix columns with a single column containing the sum.
data	A list or <code>codedata.frame</code> of named data vectors. Scalars are expanded to match the number of rows in the A matrices, or any non-scalar data vectors. An error is given if the input is inconsistent.
A	A list of observation matrices. Scalars are expanded to diagonal matrices matching the effect vector lengths. An error is given if the input is inconsistent or ambiguous.
effects	A collection of effects/predictors. Each list element corresponds to an observation matrix, and must either be a single vector, a list of vectors, or a <code>data.frame</code> . Single-element effect vectors are expanded to vectors matching the number of columns in the corresponding A matrix. An error is given if the input is inconsistent or ambiguous.
tag	A string specifying a tag for later identification.

Details

For models with a single effects collection, the outer list container for A and effects may be omitted.

Component size definitions:

- n_l effect blocks
- n_k effects

- n_i data values
- $n_{j,l}$ effect size for block l
- $n_j = \sum_{l=1}^{n_l} n_{j,l}$ total effect size

Input:

data (y^1, \dots, y^p) p vectors, each of length n_i

A (A^1, \dots, A^{n_l}) matrices of size $n_i \times n_{j,l}$

effects $((x^{1,1}, \dots, x^{n_k,1}), \dots, (x^{1,n_l}, \dots, x^{n_k,n_l}))$ collections of effect vectors of length $n_{j,l}$

$$\text{predictor}(y^1, \dots, y^p) \sim \sum_{l=1}^{n_l} A^l \sum_{k=1}^{n_k} g(k, x^{k,l}) = \tilde{A} \sum_{k=1}^{n_k} g(k, \tilde{x}^k)$$

where

$$\tilde{A} = \text{cbind}(A^1, \dots, A^{n_l})$$

and

$$\tilde{x}^k = \text{rbind}(x^{k,1}, \dots, x^{k,n_l})$$

and for each block l , any missing $x^{k,l}$ is replaced by an NA vector.

Value

A data stack of class `inla.data.stack`. Elements:

- `data` = $(y^1, \dots, y^p, \tilde{x}^1, \dots, \tilde{x}^{n_k})$
- `A` = \tilde{A}
- `data.names` List of data names, length p
- `effect.names` List of effect names, length n_k
- `n.data` Data length, n_i
- `index` List indexed by tags, each element indexing into $i = 1, \dots, n_i$

Functions

- `inla.stack.remove.unused`: Remove unused entries from an existing stack
- `inla.stack.compress`: Compress an existing stack by removing duplicates
- `inla.stack.sum`: Create data stack as a sum of predictors
- `inla.stack.join`: Join two or more data stacks
- `inla.stack.index`: Extract tagged indices
- `inla.stack.LHS`: Extract data associated with the "left hand side" of the model (e.g. the data itself, `Ntrials`, `link`, `E`)
- `inla.stack.RHS`: Extract data associated with the "right hand side" of the model (all the covariates/predictors)
- `inla.stack.data`: Extract data for an `inla` call, and optionally join with other variables
- `inla.stack.A`: Extract the "A matrix" for `control.predictor`

Functions

- `inla.stack.remove.unused`: Remove unused entries from an existing stack
- `inla.stack.compress`: Compress an existing stack by removing duplicates
- `inla.stack`: Shorthand for `inla.stack.join` and `inla.stack.sum`
- `inla.stack.sum`: Create data stack as a sum of predictors
- `inla.stack.join`: Join two or more data stacks
- `inla.stack.index`: Extract tagged indices
- `inla.stack.LHS`: Extract data associated with the "left hand side" of the model (e.g. the data itself, `Ntrials`, `link`, `E`)
- `inla.stack.RHS`: Extract data associated with the "right hand side" of the model (all the covariates/predictors)
- `inla.stack.data`: Extract data for an `inla` call, and optionally join with other variables
- `inla.stack.A`: Extract the "A matrix" for control.predictor

See Also

[inla.spde.make.A\(\)](#), [inla.spde.make.index\(\)](#)

Examples

```
n <- 200
loc <- matrix(runif(n * 2), n, 2)
mesh <- inla.mesh.2d(
  loc.domain = loc,
  max.edge = c(0.05, 0.2)
)
proj.obs <- inla.mesh.projector(mesh, loc = loc)
proj.pred <- inla.mesh.projector(mesh, loc = mesh$loc)
spde <- inla.spde2.pcmatern(mesh,
  prior.range = c(0.01, 0.01),
  prior.sigma = c(10, 0.01)
)

covar <- rnorm(n)
field <- inla.qsample(n = 1, Q = inla.spde.precision(spde, theta = log(c(0.5, 1))))[, 1]
y <- 2 * covar + inla.mesh.project(proj.obs, field)

A.obs <- inla.spde.make.A(mesh, loc = loc)
A.pred <- inla.spde.make.A(mesh, loc = proj.pred$loc)
stack.obs <-
  inla.stack(
    data = list(y = y),
    A = list(A.obs, 1),
    effects = list(c(
      list(Intercept = 1),
      inla.spde.make.index("spatial", spde$n.spde)
    )),
    covar = covar
  ),
  tag = "obs"
)
stack.pred <-
```

```

    inla.stack(
      data = list(y = NA),
      A = list(A.pred),
      effects = list(c(
        list(Intercept = 1),
        inla.spde.make.index("spatial", mesh$n)
      )),
      tag = "pred"
    )
  stack <- inla.stack(stack.obs, stack.pred)

  formula <- y ~ -1 + Intercept + covar + f(spatial, model = spde)
  result1 <- inla(formula,
    data = inla.stack.data(stack.obs, spde = spde),
    family = "gaussian",
    control.predictor = list(
      A = inla.stack.A(stack.obs),
      compute = TRUE
    )
  )

  plot(y, result1$summary.fitted.values[inla.stack.index(stack.obs, "obs")$data, "mean"],
    main = "Observations vs posterior predicted values at the data locations"
  )

  result2 <- inla(formula,
    data = inla.stack.data(stack, spde = spde),
    family = "gaussian",
    control.predictor = list(
      A = inla.stack.A(stack),
      compute = TRUE
    )
  )

  field.pred <- inla.mesh.project(
    proj.pred,
    result2$summary.fitted.values[inla.stack.index(stack, "pred")$data, "mean"]
  )
  field.pred.sd <- inla.mesh.project(
    proj.pred,
    result2$summary.fitted.values[inla.stack.index(stack, "pred")$data, "sd"]
  )

  plot(field, field.pred, main = "True vs predicted field")
  abline(0, 1)
  image(inla.mesh.project(mesh,
    field = field,
    dims = c(200, 200)
  ),
  main = "True field"
  )
  image(inla.mesh.project(mesh,
    field = field.pred,
    dims = c(200, 200)
  ),
  main = "Posterior field mean"
  )

```

```

image(inla.mesh.project(mesh,
  field = field.pred.sd,
  dims = c(200, 200)
),
main = "Prediction standard deviation"
)
plot(field, (field.pred - field) / 1,
  main = "True field vs standardised prediction residuals"
)

```

inla.surv

Create a Survival Object for INLA

Description

Create a survival object, to be used as a response variable in a model formula for the `inla` function for survival models.

Usage

```

inla.surv(time, event, time2, truncation, subject = NULL, .special = NULL)
## S3 method for class 'inla.surv'
plot(x, y, ...)
## S3 method for class 'inla.surv'
print(x, ...)
is.inla.surv(object)
as.inla.surv(object, ...)

```

Arguments

<code>time</code>	For right censored data, this is the follow up time. For interval data, this is the starting time for the interval. For in-interval event, this is the observed time (in the interval) for the event. For left censored data, this the censoring time.
<code>event</code>	The status indicator, 1=observed event, 0=right censored event, 2=left censored event, 3=interval censored event, and 4=observed event in an interval (left, right).
<code>time2</code>	Ending time for the interval censored data or an in-interval event.
<code>truncation</code>	Left truncation. If missing it is assumed to be 0. The lower limit for event=4.
<code>subject</code>	Patient number in multiple event data, not needed otherwise.
<code>.special</code>	An internal object, not for public use
<code>object</code>	Any R-object
<code>x</code>	Object to plot or print
<code>y</code>	Object to plot (not in use)
<code>...</code>	Additional argument

Value

An object of class `inla.surv`. There are methods for `print`, `plot` for `inla.surv` objects.

`is.inla.surv` returns TRUE if object inherits from class `inla.surv`, otherwise FALSE.

`as.inla.surv` returns an object of class `inla.surv`

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

update.packages

inla.version	<i>Show the version of the INLA-package</i>
--------------	---

Description

Show the version of the INLA-package

Usage

```
inla.version(what = c("default", "version", "date"))
```

Arguments

what	What to show version of
------	-------------------------

Value

inla.version display the current version information using cat with default or info, or return other specific requests through the call.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
## Summary of all
inla.version()
## The building date
inla.version("date")
```

inla.wkt_is_geocent *Handling CRS/WKT*

Description

Get and set CRS object or WKT string properties.

Usage

```

inla.wkt_is_geocent(wkt)

inla.crs_is_geocent(crs)

inla.wkt_get_ellipsoid_radius(wkt)

inla.crs_get_ellipsoid_radius(crs)

inla.wkt_set_ellipsoid_radius(wkt, radius)

inla.crs_set_ellipsoid_radius(crs, radius)

inla.wkt_unit_params()

inla.wkt_get_lengthunit(wkt)

inla.wkt_set_lengthunit(wkt, unit, params = NULL)

inla.crs_get_wkt(crs)

inla.crs_get_lengthunit(crs)

inla.crs_set_lengthunit(crs, unit, params = NULL)

```

Arguments

wkt	A WKT2 character string
crs	A <code>sp::CRS</code> or <code>inla.CRS</code> object
unit	character, name of a unit. Supported names are "metre", "kilometre", and the aliases "meter", "m", "International metre", "kilometer", and "km", as defined by <code>inla.wkt_unit_params</code> or the <code>params</code> argument. (For legacy PROJ4 use, only "m" and "km" are supported)
params	Length unit definitions, in the list format produced by <code>inla.wkt_unit_params()</code> , Default: <code>NULL</code> , which invokes <code>inla.wkt_unit_params()</code>

Value

For `inla.wkt_unit_params`, a list of named unit definitions

For `inla.wkt_get_lengthunit`, a list of length units used in the wkt string, excluding the ellipsoid radius unit.

For `inla.wkt_set_lengthunit`, a WKT2 string with altered length units. Note that the length unit for the ellipsoid radius is unchanged.

For `inla.crs_get_wkt`, WKT2 string.

For `inla.crs_get_lengthunit`, a list of length units used in the wkt string, excluding the ellipsoid radius unit. (For legacy PROJ4 code, the raw units from the proj4string are returned, if present.)

For `inla.crs_set_lengthunit`, a `sp::CRS` object with altered length units. Note that the length unit for the ellipsoid radius is unchanged.

For `inla.wkt_unit_params`, a list of named unit definitions

For `inla.wkt_get_lengthunit`, a list of length units used in the wkt string, excluding the ellipsoid radius unit.

For `inla.wkt_set_lengthunit`, a WKT2 string with altered length units. Note that the length unit for the ellipsoid radius is unchanged.

For `inla.crs_get_wkt`, WKT2 string.

For `inla.crs_get_lengthunit`, a list of length units used in the wkt string, excluding the ellipsoid radius unit. (For legacy PROJ4 code, the raw units from the proj4string are returned, if present.)

For `inla.crs_set_lengthunit`, a `sp::CRS` object with altered length units. Note that the length unit for the ellipsoid radius is unchanged.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.sp_get_crs\(\)](#)

[inla.sp_get_crs\(\)](#)

Examples

```
## Not run:
if (inla.has_PROJ6()) {
  c1 <- inla.CRS("globe")
  inla.crs_get_lengthunit(c1)
  c2 <- inla.crs_set_lengthunit(c1, "km")
  inla.crs_get_lengthunit(c2)
}

## End(Not run)

## Not run:
if (inla.has_PROJ6()) {
  c1 <- inla.CRS("globe")
  inla.crs_get_lengthunit(c1)
  c2 <- inla.crs_set_lengthunit(c1, "km")
  inla.crs_get_lengthunit(c2)
}

## End(Not run)
```

 joint.marginal

Sample, transform and evaluate from a joint marginal approximation

Description

Sample, transform and evaluate from from a joint marginal approximation as returned using argument selection in inla.

Usage

```
inla.rjmarginal(n, jmarginal, constr)
inla.rjmarginal.eval(fun, samples, ...)
inla.tjmarginal(jmarginal, A)
inla.1djmarginal(jmarginal)
```

Arguments

n	The number of samples
jmarginal	A marginal object given either by a inla object or result\$selection
constr	Optional linear constraints; see ?INLA::f and argument extraconstr
fun	A function which is evaluated for each sample, similar to inla.posterior.sample.eval: please see the documentation for this functions for details.
samples	The samples, as in the form of the output from inla.rjmarginal
A	A matrix used for the linear combination

Value

THESE FUNCTIONS ARE EXPERIMENTAL FOR THE MOMENT (JULY 2020)

inla.rjmarginal returns a list with the samples in samples (matrix) and the corresponding log-densities in log.density (vector). Each column in samples contains one sample.

inla.rjmarginal.eval returns a matrix, where each row is the (vector) function evaluated at each sample.

inla.tjmarginal returns a inla.jmarginal-object of the linear combination defined by the matrix A.

inla.1djmarginal return the marginal densities from a joint approximation.

Author(s)

Cristian Chiuchio and Havard Rue <hrue@r-inla.org>

See Also

[inla](#)

Examples

```

n = 10
x = 1+rnorm(n)
xx = 3 + rnorm(n)
y = 1 + x + xx + rnorm(n)
selection = list(xx=1, Predictor = 3:4, x=1)
r = inla(y ~ 1 + x + xx,
        data = data.frame(y, x, xx),
        selection = selection)
ns = 100
xx = inla.rjmarginal(ns, r)

print(cbind(mean = r$selection$mean, sample.mean = rowMeans(xx$samples)))
print("cov matrix")
print(round(r$selection$cov.matrix, dig=3))
print("sample cov matrix")
print(round(cov(t(xx$samples)), dig=3))

skew = function(z) mean((z-mean(z))^3)/var(z)^1.5
print(round(cbind(skew = r$selection$skewness,
                 sample.skew = apply(xx$sample, 1, skew)), dig=3))

## illustrating the eval function
n = 10
x = rnorm(n)
eta = 1 + x
y = eta + rnorm(n, sd=0.1)
selection = list(x = 1, Predictor = c(1, 2, 4, 5), '(Intercept)' = 1)
r = inla(y ~ 1 + x,
        data = data.frame(y, x),
        selection = selection)
xx = inla.rjmarginal(100, r)
xx.eval = inla.rjmarginal.eval(function() c(x, Predictor, Intercept), xx)
print(cbind(xx$samples[, 1]))
print(cbind(xx.eval[, 1]))

constr <- list(A = matrix(1, ncol = n, nrow = 1), e = 1)
x <- inla.rjmarginal(10, r, constr = constr)

A <- matrix(rnorm(n^2), n, n)
b <- inla.tjmarginal(r, A)
b.marg <- inla.1djmarginal(b)

```

jp.define

*Joint-prior models***Description**

A framework for defining joint priors in R

Usage

```
inla.jp.define(jp = NULL, ...)
```

Arguments

- jp** The `jp`-function which returns the joint log-prior as a function of argument `theta`. There is an optional second argument that is a vector of `theta`-names. If second argument is not present, argument `.theta.desc` will be added.
- ...** Named list of variables that defines the environment of `jp`

Value

- This allows joint priors to be defined in R.
- This function is for internal use only.

Author(s)

Havard Rue <hrue@r-inla.org>

Kidney

Kidney infection data

Description

Times of infection from the time to insertion of the catheter for 38 kidney patients using portable dialysis equipment

Format

A data frame with 76 observations on the following 9 variables.

time a numeric vector. Time to infection from the insertion of catheter

event a numeric vector. 1: time of infection 0: time of censoring

age a numeric vector. Age of the patient at the time of infection

sex a numeric vector. Sex of the patient 0: male 1:female

disease a numeric vector. Type of disease

dis1 a numeric vector. Dummy variable to codify the disease type.

dis2 a numeric vector. Dummy variable to codify the disease type.

dis3 a numeric vector. Dummy variable to codify the disease type.

ID a numeric vector. Patient code.

References

McGilchrist and C.W. Aisbett (1991), Regression with frailty in survival analysis, *Biometrics*, vol.47, pages 461–166.

D.J. Spiegelhalter and A. Thomas and N.G. Best and W.R. Gilks (1995) BUGS: Bayesian Inference Using Gibbs sampling, Version 0.50., MRC Biostatistics Unit, Cambridge, England.

`lattice2node`*Functions to define mapping between a lattice and nodes*

Description

These functions define mapping in between two-dimensional indices on a lattice and the one-dimensional node representation used in `inla`.

The mapping from node to lattice follows the default R behaviour (which is column based storage), and `as.vector(A)` and `matrix(a,nrow,ncol)` can be used instead of `inla.matrix2vector` and `inla.vector2matrix`.

Usage

```
inla.lattice2node.mapping(nrow, ncol)
inla.node2lattice.mapping(nrow, ncol)
inla.lattice2node(irow, icol, nrow, ncol)
inla.node2lattice(node, nrow, ncol)
inla.matrix2vector(a.matrix)
inla.vector2matrix(a.vector, nrow, ncol)
```

Arguments

<code>nrow</code>	Number of rows in the lattice.
<code>ncol</code>	Number of columns in the lattice.
<code>irow</code>	Lattice row index, between 1 and <code>nrow</code>
<code>icol</code>	Lattice column index, between 1 and <code>ncol</code>
<code>node</code>	The node index, between 1 and <code>ncol*nrow</code>
<code>a.matrix</code>	is a matrix to be mapped to a vector using internal representation defined by <code>inla.lattice2node</code>
<code>a.vector</code>	is a vector to be mapped into a matrix using the internal representation defined by <code>inla.node2lattice</code>

Value

`inla.lattice2node.mapping` returns the hole mapping as a matrix, and `inla.node2lattice.mapping` returns the hole mapping as `list(irow=..., icol=...)`. `inla.lattice2node` and `inla.node2lattice` provide the mapping for a given set of lattice indices and nodes. `inla.matrix2vector` provide the mapped vector from a matrix, and `inla.vector2matrix` provide the inverse mapped matrix from vector.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla](#)

Examples

```

## write out the mapping using the two alternatives
nrow = 2
ncol = 3
mapping = inla.lattice2node.mapping(nrow,ncol)

for (i in 1:nrow){
  for(j in 1:ncol){
    print(paste("Alt.1: lattice index [", i,",", j,"] corresponds",
               "to node [", mapping[i,j],"]", sep=""))
  }
}

for (i in 1:nrow){
  for(j in 1:ncol){
    print(paste("Alt.2: lattice index [", i,",", j,"] corresponds to node [",
               inla.lattice2node(i,j,nrow,ncol), "]", sep=""))
  }
}

inv.mapping = inla.node2lattice.mapping(nrow,ncol)
for(node in 1:(nrow*ncol))
  print(paste("Alt.1: node [", node, "] corresponds to lattice index [",
             inv.mapping$irrow[node], ",",
             inv.mapping$icol[node],"]", sep=""))

for(node in 1:(nrow*ncol))
  print(paste("Alt.2: node [", node, "] corresponds to lattice index [",
             inla.node2lattice(node,nrow,ncol)$irrow[1], ",",
             inla.node2lattice(node,nrow,ncol)$icol[1],"]", sep=""))

## apply the mapping from matrix to vector and back
n = nrow*ncol
z = matrix(1:n,nrow,ncol)
z.vector = inla.matrix2vector(z) # as.vector(z) could also be used
print(mapping)
print(z)
print(z.vector)

## the vector2matrix is the inverse, and should give us the z-matrix
## back. matrix(z.vector, nrow, ncol) could also be used here.
z.matrix = inla.vector2matrix(z.vector, nrow, ncol)
print(z.matrix)

```

Leuk

The Leukemia data

Description

This the Leukemia data from Henderson et al (2003); see source.

Format

A data frame with 1043 observations on the following 9 variables.

```
time TODO
cens TODO
xcoord TODO
ycoord TODO
age TODO
sex TODO
wbc TODO
tpi TODO
district TODO
```

Source

This is the dataset from

Henderson, R. and Shimakura, S. and Gorst, D., 2002, Modeling spatial variation in leukemia survival data, JASA, 97, 460, 965–972.

Examples

```
data(Leuk)
```

```
lines.inla.mesh.segment
```

```
Draw inla.mesh.segment objects.
```

Description

Draws a `inla.mesh.segment()` object with generic or rgl graphics.

Usage

```
## S3 method for class 'inla.mesh.segment'
lines(
  x,
  loc = NULL,
  col = NULL,
  colors = c("black", "blue", "red", "green"),
  add = TRUE,
  xlim = NULL,
  ylim = NULL,
  rgl = FALSE,
  ...
)
```

Arguments

x	An <code>inla.mesh.segment()</code> object.
loc	Point locations to be used if <code>x\$loc</code> is NULL.
col	Segment color specification.
colors	Colors to cycle through if <code>col</code> is NULL.
add	If TRUE, add to the current plot, otherwise start a new plot.
xlim	X axis limits for a new plot.
ylim	Y axis limits for a new plot.
rgl	If TRUE, use <code>rgl</code> for plotting.
...	Additional parameters, passed on to graphics methods.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.segment\(\)](#)

link

Link functions in INLA

Description

Define link-functions and its inverse

Usage

```

inla.link.log(x, inverse=FALSE)
inla.link.invlog(x, inverse=FALSE)
inla.link.neglog(x, inverse=FALSE)
inla.link.invneglog(x, inverse=FALSE)
inla.link.logit(x, inverse=FALSE)
inla.link.invlogit(x, inverse=FALSE)
inla.link.probit(x, inverse=FALSE)
inla.link.invprobit(x, inverse=FALSE)
inla.link.cloglog(x, inverse=FALSE)
inla.link.invcloglog(x, inverse=FALSE)
inla.link.loglog(x, inverse=FALSE)
inla.link.invloglog(x, inverse=FALSE)
inla.link.tan(x, inverse=FALSE)
inla.link.invtan(x, inverse=FALSE)
inla.link.cauchit(x, inverse=FALSE)
inla.link.invcauchit(x, inverse=FALSE)
inla.link.identity(x, inverse=FALSE)
inla.link.invidentity(x, inverse=FALSE)
inla.link.inverse(x, inverse=FALSE)
inla.link.ininverse(x, inverse=FALSE)

```

```

inla.link.robit(x, df=7, inverse=FALSE)
inla.link.invrobit(x, df=7, inverse=FALSE)
inla.link.sn(x, intercept=0.5, skew=0, a=0, inverse=FALSE)
inla.link.invsn(x, intercept=0.5, skew=0, a=0, inverse=FALSE)
inla.link.invalid(x, inverse=FALSE)
inla.link.ininvalid(x, inverse=FALSE)

```

Arguments

x	The argument. A numeric vector.
df	The degrees of freedom for the Student-t
inverse	Logical. Use the link (inverse=FALSE) or its inverse (inverse=TRUE)
intercept	The quantile level for the intercept in the Skew-Normal link
skew	The skewness in the Skew-Normal. Not both of skew and a can be given.
a	The a-parameter in the Skew-Normal. Not both of skew and a can be given.

Value

Return the values of the link-function or its inverse.

Note

The inv-functions are redundant, as `inla.link.invlog(x) = inla.link.log(x, inverse=TRUE)` and so on, but they are simpler to use a arguments to other functions.

Author(s)

Havard Rue <hrue@r-inla.org>

make.lincomb	<i>Create linear combinations</i>
--------------	-----------------------------------

Description

Create a linear combination or several linear combinations, as input to `inla(..., lincomb = <lincomb>)`

Usage

```

inla.make.lincomb(...)
inla.make.lincombs(...)

```

Arguments

... Arguments; see examples

Value

A structure to be passed on to `inla` argument `lincomb`

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

TODO

Examples

```
##See the worked out examples and description in the FAQ
##section on {www.r-inla.org}
```

marginal

Functions which operates on marginals

Description

Density, distribution function, quantile function, random generation, hpd-interval, interpolation, expectations, mode and transformations of marginals obtained by `inla` or `inla.hyperpar()`. These functions computes the density (`inla.dmarginal`), the distribution function (`inla.pmarginal`), the quantile function (`inla.qmarginal`), random generation (`inla.rmarginal`), spline smoothing (`inla.smarginal`), computes expected values (`inla.emarginal`), computes the mode (`inla.mmarginal`), transforms the marginal (`inla.tmarginal`), and provide summary statistics (`inla.zmarginal`).

Usage

```
inla.dmarginal(x, marginal, log = FALSE)
inla.pmarginal(q, marginal, normalize = TRUE, len = 2048L)
inla.qmarginal(p, marginal, len = 2048L)
inla.rmarginal(n, marginal)
inla.hpdmarginal(p, marginal, len = 2048L)
inla.smarginal(marginal, log = FALSE, extrapolate = 0.0, keep.type = FALSE, factor=15L)
inla.emarginal(fun, marginal, ...)
inla.mmarginal(marginal)
inla.tmarginal(fun, marginal, n=2048L, h.diff = .Machine[["double.eps"]]^(1.0/3.0),
               method = c("quantile", "linear"))
inla.zmarginal(marginal, silent = FALSE)
```

Arguments

<code>marginal</code>	A marginal object from either <code>inla</code> or <code>inla.hyperpar()</code> , which is either <code>list(x=c(), y=c())</code> with density values <code>y</code> at locations <code>x</code> , or a <code>matrix(,n,2)</code> for which the density values are the second column and the locations in the first column. The <code>inla.hpdmarginal()</code> -function assumes a unimodal density.
<code>fun</code>	A (vectorised) function like <code>function(x) exp(x)</code> to compute the expectation against, or which define the transformation <code>new = fun(old)</code>
<code>x</code>	Evaluation points
<code>q</code>	Quantiles
<code>p</code>	Probabilities

n	The number of observations. If <code>length(n) > 1</code> , the length is taken to be the number required.
h.diff	The step-length for the numerical differentiation inside <code>inla.tmargin</code>
...	Further arguments to be passed to function which expectation is to be computed.
log	Return density or interpolated density in log-scale?
normalize	Renormalise the density after interpolation?
len	Number of locations used to interpolate the distribution function.
keep.type	If FALSE then return a <code>list(x=,y=)</code> , otherwise if TRUE, then return a matrix if the input is a matrix
extrapolate	How much to extrapolate on each side when computing the interpolation. In fraction of the range.
factor	The number of points after interpolation is factor times the original number of points; which is argument n in <code>spline</code>
method	Which method should be used to layout points for where the transformation is computed.
silent	Output the result visually (TRUE) or just through the call.

Value

`inla.smarginal` returns `list=c(x=c(),y=c())` of interpolated values do extrapolation using the factor given, and the remaining function returns what they say they should do.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla](#), [inla.hyperpar](#)

Examples

```
## a simple linear regression example
n = 10
x = rnorm(n)
sd = 0.1
y = 1+x + rnorm(n,sd=sd)
res = inla(y ~ 1 + x, data = data.frame(x,y),
          control.family=list(initial = log(1/sd^2L),fixed=TRUE))

## chose a marginal and compare the with the results computed by the
## inla-program
r = res$summary.fixed["x",]
m = res$marginals.fixed$x

## compute the 95% HPD interval
inla.hpdmarginal(0.95, m)

x = seq(-6, 6, len = 1000)
y = dnorm(x)
inla.hpdmarginal(0.95, list(x=x, y=y))
```

```
## compute the the density for exp(r), version 1
r.exp = inla.tmarginal(exp, m)
## or version 2
r.exp = inla.tmarginal(function(x) exp(x), m)

## to plot the marginal, we use the inla.smarginal, which interpolates (in
## log-scale). Compare with some samples.
plot(inla.smarginal(m), type="l")
s = inla.rmarginal(1000, m)
hist(inla.rmarginal(1000, m), add=TRUE, prob=TRUE)
lines(density(s), lty=2)

m1 = inla.emarginal(function(x) x, m)
m2 = inla.emarginal(function(x) x^2L, m)
stdev = sqrt(m2 - m1^2L)
q = inla.qmarginal(c(0.025,0.975), m)

## inla-program results
print(r)

## inla.marginal-results (they shouldn't be perfect!)
print(c(mean=m1, sd=stdev, "0.025quant" = q[1], "0.975quant" = q[2L]))
## using the buildt-in function
inla.zmarginal(m)
```

meshbuilder

Interactive mesh building and diagnostics

Description

Interactively design and build a triangle mesh for use with SPDE models, and assess the finite element approximation errors. The R code needed to recreate the mesh outside the interactive Shiny app is also generated. Spatial objects can be imported from the global workspace.

Usage

```
meshbuilder()
```

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.mesh.2d\(\)](#), [inla.mesh.create\(\)](#)

Examples

```
## Not run:
meshbuilder()

## End(Not run)
```

Munich

The Munich rent data

Description

The Munich rent data

Format

A data frame with 2035 observations on the following 17 variables.

"rent Net rent per square meter.

"floor.size Size of the flat in square meters.

"year Year of construction of the building in which the flat is located.

location Location index (in terms of subquarters).

Gute.Wohnlage Dummy variable for good locations / good neighborhoods.

Beste.Wohnlage Dummy variable for very good locations / very good neighborhoods.

Keine.Wvv Dummy for absence of warm water supply.

Keine.Zh Dummy for absence of central heating system.

Kein.Badkach Dummy for absence of flagging in the bathroom.

Besond.Bad Dummy for special features of the bathroom.

Gehobene.Kueche Dummy for more refined kitchen equipment.

zim1 Dummy for a flat with 1 room.

zim2 Dummy for a flat with 2 rooms.

zim3 Dummy for a flat with 3 rooms.

zim4 Dummy for a flat with 4 rooms.

zim5 Dummy for a flat with 5 rooms.

zim6 Dummy for a flat with 6 rooms.

Source

See Rue and Held (2005), Chapter 4.

References

Rue, H and Held, L. (2005) *Gaussian Markov Random Fields - Theory and Applications* Chapman and Hall

 nwEngland

The New England map

Description

This map is used in association to the Leukemia data from Henderson et al (2003); see source.

Format

A SpatialPolygons object.

Source

This map are used to analyse the Leukaemia dataset from

Henderson, R. and Shimakura, S. and Gorst, D., 2002, Modeling spatial variation in leukemia survival data, *JASA*, 97, 460, 965–972.

Examples

```
data(Leuk)
plot(nwEngland)
```

 Oral

~~ data name/kind ... ~~

Description

~~ A concise (1-5 lines) description of the dataset. ~~

Format

A data frame with 544 observations on the following 3 variables.

region a numeric vector

E a numeric vector

Y a numeric vector

References

Rue, H and Held, L. (2005) *Gaussian Markov Random Fields - Theory and Applications* Chapman and Hall

param2.matern.orig *Parameter settings for inla.spde2.matern models.*

Description

Construct parameter settings for `inla.spde2.matern` models.

Usage

```
param2.matern.orig(  
  mesh,  
  alpha = 2,  
  B.tau = matrix(c(0, 1, 0), 1, 3),  
  B.kappa = matrix(c(0, 0, 1), 1, 3),  
  prior.variance.nominal = 1,  
  prior.range.nominal = NULL,  
  prior.tau = NULL,  
  prior.kappa = NULL,  
  theta.prior.mean = NULL,  
  theta.prior.prec = 0.1  
)
```

Arguments

<code>mesh</code>	The mesh to build the model on, as an <code>inla.mesh()</code> object.
<code>alpha</code>	Fractional operator order, $0 < \alpha \leq 2$ supported. ($\nu = \alpha - d/2$)
<code>B.tau</code>	Matrix with specification of log-linear model for τ .
<code>B.kappa</code>	Matrix with specification of log-linear model for κ .
<code>prior.variance.nominal</code>	Nominal prior mean for the field variance
<code>prior.range.nominal</code>	Nominal prior mean for the spatial range
<code>prior.tau</code>	Prior mean for tau (overrides <code>prior.variance.nominal</code>)
<code>prior.kappa</code>	Prior mean for kappa (overrides <code>prior.range.nominal</code>)
<code>theta.prior.mean</code>	(overrides <code>prior.*</code>)
<code>theta.prior.prec</code>	Scalar, vector or matrix, specifying the joint prior precision for <i>theta</i> .

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[inla.spde2.matern\(\)](#)

pc.alphaw	<i>Utility functions for the PC prior for the alpha parameter in the Weibull likelihood</i>
-----------	---

Description

Functions to evaluate, sample, compute quantiles and percentiles of the PC prior for the alpha parameter in the Weibull likelihood

Usage

```
inla.pc.ralphaw(n, lambda = 5)
inla.pc.dalphaw(alpha, lambda = 5, log = FALSE)
inla.pc.qalphaw(p, lambda = 5)
inla.pc.palphaw(q, lambda = 5)
```

Arguments

n	Number of observations
lambda	The rate parameter in the PC-prior
alpha	Vector of evaluation points, where $\alpha > 0$.
log	Logical. Return the density in natural or log-scale.
p	Vector of probabilities
q	Vector of quantiles

Details

This gives the PC prior for the alpha parameter for the Weibull likelihood, where $\alpha=1$ is the base model.

Value

`inla.pc.dalphaw` gives the density, `inla.pc.palphaw` gives the distribution function, `inla.pc.qalphaw` gives the quantile function, and `inla.pc.ralphaw` generates random deviates.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

`inla.doc("pc.alphaw")`

Examples

```
x = inla.pc.ralphaw(100, lambda = 5)
d = inla.pc.dalphaw(x, lambda = 5)
x = inla.pc.qalphaw(0.5, lambda = 5)
inla.pc.palphaw(x, lambda = 5)
```

pc.ar

*Utility functions for the PC prior for a an AR(p) model***Description**

Functions to evaluate and sample from the PC prior for an AR(p) model

Usage

```
inla.pc.ar.rpacf(n=1, p, lambda = 1)
inla.pc.ar.dpacf(pac, lambda = 1, log = TRUE)
```

Arguments

p	The order of the AR-model
pac	A vector of partial autocorrelation coefficients
n	Number of observations
lambda	The rate parameter in the prior
log	Logical. Return the density in natural or log-scale.

Value

inla.pc.ar.rpac generate samples from the prior, returning a matrix where each row is a sample of theta. inla.pc.ar.dpac evaluates the density of pac. Use inla.ar.pacf2phi, inla.ar.phi2pacf, inla.ar.pacf2acf and inla.ar.acf2pacf to convert between various parameterisations.

Author(s)

Havard Rue <hrue@r-inla.org>

pc.cor0

*Utility functions for the PC prior for correlation in AR(1)***Description**

Functions to evaluate, sample, compute quantiles and percentiles of the PC prior for the correlation in the Gaussian AR(1) model where the base-model is zero correlation.

Usage

```
inla.pc.rcor0(n, u, alpha, lambda)
inla.pc.dcor0(cor, u, alpha, lambda, log = FALSE)
inla.pc.qcor0(p, u, alpha, lambda)
inla.pc.pcor0(q, u, alpha, lambda)
```

Arguments

n	Number of observations
u	The upper limit (see Details)
alpha	The probability going above the upper limit (see Details)
lambda	The rate parameter (see Details)
cor	Vector of correlations
log	Logical. Return the density in natural or log-scale.
p	Vector of probabilities
q	Vector of quantiles

Details

The statement $\text{Prob}(|\text{cor}| > u) = \text{alpha}$ is used to determine lambda unless lambda is given. Either lambda must be given, or u AND alpha. The density is symmetric around zero.

Value

`inla.pc.dcor0` gives the density, `inla.pc.pcor0` gives the distribution function, `inla.pc.qcor0` gives the quantile function, and `inla.pc.rcor0` generates random deviates.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

`inla.doc("pc.rho0")`

Examples

```
cor = inla.pc.rcor0(100, lambda = 1)
d = inla.pc.dcor0(cor, lambda = 1)
cor = inla.pc.qcor0(c(0.3, 0.7), u = 0.5, alpha=0.01)
inla.pc.pcor0(cor, u = 0.5, alpha=0.01)
```

pc.cor1

Utility functions for the PC prior for correlation in AR(1)

Description

Functions to evaluate, sample, compute quantiles and percentiles of the PC prior for the correlation in the Gaussian AR(1) model where the base-model is correlation one.

Usage

```
inla.pc.rcor1(n, u, alpha, lambda)
inla.pc.dcor1(cor, u, alpha, lambda, log = FALSE)
inla.pc.qcor1(p, u, alpha, lambda)
inla.pc.pcor1(q, u, alpha, lambda)
```

Arguments

n	Number of observations
u	The upper limit (see Details)
alpha	The probability going above the upper limit (see Details)
lambda	The rate parameter (see Details)
cor	Vector of correlations
log	Logical. Return the density in natural or log-scale.
p	Vector of probabilities
q	Vector of quantiles

Details

The statement $\text{Prob}(\text{cor} > u) = \text{alpha}$ is used to determine lambda unless lambda is given. Either lambda must be given, or u AND alpha.

Value

inla.pc.dcor1 gives the density, inla.pc.pcor1 gives the distribution function, inla.pc.qcor1 gives the quantile function, and inla.pc.rcor1 generates random deviates.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

inla.doc("pc.rho1")

Examples

```
cor = inla.pc.rcor1(100, lambda = 1)
d = inla.pc.dcor1(cor, lambda = 1)
cor = inla.pc.qcor1(c(0.3, 0.7), u = 0.5, alpha=0.75)
inla.pc.pcor1(cor, u = 0.5, alpha=0.75)
```

pc.cormat

Utility functions for the PC prior for a correlation matrix

Description

Functions to evaluate and sample from the PC prior for a correlation matrix.

Usage

```

inla.pc.cormat.dim2p(dim)
inla.pc.cormat.p2dim(p)
inla.pc.cormat.theta2R(theta)
inla.pc.cormat.R2theta(R)
inla.pc.cormat.r2R(r)
inla.pc.cormat.R2r(R)
inla.pc.cormat.r2theta(r)
inla.pc.cormat.theta2r(theta)
inla.pc.cormat.permute(R)
inla.pc.cormat.rtheta(n=1, p, lambda = 1)
inla.pc.cormat.dtheta(theta, lambda = 1, log = FALSE)

```

Arguments

dim	The dimension of theta, the parameterisation of the correlation matrix
p	The dimension the correlation matrix
theta	A vector of parameters for the correlation matrix
r	The off diagonal elements of a correlation matrix
R	A correlation matrix
n	Number of observations
lambda	The rate parameter in the prior
log	Logical. Return the density in natural or log-scale.

Details

The parameterisation of a correlation matrix of dimension p has dim parameters: θ which are in the interval $-\pi$ to π . The alternative parameterisation is through the off-diagonal elements r of the correlation matrix R . The functions `inla.pc.cormat.<A>2` convert between parameterisations $\langle A \rangle$ to parameterisations $\langle B \rangle$, where both $\langle A \rangle$ and $\langle B \rangle$ are one of θ , r and R , and p and dim .

Value

`inla.pc.cormat.rtheta` generate samples from the prior, returning a matrix where each row is a sample of θ . `inla.pc.cormat.dtheta` evaluates the density of θ . `inla.pc.cormat.permute` randomly permutes a correlation matrix, which is useful if an exchangeable sample of a correlation matrix is required.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```

p = 4
print(paste("theta has length", inla.pc.cormat.p2dim(p)))
theta = inla.pc.cormat.rtheta(n=1, p=4, lambda = 1)
print("sample theta:")
print(theta)
print(paste("log.dens", inla.pc.cormat.dtheta(theta, log=TRUE)))

```

```

print("r:")
r = inla.pc.cormat.theta2r(theta)
print(r)
print("A sample from the non-exchangable prior, R:")
R = inla.pc.cormat.r2R(r)
print(R)
print("A sample from the exchangable prior, R:")
R = inla.pc.cormat.permute(R)
print(R)

```

pc.ddof

*PC-prior for dof in a standardized Student-t***Description**

A function to evaluate the PC-prior for the degrees of freedom in a standardized Student-t distribution

Usage

```
inla.pc.ddof(dof, lambda, u, alpha, log=FALSE)
```

Arguments

dof	Degrees of freedom
log	Logical. Return the density or the log-density
lambda	The optional value of lambda, instead of defining it implicitly through u and alpha
u	The upper value of dof used to elicitate lambda, $\text{Prob}(\text{dof} < u) = \alpha$
alpha	The probability alpha used to elicitate lambda

Details

These functions implements the PC-prior for the dof in a standardized Student-t distribution (ie. with unit variance and $\text{dof} > 2$). Either lambda, or u AND alpha must be given. Due the internal tabulation, dof must be larger than 2.0025.

Value

inla.pc.ddof returns the prior density for given dof.

Author(s)

Havard Rue <hrue@r-inla.org>

`pc.gamma`*Utility functions for the PC prior for Gamma(1/a, 1/a)*

Description

Functions to evaluate, sample, compute quantiles and percentiles of the PC prior for Gamma(1/a, 1/a)

Usage

```
inla.pc.rgamma(n, lambda = 1)
inla.pc.dgamma(x, lambda = 1, log = FALSE)
inla.pc.qgamma(p, lambda = 1)
inla.pc.pgamma(q, lambda = 1)
```

Arguments

<code>n</code>	Number of observations
<code>lambda</code>	The rate parameter (see Details)
<code>x</code>	Evaluation points
<code>log</code>	Logical. Return the density in natural or log-scale.
<code>p</code>	Vector of probabilities
<code>q</code>	Vector of quantiles

Details

This gives the PC prior for the Gamma(1/a, 1/a) case, where a=0 is the base model.

Value

`inla.pc.dgamma` gives the density, `inla.pc.pgamma` gives the distribution function, `inla.pc.qgamma` gives the quantile function, and `inla.pc.rgamma` generates random deviates.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

`inla.doc("pc.gamma")`

Examples

```
x = inla.pc.rgamma(100, lambda = 1)
d = inla.pc.dgamma(x, lambda = 1)
x = inla.pc.qgamma(0.5, lambda = 1)
inla.pc.pgamma(x, lambda = 1)
```

`pc.gammacount`*Utility functions for the PC prior for the gammacount likelihood*

Description

Functions to evaluate, sample, compute quantiles and percentiles of the PC prior for the gammacount likelihood

Usage

```
inla.pc.rgammacount(n, lambda = 1)
inla.pc.dgammacount(x, lambda = 1, log = FALSE)
inla.pc.qgammacount(p, lambda = 1)
inla.pc.pgammacount(q, lambda = 1)
```

Arguments

<code>n</code>	Number of observations
<code>lambda</code>	The rate parameter (see Details)
<code>x</code>	Evaluation points
<code>log</code>	Logical. Return the density in natural or log-scale.
<code>p</code>	Vector of probabilities
<code>q</code>	Vector of quantiles

Details

This gives the PC prior for the gammacount likelihood, which is the PC prior for a in $\text{Gamma}(a, 1)$ where $\text{Gamma}(1, 1)$ is the base model.

Value

`inla.pc.dgammacount` gives the density, `inla.pc.pgammacount` gives the distribution function, `inla.pc.qgammacount` gives the quantile function, and `inla.pc.rgammacount` generates random deviates.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

`inla.doc("pc.gammacount")`

Examples

```
x = inla.pc.rgammacount(100, lambda = 1)
d = inla.pc.dgammacount(x, lambda = 1)
x = inla.pc.qgammacount(0.5, lambda = 1)
inla.pc.pgammacount(x, lambda = 1)
```

pc.gevtail	<i>Utility functions for the PC prior for the tail parameter in the GEV likelihood</i>
------------	--

Description

Functions to evaluate, sample, compute quantiles and percentiles of the PC prior for the tail parameter in the GEV likelihood

Usage

```
inla.pc.rgevtail(n, lambda = 7)
inla.pc.dgevtail(xi, lambda = 7, log = FALSE)
inla.pc.qgevtail(p, lambda = 7)
inla.pc.pgevtail(q, lambda = 7)
```

Arguments

n	Number of observations
lambda	The rate parameter in the PC-prior
xi	Vector of evaluation points, where $1 > xi > 0$.
log	Logical. Return the density in natural or log-scale.
p	Vector of probabilities
q	Vector of quantiles

Details

This gives the PC prior for the tail parameter for the GEV likelihood, where $xi=0$ is the base model.

Value

inla.pc.dgevtail gives the density, inla.pc.pgevtail gives the distribution function, inla.pc.qgevtail gives the quantile function, and inla.pc.rgevtail generates random deviates.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

inla.doc("pc.gevtail")

Examples

```
xi = inla.pc.rgevtail(100, lambda = 7)
d = inla.pc.dgevtail(xi, lambda = 7)
xi = inla.pc.qgevtail(0.5, lambda = 7)
inla.pc.pgevtail(xi, lambda = 7)
```

pc.multvar *Multivariate PC priors*

Description

Functions to evaluate and simulate from multivariate PC priors: The simplex and sphere case

Usage

```
inla.pc.multvar.h.default(x, inverse = FALSE, derivative = FALSE)
inla.pc.multvar.simplex.r(n = NULL, lambda = 1, h = inla.pc.multvar.h.default, b = NULL)
inla.pc.multvar.simplex.d(x = NULL, lambda = 1, log = FALSE, h = inla.pc.multvar.h.default, b = NULL)
inla.pc.multvar.sphere.r(n = NULL, lambda = 1, h = inla.pc.multvar.h.default, H = NULL)
inla.pc.multvar.sphere.d(x = NULL, lambda = 1, log = FALSE, h = inla.pc.multvar.h.default, H = NULL)
```

Arguments

x	Samples to evaluate. If input is a matrix then each row is a sample. If input is a vector then this is the sample.
inverse	Compute the inverse of the h()-function.
derivative	Compute the derivative of the h()-function. (derivative of the inverse function is not used).
n	Number of samples to generate.
lambda	The lambda-parameter in the PC-prior.
log	Evaluate the density in log-scale or ordinary scale.
h	The h()-function, defaults to <code>inla.pc.multvar.h.default</code> . See that code for an example of how to write a user-specific function.
b	The b-vector (gradient) in the expression for the simplex option, $d(x_i) = h(b^T x_i)$
H	The H(essian)-matrix in the expression for the sphere option, $d(x_i) = h(1/2 * x_i^T H x_i)$. If H is a vector, then it is interpreted as the diagonal of a (sparse) diagonal matrix.

Details

These functions implements multivariate PC-priors of the simplex and sphere type.

Value

`inla.pc.multvar.simplex.r` generate samples from the simplex case, and `inla.pc.multvar.simplex.d` evaluate the density. `inla.pc.multvar.sphere.r` generate samples from the sphere case, and `inla.pc.multvar.sphere.d` evaluate the density. `inla.pc.multvar.h.default` implements the default h()-function and illustrate how to code your own spesific one, if needed.

Author(s)

Havard Rue <hrue@r-inla.org>

pc.prec

Utility functions for the PC prior for the precision

Description

Functions to evaluate, sample, compute quantiles and percentiles of the PC prior for the precision in the Gaussian distribution.

Usage

```
inla.pc.rprec(n, u, alpha, lambda)
inla.pc.dprec(prec, u, alpha, lambda, log = FALSE)
inla.pc.qprec(p, u, alpha, lambda)
inla.pc.pprec(q, u, alpha, lambda)
```

Arguments

n	Number of observations
u	The upper limit (see Details)
alpha	The probability going above the upper limit (see Details)
lambda	The rate parameter (see Details)
prec	Vector of precisions
log	Logical. Return the density in natural or log-scale.
p	Vector of probabilities
q	Vector of quantiles

Details

The statement $\text{Prob}(1/\sqrt{\text{prec}} > u) = \alpha$ is used to determine lambda unless lambda is given. Either lambda must be given, or u AND alpha.

Value

inla.pc.dprec gives the density, inla.pc.pprec gives the distribution function, inla.pc.qprec gives the quantile function, and inla.pc.rprec generates random deviates.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

inla.doc("pc.prec")

Examples

```
prec = inla.pc.rprec(100, lambda = 1)
d = inla.pc.dprec(prec, lambda = 1)
prec = inla.pc.qprec(0.5, u = 1, alpha=0.01)
inla.pc.pprec(prec, u = 1, alpha=0.01)
```

pc.sn	<i>Utility functions for the PC prior for skewness in the skew-normal linkfunction and likelihood</i>
-------	---

Description

Functions to evaluate, sample, compute quantiles and percentiles of the PC prior for the skewness in the skew-normal link-function and likelihood

Usage

```
inla.pc.rsn(n, lambda = 40)
inla.pc.dsn(skew, lambda = 40, log = FALSE)
inla.pc.qsn(p, lambda = 40)
inla.pc.psn(q, lambda = 40)
```

Arguments

n	number of observations
lambda	the rate parameter in the PC prior
skew	vector of evaluation points
log	logical. return the density in natural or log-scale.
p	vector of probabilities
q	vector of quantiles

Details

Defines the PC prior for the skewness for the skew-normal linkfunction and likelihood, where skew=0 is the base model. The skewness range from -0.99527... to 0.99527.... ca.

Value

inla.pc.dsn gives the density, inla.pc.psn gives the distribution function, inla.pc.qsn gives the quantile function, and inla.pc.rsn generates random deviates.

Author(s)

havard rue <hrue@r-inla.org>

See Also

inla.doc("pc.sn")

Examples

```
x = inla.pc.rsn(100, lambda = 40)
d = inla.pc.dsn(x, lambda = 40)
x = inla.pc.qsn(0.5, lambda = 40)
inla.pc.psn(x, lambda = 40)
```

plot.inla

*Default INLA plotting***Description**

Takes an inla object produced by inla and plot the results

Usage

```
## S3 method for class 'inla'
plot(x,
      plot.fixed.effects = TRUE,
      plot.lincomb = TRUE,
      plot.random.effects = TRUE,
      plot.hyperparameters = TRUE,
      plot.predictor = TRUE,
      plot.q = TRUE,
      plot.cpo = TRUE,
      plot.prior = FALSE,
      single = FALSE,
      postscript = FALSE,
      pdf = FALSE,
      prefix = "inla.plots/figure-",
      intern = FALSE,
      debug = FALSE,
      cex = 1.75,
      ...)
```

Arguments

x	A fitted inla object produced by inla
plot.fixed.effects	Boolean indicating if posterior marginals for the fixed effects in the model should be plotted
plot.lincomb	Boolean indicating if posterior marginals for the linear combinations should be plotted
plot.random.effects	Boolean indicating if posterior mean and quantiles for the random effects in the model should be plotted
plot.hyperparameters	Boolean indicating if posterior marginals for the hyperparameters in the model should be plotted
plot.predictor	Boolean indicating if posterior mean and quantiles for the linear predictor in the model should be plotted
plot.q	Boolean indicating if precision matrix should be displayed
plot.cpo	Boolean indicating if CPO/PIT values should be plotted
plot.prior	Plot also the prior density for the hyperparameters

single	Boolean indicating if there should be more than one plot per page (FALSE) or just one (TRUE)
postscript	Boolean indicating if postscript files should be produced instead
pdf	Boolean indicating if PDF files should be produced instead
prefix	The prefix for the created files. Additional numbering and suffix is added.
intern	Plot also the hyperparameters in its internal scale.
debug	Write some debug information
cex	The cex parameter in par(). If set to NULL or 0, then default values will be used for graphics parameters
...	Additional arguments to postscript(), pdf() or dev.new().

Value

The return value is a list of the files created (if any).

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla](#)

Examples

```
## Not run:
result = inla(...)
plot(result)
plot(result, single = TRUE, plot.prior = TRUE)
plot(result, single = TRUE, pdf = TRUE, paper = "a4")

## End(Not run)
```

plot.inla.CRS

Plot CRS and inla.CRS objects

Description

Plot the outline of a CRS or inla.CRS projection, with optional graticules (transformed parallels and meridians) and Tissot indicatrices.

Usage

```
## S3 method for class 'inla.CRS'
plot(
  x,
  xlim = NULL,
  ylim = NULL,
  outline = TRUE,
```

```

    graticule = c(15, 15, 45),
    tissot = c(30, 30, 30),
    asp = 1,
    add = FALSE,
    eps = 0.05,
    ...
)

## S3 method for class 'CRS'
plot(
  x,
  xlim = NULL,
  ylim = NULL,
  outline = TRUE,
  graticule = c(15, 15, 45),
  tissot = c(30, 30, 30),
  asp = 1,
  add = FALSE,
  eps = 0.05,
  ...
)

```

Arguments

x	A CRS or inla.CRS() object.
xlim	Optional x-axis limits.
ylim	Optional y-axis limits.
outline	Logical, if TRUE, draw the outline of the projection.
graticule	Vector of length at most 3, to plot meridians with spacing <code>graticule[1]</code> degrees and parallels with spacing <code>graticule[2]</code> degrees. <code>graticule[3]</code> optionally specifies the spacing above and below the first and last parallel. When <code>graticule[1]==0</code> no meridians are drawn, and when <code>graticule[2]==0</code> no parallels are drawn. Use <code>graticule=NULL</code> to skip drawing a graticule.
tissot	Vector of length at most 3, to plot Tissot's indicatrices with spacing <code>tissot[1]</code> degrees and parallels with spacing <code>tissot[2]</code> degrees. <code>tissot[3]</code> specifies a scaling factor. Use <code>tissot=NULL</code> to skip drawing a Tissot's indicatrices.
asp	The aspect ratio for the plot, default 1.
add	If TRUE, add the projecton plot to an existing plot.
eps	Clipping tolerance for rudimentary boundary clipping
...	Additional arguments passed on to the internal calls to <code>plot</code> and <code>lines</code> .

Author(s)

Finn Lindgren finn.lindgren@gmail.com

See Also

[inla.CRS\(\)](#)

Examples

```

if (require(rgdal)) {
  oblique <- c(0, 45, 45, 0)
  for (projtype in c("longlat", "lambert", "mollweide", "hammer")) {
    plot(inla.CRS(projtype), main = projtype)
    plot(inla.CRS(projtype, oblique = oblique), main = paste("oblique", projtype))
  }
}

```

plot.inla.mesh

Draw a triangulation mesh object

Description

Plots an `inla.mesh()` object using either standard graphics or with `rgl`.

Usage

```

## S3 method for class 'inla.mesh'
plot(
  x,
  col = "white",
  t.sub = 1:nrow(mesh$graph$tv),
  add = FALSE,
  lwd = 1,
  xlim = range(mesh$loc[, 1]),
  ylim = range(mesh$loc[, 2]),
  main = NULL,
  rgl = FALSE,
  size = 2,
  draw.vertices = FALSE,
  vertex.color = "black",
  draw.edges = TRUE,
  edge.color = rgb(0.3, 0.3, 0.3),
  draw.segments = draw.edges,
  ...
)

```

Arguments

<code>x</code>	An <code>inla.mesh()</code> object.
<code>col</code>	Color specification. A single named color, a vector of scalar values, or a matrix of RGB values. Requires <code>rgl=TRUE</code> .
<code>t.sub</code>	Optional triangle index subset to be drawn.
<code>add</code>	If <code>TRUE</code> , adds to the current plot instead of starting a new one.
<code>lwd</code>	Line width for triangle edges.
<code>xlim</code>	X-axis limits.
<code>ylim</code>	Y-axis limits.

main	The main plot title. If not specified, a default title is generated based on the mesh type.
rgl	When TRUE, generates an rgl plot instead of a generic graphics plot. Allows 3D plotting and color surface plotting.
size	Size of vertex points in rgl plotting. See rgl.material.
draw.vertices	If TRUE, draw triangle vertices.
vertex.color	Color specification for all vertices.
draw.edges	If TRUE, draw triangle edges.
edge.color	Color specification for all edges.
draw.segments	If TRUE, draw boundary and interior constraint edges more prominently.
...	Further graphics parameters, interpreted by the respective plotting systems.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[plot.inla.trimesh\(\)](#)

Examples

```
mesh <- inla.mesh.create(globe = 10)
plot(mesh)

if (require(rgl)) {
  plot(mesh, rgl = TRUE, col = mesh$loc[, 1])
}
```

plot.inla.trimesh *Low level triangulation mesh plotting*

Description

Plots a triangulation mesh using rgl.

Usage

```
## S3 method for class 'inla.trimesh'
plot(
  x,
  S,
  color = NULL,
  color.axis = NULL,
  color.n = 512,
  color.palette = cm.colors,
  color.truncate = FALSE,
  alpha = NULL,
```

```

    lwd = 1,
    specular = "black",
    draw.vertices = TRUE,
    draw.edges = TRUE,
    edge.color = rgb(0.3, 0.3, 0.3),
    ...
  )

```

Arguments

<code>x</code>	A 3-column triangle-to-vertex index map matrix.
<code>S</code>	A 3-column vertex coordinate matrix.
<code>color</code>	Color specification. A single named color, a vector of scalar values, or a matrix of RGB values.
<code>color.axis</code>	The min/max limit values for the color mapping.
<code>color.n</code>	The number of colors to use in the color palette.
<code>color.palette</code>	A color palette function.
<code>color.truncate</code>	If TRUE, truncate the colors at the color axis limits.
<code>alpha</code>	Transparency/opaqueness values. See <code>rgl.material</code> .
<code>lwd</code>	Line width for edges. See <code>rgl.material</code> .
<code>specular</code>	Specular color. See <code>rgl.material</code> .
<code>draw.vertices</code>	If TRUE, draw triangle vertices.
<code>draw.edges</code>	If TRUE, draw triangle edges.
<code>edge.color</code>	Edge color specification.
<code>...</code>	Additional parameters passed to and from other methods.

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

See Also

[plot.inla.mesh\(\)](#)

PRborder

The PRborder data

Description

A data matrix with Longitude and Latitude coordinates for the boundary of Parana State.

Format

Longitude The Longitude coordinate

Latitude The Latitude coordinate

See Also

PRprec

`print.inla` *Print a INLA fit*

Description

Print a INLA fit

Usage

```
## S3 method for class 'inla'  
print(x, digits = 3L, ...)
```

Arguments

<code>x</code>	An inla-object (output from an inla -call).
<code>digits</code>	Number of digits to print
<code>...</code>	other arguments.

Details

None

Value

None

Author(s)

Havard Rue

See Also

[inla](#)

Examples

```
## None
```

`PRprec` *The PRprec data*

Description

A data frame with daily rainfall in the Parana State.

Format

A data frame TODO

Altitude TODO

Latitude TODO

Longitude TODO

d0101 Daily rainfall at day "mmdd"

d0102 Daily rainfall at day "mmdd"

d0103 Daily rainfall at day "mmdd"

d0104 Daily rainfall at day "mmdd"

d0105 Daily rainfall at day "mmdd"

d0106 Daily rainfall at day "mmdd"

d0107 Daily rainfall at day "mmdd"

d0108 Daily rainfall at day "mmdd"

d0109 Daily rainfall at day "mmdd"

d0110 Daily rainfall at day "mmdd"

d0111 Daily rainfall at day "mmdd"

d0112 Daily rainfall at day "mmdd"

d0113 Daily rainfall at day "mmdd"

d0114 Daily rainfall at day "mmdd"

d0115 Daily rainfall at day "mmdd"

d0116 Daily rainfall at day "mmdd"

d0117 Daily rainfall at day "mmdd"

d0118 Daily rainfall at day "mmdd"

d0119 Daily rainfall at day "mmdd"

d0120 Daily rainfall at day "mmdd"

d0121 Daily rainfall at day "mmdd"

d0122 Daily rainfall at day "mmdd"

d0123 Daily rainfall at day "mmdd"

d0124 Daily rainfall at day "mmdd"

d0125 Daily rainfall at day "mmdd"

d0126 Daily rainfall at day "mmdd"

d0127 Daily rainfall at day "mmdd"

d0128 Daily rainfall at day "mmdd"

d0129 Daily rainfall at day "mmdd"

d0130 Daily rainfall at day "mmdd"

d0131 Daily rainfall at day "mmdd"

d0201 Daily rainfall at day "mmdd"

d0202 Daily rainfall at day "mmdd"

d0203 Daily rainfall at day "mmdd"

d0204 Daily rainfall at day "mmdd"

d0205 Daily rainfall at day "mmdd"
d0206 Daily rainfall at day "mmdd"
d0207 Daily rainfall at day "mmdd"
d0208 Daily rainfall at day "mmdd"
d0209 Daily rainfall at day "mmdd"
d0210 Daily rainfall at day "mmdd"
d0211 Daily rainfall at day "mmdd"
d0212 Daily rainfall at day "mmdd"
d0213 Daily rainfall at day "mmdd"
d0214 Daily rainfall at day "mmdd"
d0215 Daily rainfall at day "mmdd"
d0216 Daily rainfall at day "mmdd"
d0217 Daily rainfall at day "mmdd"
d0218 Daily rainfall at day "mmdd"
d0219 Daily rainfall at day "mmdd"
d0220 Daily rainfall at day "mmdd"
d0221 Daily rainfall at day "mmdd"
d0222 Daily rainfall at day "mmdd"
d0223 Daily rainfall at day "mmdd"
d0224 Daily rainfall at day "mmdd"
d0225 Daily rainfall at day "mmdd"
d0226 Daily rainfall at day "mmdd"
d0227 Daily rainfall at day "mmdd"
d0228 Daily rainfall at day "mmdd"
d0301 Daily rainfall at day "mmdd"
d0302 Daily rainfall at day "mmdd"
d0303 Daily rainfall at day "mmdd"
d0304 Daily rainfall at day "mmdd"
d0305 Daily rainfall at day "mmdd"
d0306 Daily rainfall at day "mmdd"
d0307 Daily rainfall at day "mmdd"
d0308 Daily rainfall at day "mmdd"
d0309 Daily rainfall at day "mmdd"
d0310 Daily rainfall at day "mmdd"
d0311 Daily rainfall at day "mmdd"
d0312 Daily rainfall at day "mmdd"
d0313 Daily rainfall at day "mmdd"
d0314 Daily rainfall at day "mmdd"
d0315 Daily rainfall at day "mmdd"
d0316 Daily rainfall at day "mmdd"

d0317 Daily rainfall at day "mmdd"
d0318 Daily rainfall at day "mmdd"
d0319 Daily rainfall at day "mmdd"
d0320 Daily rainfall at day "mmdd"
d0321 Daily rainfall at day "mmdd"
d0322 Daily rainfall at day "mmdd"
d0323 Daily rainfall at day "mmdd"
d0324 Daily rainfall at day "mmdd"
d0325 Daily rainfall at day "mmdd"
d0326 Daily rainfall at day "mmdd"
d0327 Daily rainfall at day "mmdd"
d0328 Daily rainfall at day "mmdd"
d0329 Daily rainfall at day "mmdd"
d0330 Daily rainfall at day "mmdd"
d0331 Daily rainfall at day "mmdd"
d0401 Daily rainfall at day "mmdd"
d0402 Daily rainfall at day "mmdd"
d0403 Daily rainfall at day "mmdd"
d0404 Daily rainfall at day "mmdd"
d0405 Daily rainfall at day "mmdd"
d0406 Daily rainfall at day "mmdd"
d0407 Daily rainfall at day "mmdd"
d0408 Daily rainfall at day "mmdd"
d0409 Daily rainfall at day "mmdd"
d0410 Daily rainfall at day "mmdd"
d0411 Daily rainfall at day "mmdd"
d0412 Daily rainfall at day "mmdd"
d0413 Daily rainfall at day "mmdd"
d0414 Daily rainfall at day "mmdd"
d0415 Daily rainfall at day "mmdd"
d0416 Daily rainfall at day "mmdd"
d0417 Daily rainfall at day "mmdd"
d0418 Daily rainfall at day "mmdd"
d0419 Daily rainfall at day "mmdd"
d0420 Daily rainfall at day "mmdd"
d0421 Daily rainfall at day "mmdd"
d0422 Daily rainfall at day "mmdd"
d0423 Daily rainfall at day "mmdd"
d0424 Daily rainfall at day "mmdd"
d0425 Daily rainfall at day "mmdd"

d0426 Daily rainfall at day "mmdd"
d0427 Daily rainfall at day "mmdd"
d0428 Daily rainfall at day "mmdd"
d0429 Daily rainfall at day "mmdd"
d0430 Daily rainfall at day "mmdd"
d0501 Daily rainfall at day "mmdd"
d0502 Daily rainfall at day "mmdd"
d0503 Daily rainfall at day "mmdd"
d0504 Daily rainfall at day "mmdd"
d0505 Daily rainfall at day "mmdd"
d0506 Daily rainfall at day "mmdd"
d0507 Daily rainfall at day "mmdd"
d0508 Daily rainfall at day "mmdd"
d0509 Daily rainfall at day "mmdd"
d0510 Daily rainfall at day "mmdd"
d0511 Daily rainfall at day "mmdd"
d0512 Daily rainfall at day "mmdd"
d0513 Daily rainfall at day "mmdd"
d0514 Daily rainfall at day "mmdd"
d0515 Daily rainfall at day "mmdd"
d0516 Daily rainfall at day "mmdd"
d0517 Daily rainfall at day "mmdd"
d0518 Daily rainfall at day "mmdd"
d0519 Daily rainfall at day "mmdd"
d0520 Daily rainfall at day "mmdd"
d0521 Daily rainfall at day "mmdd"
d0522 Daily rainfall at day "mmdd"
d0523 Daily rainfall at day "mmdd"
d0524 Daily rainfall at day "mmdd"
d0525 Daily rainfall at day "mmdd"
d0526 Daily rainfall at day "mmdd"
d0527 Daily rainfall at day "mmdd"
d0528 Daily rainfall at day "mmdd"
d0529 Daily rainfall at day "mmdd"
d0530 Daily rainfall at day "mmdd"
d0531 Daily rainfall at day "mmdd"
d0601 Daily rainfall at day "mmdd"
d0602 Daily rainfall at day "mmdd"
d0603 Daily rainfall at day "mmdd"
d0604 Daily rainfall at day "mmdd"

d0605 Daily rainfall at day "mmdd"
d0606 Daily rainfall at day "mmdd"
d0607 Daily rainfall at day "mmdd"
d0608 Daily rainfall at day "mmdd"
d0609 Daily rainfall at day "mmdd"
d0610 Daily rainfall at day "mmdd"
d0611 Daily rainfall at day "mmdd"
d0612 Daily rainfall at day "mmdd"
d0613 Daily rainfall at day "mmdd"
d0614 Daily rainfall at day "mmdd"
d0615 Daily rainfall at day "mmdd"
d0616 Daily rainfall at day "mmdd"
d0617 Daily rainfall at day "mmdd"
d0618 Daily rainfall at day "mmdd"
d0619 Daily rainfall at day "mmdd"
d0620 Daily rainfall at day "mmdd"
d0621 Daily rainfall at day "mmdd"
d0622 Daily rainfall at day "mmdd"
d0623 Daily rainfall at day "mmdd"
d0624 Daily rainfall at day "mmdd"
d0625 Daily rainfall at day "mmdd"
d0626 Daily rainfall at day "mmdd"
d0627 Daily rainfall at day "mmdd"
d0628 Daily rainfall at day "mmdd"
d0629 Daily rainfall at day "mmdd"
d0630 Daily rainfall at day "mmdd"
d0701 Daily rainfall at day "mmdd"
d0702 Daily rainfall at day "mmdd"
d0703 Daily rainfall at day "mmdd"
d0704 Daily rainfall at day "mmdd"
d0705 Daily rainfall at day "mmdd"
d0706 Daily rainfall at day "mmdd"
d0707 Daily rainfall at day "mmdd"
d0708 Daily rainfall at day "mmdd"
d0709 Daily rainfall at day "mmdd"
d0710 Daily rainfall at day "mmdd"
d0711 Daily rainfall at day "mmdd"
d0712 Daily rainfall at day "mmdd"
d0713 Daily rainfall at day "mmdd"
d0714 Daily rainfall at day "mmdd"

d0715 Daily rainfall at day "mmdd"
d0716 Daily rainfall at day "mmdd"
d0717 Daily rainfall at day "mmdd"
d0718 Daily rainfall at day "mmdd"
d0719 Daily rainfall at day "mmdd"
d0720 Daily rainfall at day "mmdd"
d0721 Daily rainfall at day "mmdd"
d0722 Daily rainfall at day "mmdd"
d0723 Daily rainfall at day "mmdd"
d0724 Daily rainfall at day "mmdd"
d0725 Daily rainfall at day "mmdd"
d0726 Daily rainfall at day "mmdd"
d0727 Daily rainfall at day "mmdd"
d0728 Daily rainfall at day "mmdd"
d0729 Daily rainfall at day "mmdd"
d0730 Daily rainfall at day "mmdd"
d0731 Daily rainfall at day "mmdd"
d0801 Daily rainfall at day "mmdd"
d0802 Daily rainfall at day "mmdd"
d0803 Daily rainfall at day "mmdd"
d0804 Daily rainfall at day "mmdd"
d0805 Daily rainfall at day "mmdd"
d0806 Daily rainfall at day "mmdd"
d0807 Daily rainfall at day "mmdd"
d0808 Daily rainfall at day "mmdd"
d0809 Daily rainfall at day "mmdd"
d0810 Daily rainfall at day "mmdd"
d0811 Daily rainfall at day "mmdd"
d0812 Daily rainfall at day "mmdd"
d0813 Daily rainfall at day "mmdd"
d0814 Daily rainfall at day "mmdd"
d0815 Daily rainfall at day "mmdd"
d0816 Daily rainfall at day "mmdd"
d0817 Daily rainfall at day "mmdd"
d0818 Daily rainfall at day "mmdd"
d0819 Daily rainfall at day "mmdd"
d0820 Daily rainfall at day "mmdd"
d0821 Daily rainfall at day "mmdd"
d0822 Daily rainfall at day "mmdd"
d0823 Daily rainfall at day "mmdd"

d0824 Daily rainfall at day "mmdd"
d0825 Daily rainfall at day "mmdd"
d0826 Daily rainfall at day "mmdd"
d0827 Daily rainfall at day "mmdd"
d0828 Daily rainfall at day "mmdd"
d0829 Daily rainfall at day "mmdd"
d0830 Daily rainfall at day "mmdd"
d0831 Daily rainfall at day "mmdd"
d0901 Daily rainfall at day "mmdd"
d0902 Daily rainfall at day "mmdd"
d0903 Daily rainfall at day "mmdd"
d0904 Daily rainfall at day "mmdd"
d0905 Daily rainfall at day "mmdd"
d0906 Daily rainfall at day "mmdd"
d0907 Daily rainfall at day "mmdd"
d0908 Daily rainfall at day "mmdd"
d0909 Daily rainfall at day "mmdd"
d0910 Daily rainfall at day "mmdd"
d0911 Daily rainfall at day "mmdd"
d0912 Daily rainfall at day "mmdd"
d0913 Daily rainfall at day "mmdd"
d0914 Daily rainfall at day "mmdd"
d0915 Daily rainfall at day "mmdd"
d0916 Daily rainfall at day "mmdd"
d0917 Daily rainfall at day "mmdd"
d0918 Daily rainfall at day "mmdd"
d0919 Daily rainfall at day "mmdd"
d0920 Daily rainfall at day "mmdd"
d0921 Daily rainfall at day "mmdd"
d0922 Daily rainfall at day "mmdd"
d0923 Daily rainfall at day "mmdd"
d0924 Daily rainfall at day "mmdd"
d0925 Daily rainfall at day "mmdd"
d0926 Daily rainfall at day "mmdd"
d0927 Daily rainfall at day "mmdd"
d0928 Daily rainfall at day "mmdd"
d0929 Daily rainfall at day "mmdd"
d0930 Daily rainfall at day "mmdd"
d1001 Daily rainfall at day "mmdd"
d1002 Daily rainfall at day "mmdd"

d1003 Daily rainfall at day "mmdd"
d1004 Daily rainfall at day "mmdd"
d1005 Daily rainfall at day "mmdd"
d1006 Daily rainfall at day "mmdd"
d1007 Daily rainfall at day "mmdd"
d1008 Daily rainfall at day "mmdd"
d1009 Daily rainfall at day "mmdd"
d1010 Daily rainfall at day "mmdd"
d1011 Daily rainfall at day "mmdd"
d1012 Daily rainfall at day "mmdd"
d1013 Daily rainfall at day "mmdd"
d1014 Daily rainfall at day "mmdd"
d1015 Daily rainfall at day "mmdd"
d1016 Daily rainfall at day "mmdd"
d1017 Daily rainfall at day "mmdd"
d1018 Daily rainfall at day "mmdd"
d1019 Daily rainfall at day "mmdd"
d1020 Daily rainfall at day "mmdd"
d1021 Daily rainfall at day "mmdd"
d1022 Daily rainfall at day "mmdd"
d1023 Daily rainfall at day "mmdd"
d1024 Daily rainfall at day "mmdd"
d1025 Daily rainfall at day "mmdd"
d1026 Daily rainfall at day "mmdd"
d1027 Daily rainfall at day "mmdd"
d1028 Daily rainfall at day "mmdd"
d1029 Daily rainfall at day "mmdd"
d1030 Daily rainfall at day "mmdd"
d1031 Daily rainfall at day "mmdd"
d1101 Daily rainfall at day "mmdd"
d1102 Daily rainfall at day "mmdd"
d1103 Daily rainfall at day "mmdd"
d1104 Daily rainfall at day "mmdd"
d1105 Daily rainfall at day "mmdd"
d1106 Daily rainfall at day "mmdd"
d1107 Daily rainfall at day "mmdd"
d1108 Daily rainfall at day "mmdd"
d1109 Daily rainfall at day "mmdd"
d1110 Daily rainfall at day "mmdd"
d1111 Daily rainfall at day "mmdd"

d1112 Daily rainfall at day "mmdd"
d1113 Daily rainfall at day "mmdd"
d1114 Daily rainfall at day "mmdd"
d1115 Daily rainfall at day "mmdd"
d1116 Daily rainfall at day "mmdd"
d1117 Daily rainfall at day "mmdd"
d1118 Daily rainfall at day "mmdd"
d1119 Daily rainfall at day "mmdd"
d1120 Daily rainfall at day "mmdd"
d1121 Daily rainfall at day "mmdd"
d1122 Daily rainfall at day "mmdd"
d1123 Daily rainfall at day "mmdd"
d1124 Daily rainfall at day "mmdd"
d1125 Daily rainfall at day "mmdd"
d1126 Daily rainfall at day "mmdd"
d1127 Daily rainfall at day "mmdd"
d1128 Daily rainfall at day "mmdd"
d1129 Daily rainfall at day "mmdd"
d1130 Daily rainfall at day "mmdd"
d1201 Daily rainfall at day "mmdd"
d1202 Daily rainfall at day "mmdd"
d1203 Daily rainfall at day "mmdd"
d1204 Daily rainfall at day "mmdd"
d1205 Daily rainfall at day "mmdd"
d1206 Daily rainfall at day "mmdd"
d1207 Daily rainfall at day "mmdd"
d1208 Daily rainfall at day "mmdd"
d1209 Daily rainfall at day "mmdd"
d1210 Daily rainfall at day "mmdd"
d1211 Daily rainfall at day "mmdd"
d1212 Daily rainfall at day "mmdd"
d1213 Daily rainfall at day "mmdd"
d1214 Daily rainfall at day "mmdd"
d1215 Daily rainfall at day "mmdd"
d1216 Daily rainfall at day "mmdd"
d1217 Daily rainfall at day "mmdd"
d1218 Daily rainfall at day "mmdd"
d1219 Daily rainfall at day "mmdd"
d1220 Daily rainfall at day "mmdd"
d1221 Daily rainfall at day "mmdd"

d1222 Daily rainfall at day "mmdd"
d1223 Daily rainfall at day "mmdd"
d1224 Daily rainfall at day "mmdd"
d1225 Daily rainfall at day "mmdd"
d1226 Daily rainfall at day "mmdd"
d1227 Daily rainfall at day "mmdd"
d1228 Daily rainfall at day "mmdd"
d1229 Daily rainfall at day "mmdd"
d1230 Daily rainfall at day "mmdd"
d1231 Daily rainfall at day "mmdd"

See Also

PRborder

qinv

Computes (parts of) the inverse of a SPD sparse matrix

Description

This routine use the GMRFLib implementation which compute parts of the inverse of a SPD sparse matrix. The diagonal and values for the neighbours in the inverse, are provided.

Usage

```
inla.qinv(Q, constr, reordering = INLA::inla.reorderings(), num.threads = NULL)
```

Arguments

Q	A SPD matrix, either as a (dense) matrix or sparseMatrix.
constr	Optional linear constraints; see ?INLA::f and argument extraconstr
reordering	The type of reordering algorithm to be used for TAUCS; either one of the names listed in inla.reorderings() or the output from inla.qreordering(Q). The default is "auto" which try several reordering algorithm and use the best one for this particular matrix.
num.threads	Maximum number of threads the inla-program will use, or as 'A:B' defining the number threads in the outer (A) and inner (B) layer for nested parallelism.

Value

inla.qinv returns a sparseMatrix of type dgTMatrix with the diagonal and values for the neighbours in the inverse. Note that the full inverse is NOT provided!

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```

## dense matrix example
n = 10
A = matrix(runif(n^2), n, n)
Q = A %*% t(A)
print(mean(abs(inla.qinv(Q) - solve(Q))))

## sparse matrix example
rho = 0.9
Q = toeplitz(c(1+rho^2, -rho, rep(0, n-3), -rho)) / (1-rho^2)
Q = inla.as.dgTMatrix(Q)
Q.inv = inla.qinv(Q)

## compute the marginal variances as a vector from a precision matrix
marginal.variances = diag(inla.qinv(Q))

## read the sparse matrix from a file in the 'i, j, value' format
filename = tempfile()
write(t(cbind(Q@i+1L, Q@j+1L, Q@x)), ncol=3, file=filename)
Qinv = inla.qinv(filename)
unlink(filename)

```

qreordering

*Compute the reordering using the GMRFLib implementation***Description**

This function compute the reordering (or find the best reordering) using the GMRFLib implementation

Usage

```
inla.qreordering(graph, reordering = inla.reorderings())
```

Arguments

graph	A (inla-)graph object
reordering	The name of the reordering algorithm to be used; either one of the names listed in <code>inla.reorderings()</code> . The default is "auto" which try several reordering algorithm and use the best one for this particular matrix.

Value

`inla.qreordering` returns a list with the name of the reordering algorithm used or found, the reordering code for the reordering algorithm, the actual reordering and its inverse.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
g = system.file("demodata/germany.graph", package="INLA")
r = inla.qreordering(g)
m = inla.graph2matrix(g)
r = inla.qreordering(m)
```

qsample*Generate samples from a GMRF using the GMRFLib implementation*

Description

This function generate samples from a GMRF using the GMRFLib implementation

Usage

```
inla.qsample(
  n = 1L,
  Q,
  b,
  mu,
  sample,
  constr,
  reordering = INLA::inla.reorderings(),
  seed = 0L,
  logdens = ifelse(missing(sample), FALSE, TRUE),
  compute.mean = ifelse(missing(sample), FALSE, TRUE),
  num.threads = if (seed == 0L) "0:0" else NULL,
  selection = NULL, verbose = FALSE)
```

Arguments

n	Number of samples. Only used if <code>missing(sample)</code>
Q	The precision matrix or a filename containing it.
b	The linear term
mu	The mu term
sample	A matrix of optional samples where each column is a sample. If set, then evaluate the log-density for each sample only.
constr	Optional linear constraints; see <code>?INLA::f</code> and argument <code>extraconstr</code>
reordering	The type of reordering algorithm to be used for TAUCS; either one of the names listed in <code>inla.reorderings()</code> or the output from <code>inla.qreordering(Q)</code> . The default is "auto" which try several reordering algorithm and use the best one for this particular matrix.
seed	Control the RNG. If <code>seed=0L</code> then GMRFLib will set the seed intelligently/at 'random', and this is and should be the default behaviour. If <code>seed < 0L</code> then the saved state of the RNG will be reused if possible, otherwise, GMRFLib will set the seed intelligently/at 'random'. If <code>seed > 0L</code> then this value is used as the seed for the RNG.

PLEASE NOTE1: If `seed!=0` then the computations will run in serial mode, over-riding whatever is set in `num.threads` (a warning might be issued).

PLEASE NOTE2: If the PARDISO sparse matrix library is used, continuity of the samples with respect to small changes in the precision matrix, can be expected but is not guaranteed. If this feature is required, please use the TAUCS sparse matrix library.

<code>logdens</code>	If TRUE, compute also the log-density of each sample. Note that the output format then change.
<code>compute.mean</code>	If TRUE, compute also the (constrained) mean. Note that the output format then change.
<code>num.threads</code>	Maximum number of threads the <code>inla</code> -program will use, or as 'A:B' defining the number threads in the outer (A) and inner (B) layer for nested parallelism. <code>seed!=0</code> requires serial computations.
<code>selection</code>	A vector of indices of each sample to return. NULL means return the whole sample. (Note that the log-density returned, is for the whole sample.) The use of <code>selection</code> cannot be combined with the use of <code>sample</code> .
<code>verbose</code>	Logical. Run in verbose mode or not.

Value

The log-density has form $-1/2(x-\mu)^T Q (x-\mu) + b^T x$

If `logdens` is FALSE, then `inla.qsample` returns the samples in a matrix, where each column is a sample. If `logdens` or `compute.mean` is TRUE, then a list with names `sample`, `logdens` and `mean` is returned. The samples are stored in the matrix `sample` where each column is a sample, and the log densities of each sample are stored as the vector `logdens`. The mean (include corrections for the constraints, if any) is store in the vector `mean`.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
g = system.file("demodata/germany.graph", package="INLA")
Q = inla.graph2matrix(g)
diag(Q) = dim(Q)[1L]
x = inla.qsample(10, Q)
## Not run: matplot(x)
x = inla.qsample(10, Q, logdens=TRUE)
## Not run: matplot(x$sample)

n = 3
Q = diag(n)
ns = 2

## sample and evaluate a sample
x = inla.qsample(n, Q=Q, logdens=TRUE)
xx = inla.qsample(Q=Q, sample = x$sample)
print(x$logdens - xx$logdens)

## the use of a constraint
constr = list(A = matrix(rep(1, n), 1, n), e = 0)
x = inla.qsample(n, Q=Q, constr=constr)
```

```

print(constr$A %*% x)

## control the RNG (require serial mode)
x = inla.qsample(n, Q=Q, seed = 123, num.threads="1:1")
## restart from same seed, only sample 1
xx = inla.qsample(n=1, Q=Q, seed = 123, num.threads="1:1")
## continue from the save state, sample the remaining 2
xxx = inla.qsample(n=n-1, Q=Q, seed = -1, num.threads="1:1")
## should be 0
print(x - cbind(xx, xxx))

```

qsolve

Solves linear SPD systems

Description

This routine use the GMRFLib implementation to solve linear systems with a SPD matrix.

Usage

```
inla.qsolve(Q, B, reordering = inla.reorderings(), method = c("solve", "forward", "backward"))
```

Arguments

Q	A SPD matrix, either as a (dense) matrix or sparse-matrix.
B	The right hand side matrix, either as a (dense) matrix or sparse-matrix.
reordering	The type of reordering algorithm to be used for TAUCS; either one of the names listed in <code>inla.reorderings()</code> or the output from <code>inla.qreordering(Q)</code> . The default is "auto" which try several reordering algorithm and use the best one for this particular matrix (using the TAUCS library).
method	The system to solve, one of "solve", "forward" or "backward". Let $Q = L L^T$, where L is lower triangular (the Cholesky triangle), then <code>method="solve"</code> solves $L L^T X = B$ or equivalently $Q X = B$, <code>method="forward"</code> solves $L X = B$, and <code>method="backward"</code> solves $L^T X = B$.

Value

`inla.qsolve` returns a matrix X, which is the solution of $Q X = B$, $L X = B$ or $L^T X = B$ depending on the value of `method`.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```

n = 10
nb <- n-1
QQ = matrix(rnorm(n^2), n, n)
QQ <- QQ %%% t(QQ)

Q = inla.as.sparse(QQ)
B = matrix(rnorm(n*nb), n, nb)

X = inla.qsolve(Q, B, method = "solve")
XX = inla.qsolve(Q, B, method = "solve", reordering = inla.qreordering(Q))
print(paste("err solve1", sum(abs( Q %%% X - B))))
print(paste("err solve2", sum(abs( Q %%% XX - B))))

## the forward and backward solve is tricky, as after permutation and with Q=LL', then L is
## lower triangular, but L in the original ordering is not lower triangular. if the rhs is iid
## noise, this is not important. to control the reordering, then the 'taucs' library must be
## used.
inla.setOption(smtp = 'taucs')

## case 1. use the matrix as is, no reordering
r <- "identity"
L = t(chol(Q))
X = inla.qsolve(Q, B, method = "forward", reordering = r)
XX = inla.qsolve(Q, B, method = "backward", reordering = r)
print(paste("err forward ", sum(abs(L %%% X - B))))
print(paste("err backward", sum(abs(t(L) %%% XX - B))))

## case 2. use a reordering from the library
r <- inla.qreordering(Q)
im <- r$ireordering
m <- r$reordering
print(cbind(idx = 1:n, m, im) )
Qr <- Q[im, im]
L = t(chol(Qr))[m, m]

X = inla.qsolve(Q, B, method = "forward", reordering = r)
XX = inla.qsolve(Q, B, method = "backward", reordering = r)
print(paste("err forward ", sum(abs( L %%% X - B))))
print(paste("err backward", sum(abs( t(L) %%% XX - B))))

```

read.graph

*Read and write a graph-object***Description**

Construct a graph-object from a file or a matrix; write graph-object to file

Usage

```

inla.read.graph(..., size.only = FALSE)
inla.write.graph(graph, filename = "graph.dat", mode = c("binary", "ascii"), ...)

## S3 method for class 'inla.graph'

```

```
summary(object, ...)
## S3 method for class 'inla.graph'
plot(x, y, ...)
## S3 method for class 'inla.graph.summary'
print(x, ...)
```

Arguments

filename	The filename of the graph.
graph	An <code>inla.graph</code> -object, a (sparse) symmetric matrix, a filename containing the graph, a list or collection of characters and/or numbers defining the graph, or a neighbours list with class <code>nb</code> (see <code>spdep::card</code> and <code>spdep::poly2nb</code> for details of <code>nb</code> and an example a function returning an <code>nb</code> object)
mode	The mode of the file; <code>ascii-file</code> or a (gzip-compressed) binary.
object	An <code>inla.graph</code> -object
x	An <code>inla.graph</code> -object
y	Not used
size.only	Only read the size of the graph
...	Additional arguments. In <code>inla.read.graph</code> , then it is the graph definition (object, matrix, character, filename), plus extra arguments. In <code>inla.write.graph</code> it is extra arguments to <code>inla.read.graph</code> .

Value

The output of `inla.read.graph`, is an `inla.graph` object, with elements

n	is the size of the graph
nnbs	is a vector with the number of neighbours
nbs	is a list-list with the neighbours
cc	list with connected component information <ul style="list-style-type: none"> • <code>idis</code> is a vector with the connected component id for each node (starting from 1) • <code>nis</code> is the number of connected components • <code>nodesis</code> is a list-list of nodes belonging to each connected component

Methods implemented for `inla.graph` are `summary` and `plot`. The method `plot` require the libraries `Rgraphviz` and `graph` from the Bioconductor-project, see <https://www.bioconductor.org>.

Author(s)

Havard Rue <hrue@r-inla.org>

See Also

[inla.spy](#)

Examples

```

## a graph from a file
cat("3 1 1 2 2 1 1 3 0\n", file="g.dat")
g = inla.read.graph("g.dat")
## writing an inla.graph-object to file
g.file = inla.write.graph(g, mode="binary")
## re-reading it from that file
gg = inla.read.graph(g.file)
summary(g)
##
Not run:
plot(g)
inla.spy(g)
## when defining the graph directly in the call,
## we can use a mix of character and numbers
g = inla.read.graph(c(3, 1, "1 2 2 1 1 3", 0))
inla.spy(c(3, 1, "1 2 2 1 1 3 0"))
inla.spy(c(3, 1, "1 2 2 1 1 3 0"), reordering=3:1)
inla.write.graph(c(3, 1, "1 2 2 1 1 3 0"))

## building a graph from adjacency matrix
adjacent = matrix(0, nrow = 4, ncol = 4)
adjacent[1,4] = adjacent[4,1] = 1
adjacent[2,4] = adjacent[4,2] = 1
adjacent[2,3] = adjacent[3,2] = 1
adjacent[3,4] = adjacent[4,3] = 1
g = inla.read.graph(adjacent)
plot(g)
summary(g)
End(Not run)

```

rgeneric.define

rgeneric models

Description

A framework for defining latent models in R

Usage

```

inla.rgeneric.define(model = NULL, debug = FALSE, compile = TRUE, optimize = FALSE, ...)
inla.rgeneric.iid.model(
  cmd = c("graph", "Q", "mu", "initial", "log.norm.const", "log.prior", "quit"),
  theta = NULL)
inla.rgeneric.ar1.model(
  cmd = c("graph", "Q", "mu", "initial", "log.norm.const", "log.prior", "quit"),
  theta = NULL)
inla.rgeneric.ar1.model.opt(
  cmd = c("graph", "Q", "mu", "initial", "log.norm.const", "log.prior", "quit"),
  theta = NULL)
inla.rgeneric.wrapper(
  cmd = c("graph", "Q", "mu", "initial", "log.norm.const", "log.prior", "quit"),
  model, theta = NULL)

```

```
inla.rgeneric.q(
  rmodel,
  cmd = c("graph", "Q", "mu", "initial", "log.norm.const", "log.prior", "quit"),
  theta = NULL)
```

Arguments

<code>model</code>	The definition of the model; see <code>inla.rgeneric.ar1.model</code>
<code>rmodel</code>	The rgeneric model-object, the output of <code>inla.rgeneric.define</code>
<code>debug</code>	Logical. Turn on/off debugging
<code>compile</code>	Logical. Compile the definition of the model or not.
<code>optimize</code>	Logical. With this option TRUE, then <code>model</code> pass only the values of Q and not the whole matrix. Please see the vignette for details and <code>inla.rgeneric.ar1.model.opt</code> for an example.
<code>cmd</code>	An allowed request
<code>theta</code>	Values of theta
<code>...</code>	Named list of variables that defines the environment of <code>model</code>
<code>debug</code>	Logical. Enable debug output

Value

This allows a latent model to be defined in R. See `inla.rgeneric.ar1.model` and `inla.rgeneric.iid.model` and the documentation for worked out examples of how to define latent models in this way. This will be somewhat slow and is intended for special cases and prototyping. The function `inla.rgeneric.wrapper` is for internal use only.

Author(s)

Havard Rue <hrue@r-inla.org>

Salm

Extra-Poisson variation in dose-response study

Description

Breslow (1984) analyses some mutagenicity assay data (shown below) on salmonella in which three plates have been processed at each dose i of quinoline and the number of revertant colonies of TA98 Salmonella measured

Format

A data frame with 18 observations on the following 3 variables.

y number of salmonella bacteria
dose dose of quinoline (mg per plate)
rand indicator

Source

WinBUGS/OpenBUGS manual Examples VOI.I

Examples

```
data(Salm)
```

```
scale.model
```

```
Scale an intrinsic GMRF model
```

Description

This function scales an intrinsic GMRF model so the geometric mean of the marginal variances is one

Usage

```
inla.scale.model(Q, constr = NULL, eps = sqrt(.Machine$double.eps))
```

Arguments

Q	A SPD matrix, either as a (dense) matrix or sparseMatrix
constr	Linear constraints spanning the null-space of Q; see ?INLA::f and argument extraconstr
eps	A small constant added to the diagonal of Q if constr

Value

inla.scale.model returns a sparseMatrix of type dgTMatrix scaled so the geometric mean of the marginal variances (of the possible non-singular part of Q) is one, for each connected component of the matrix.

Author(s)

Havard Rue <hrue@r-inla.org>

Examples

```
## Q is singular
data(Germany)
g = system.file("demodata/germany.graph", package="INLA")
Q = -inla.graph2matrix(g)
diag(Q) = 0
diag(Q) = -rowSums(Q)
n = dim(Q)[1]
Q.scaled = inla.scale.model(Q, constr = list(A = matrix(1, 1, n), e=0))
print(diag(MASS::ginv(Q.scaled)))

## Q is singular with 3 connected components
g = inla.read.graph("6 1 2 2 3 2 2 1 3 3 2 1 2 4 1 5 5 1 4 6 0")
print(paste("Number of connected components", g$cc$n))
Q = -inla.graph2matrix(g)
diag(Q) = 0
diag(Q) = -rowSums(Q)
n = dim(Q)[1]
```

```

Q.scaled = inla.scale.model(Q, constr = list(A = matrix(1, 1, n), e=0))
print(diag(MASS::ginv(Q.scaled)))

## Q is non-singular with 3 connected components. no constraints needed
diag(Q) = diag(Q) + 1
Q.scaled = inla.scale.model(Q)
print(diag(MASS::ginv(Q.scaled)))

```

 Scotland

Conditional Autoregressive (CAR) model for disease mapping

Description

The rate of lip cancer in 56 counties in Scotland is recorder. The data set includes the observed and expected cases (based on the population and its age and sex distribution in the country), a covariate measuring the percentage of the population engaged in agriculture, fishing or forestry and the "position" of each county expressed as a list of adjacent counties

Format

A data frame with 56 observations on the following 4 variables.

Counts The number of lip cancer registered

E The expected number of lip cancer

X The percentage of the population engaged in agriculture, fishing or forestry

Region The county

Source

OpenBUGS Example manual, GeoBUGS

References

Clayton and Kaldor (1987) and Breslow and Clayton (1993)

Examples

```
data(Scotland)
```

Seeds

Factorial design

Description

Proportion of seeds that germinated on each of 21 plates arranged according to a 2 by 2 factorial layout by seed and type of root extract

Format

A data frame with 21 observations on the following 5 variables.

r number of germinated seeds per plate

n number of total seeds per plate

x1 seed type

x2 root extracted

plate indicator for the plate

Source

WinBUGS/OpenBUGS Manual Example, Vol. I

Examples

```
data(Seeds)
```

SPDEtoy

toy simulated data set for the SPDE tutorial

Description

Simulated data set on 200 location points. The simulation process is made at the introduction of the SPDE tutorial.

Format

A data frame with 200 observations on the following 3 variables.

s1 First element of the coordinates

s2 Second element of the coordinates

y data simulated at the locations

Source

SPDE tutorial

Examples

```
data(SPDEtoy)
```

`summary.inla`*Summary for a INLA fit*

Description

Takes a fitted `inla` or `surv.inla` object produced by `inla` or `surv.inla` and produces a summary from it.

Usage

```
## S3 method for class 'inla'  
summary(object, digits = 3L, include.lincomb = TRUE, ...)  
## S3 method for class 'summary.inla'  
print(x, digits = 3L, ...)
```

Arguments

<code>object</code>	a fitted <code>inla</code> object as produced by <code>inla</code> .
<code>x</code>	a <code>summary.inla</code> object produced by <code>summary.inla</code>
<code>digits</code>	Integer Number of digits
<code>include.lincomb</code>	Logical Include the summary for the the linear combinations or not
<code>...</code>	other arguments.

Details

Posterior mean and standard deviation (together with quantiles or cdf) are printed for the fixed effects in the model.

For the random effects the function `summary()` prints the posterior mean and standard deviations for the hyperparameters

If the option `short.summary` is set to `TRUE` using `inla.setOption`, then a less verbose summary variant will be used, which might be more suitable for Markdown documents.

Value

`summary.inla` returns an object of class `summary.inla`, a list of components to print.

Author(s)

Sara Martino and Havard Rue

See Also

[inla](#)

summary.inla.mesh *Summarizing triangular mesh objects*

Description

Construct and print inla.mesh object summaries

Usage

```
## S3 method for class 'inla.mesh'
summary(object, verbose = FALSE, ...)

## S3 method for class 'summary.inla.mesh'
print(x, ...)
```

Arguments

object	an object of class "inla.mesh", usually a result of a call to inla.mesh.create() or inla.mesh.2d() .
verbose	If TRUE, produce a more detailed output.
...	further arguments passed to or from other methods.
x	an object of class "summary.inla.mesh", usually a result of a call to summary.inla.mesh() .

Author(s)

Finn Lindgren <finn.lindgren@gmail.com>

Surg *Surgical: Institutional ranking*

Description

This example considers mortality rates in 12 hospitals performing cardiac surgery in babies

Format

A data frame with 12 observations on the following 3 variables.

n Number of deaths

r Total number of cases

hospital a factor with levels A B C D E F G H I J K L

Source

WinBUGS/OpenBUGS Manual Examples Vol. I

Examples

```
data(Surg)
```

SurvSim

Survival data

Description

Simulated data set for Weibull survival model

Format

A data frame with 100 observations on the following 3 variables.

y a numeric vector of survival times

cens a numeric vector of event indicator (0=censored 1=failure)

x a numeric vector of covariate

Tokyo

Binomial time series

Description

Recorded days of rain above 1 mm in Tokyo for 2 years, 1983:84

Format

A data frame with 366 observations on the following 3 variables.

y number of days with rain

n total number of days

time day of the year

Source

<http://www.math.ntnu.no/~hrue/GMRF-book/tokyo.rainfall.data.dat>

References

Rue, H and Held, L. (2005) *Gaussian Markov Random Fields - Theory and Applications* Chapman and Hall

Examples

`data(Tokyo)`

Zambia

Semiparametric regression

Description

Undernutrition of children in each region of Zambia is measured through a score computed on the basis of some anthropometric measures. The data set contains also other information about each child.

Format

A data frame with 4847 observations on the following 10 variables.

hazstd standardised Z score of stunting

bmi body mass index of the mother

age age of the child in months

district district where the child lives

rcw mother employment status with categories "working" (1) and "not working" (-1)

edu1 mother's education status with categories "complete primary but incomplete secondary" (edu1=1), "complete secondary or higher" (edu2=1) and "no education or incomplete primary" (edu1=edu2=-1)

edu2 see above

tpr locality of the domicile with categories "urban" (1) and "rural" (-1)

sex gender of the child with categories "male" (1) and "female" (-1)

edu DO NOT KNOW; check source

Source

BayesX Manual <http://www.stat.uni-muenchen.de/~bayesx/bayesx.html>

Examples

```
data(Zambia)
```

Index

- * **Survival models**
 - inla.surv, 275
- * **datasets**
 - BivMetaAnalysis, 10
 - Cancer, 11
 - Drivers, 30
 - Epil, 30
 - Germany, 38
 - Kidney, 282
 - Leuk, 284
 - Munich, 291
 - nwEngland, 292
 - Oral, 292
 - PRborder, 311
 - PRprec, 312
 - Salm, 330
 - Scotland, 332
 - Seeds, 333
 - SPDEtoy, 333
 - Surg, 335
 - SurvSim, 336
 - Tokyo, 336
 - Zambia, 337
- * **fmesher**
 - inla.stack.remove.unused, 270
- * **graph**
 - geobugs2inla, 37
- * **plot**
 - plot.inla, 306
- 1djmarginal (joint.marginal), 280
- agaussian (aggregate.gaussian), 8
- aggregate.gaussian, 8
- ar.dpacf (pc.ar), 295
- ar.pacf2acf (inla.ar.pacf2phi), 45
- ar.pacf2phi (inla.ar.pacf2phi), 45
- ar.phi2acf (inla.ar.pacf2phi), 45
- ar.phi2pacf (inla.ar.pacf2phi), 45
- ar.rpacf (pc.ar), 295
- as.inla.mdata (inla.mdata), 74
- as.inla.mesh.segment, 9
- as.inla.surv (inla.surv), 275
- barrier (inla.barrier.pcmatern), 49
- barrier (inla.barrier), 48
- binary.install (inla.binary.install), 50
- BivMetaAnalysis, 10
- Cancer, 11
- cbind.data.frames (inla.coxph), 53
- changelog (inla.changelog), 51
- collect.results (inla.collect.results), 51
- compare.results (inla.compare.results), 52
- contour(), 95
- control.bgev.default, 11, 12, 13, 15–18, 21–27
- control.compute, 12, 12, 13, 15–18, 21–27
- control.expert, 12, 13, 13, 15–18, 21–27
- control.family, 12, 13, 14, 15–18, 21–27
- control.fixed, 12, 13, 15, 15, 16–18, 21–27
- control.group, 12, 13, 15, 16, 16, 17, 18, 21–27
- control.hazard, 12, 13, 15–17, 17, 18, 21–27
- control.inla, 12, 13, 15–18, 18, 21–27
- control.lincomb, 12, 13, 15–18, 21, 21, 22–27
- control.link, 12, 13, 15–18, 21, 22, 22, 23–27
- control.mix, 12, 13, 15–18, 21–23, 23, 24–27
- control.mode, 12, 13, 15–18, 21–24, 24, 25–27
- control.pardiso.default, 12, 13, 15–18, 21–24, 24, 25–27
- control.predictor, 12, 13, 15–18, 21–25, 25, 26, 27
- control.results, 26, 27
- control.update, 12, 13, 15–18, 21–27, 27
- cormat.dim2p (pc.cormat), 297
- cormat.dtheta (pc.cormat), 297
- cormat.p2dim (pc.cormat), 297
- cormat.permute (pc.cormat), 297
- cormat.R2r (pc.cormat), 297
- cormat.r2R (pc.cormat), 297
- cormat.R2theta (pc.cormat), 297
- cormat.r2theta (pc.cormat), 297
- cormat.rtheta (pc.cormat), 297

- cormat.theta2R (pc.cormat), 297
- cormat.theta2r (pc.cormat), 297
- coxph (inla.coxph), 53
- cpo.inla (inla.cpo), 54
- crs_wkt (inla.wkt_is_geocent), 278
- crs_wkt(), 57
- cut, 28

- debug.graph, 29
- dev.new, 58
- dmarginal (marginal), 288
- Drivers, 30

- emarginal (marginal), 288
- Epil, 30
- extract.groups, 31

- f, 32, 41, 45, 64, 68, 69
- fgn, 36

- geobugs2inla, 37
- Germany, 38
- get.inlaEnv (inla.get.inlaEnv), 63
- graph2matrix, 38

- hpdmarginal (marginal), 288
- hyperpar.inla, 36, 37
- hyperpar.inla (inla.hyperpar), 65
- hyperpar.sample (inla.hyperpar.sample), 66
- hyperpar.sampler (inla.hyperpar.sample), 66

- idx, 39
- INLA (INLA-package), 7
- inla, 12, 13, 15–18, 21–27, 36, 37, 40, 45, 51, 55, 66, 69, 75, 240–242, 275, 276, 280, 283, 287, 289, 307, 312, 334
- inla(), 254
- inla-class, 45
- INLA-package, 7
- inla.1djmarginal (joint.marginal), 280
- inla.agaussian (aggregate.gaussian), 8
- inla.ar.pacf2acf (inla.ar.pacf2phi), 45
- inla.ar.pacf2phi, 45
- inla.ar.phi2acf (inla.ar.pacf2phi), 45
- inla.ar.phi2pacf (inla.ar.pacf2phi), 45
- inla.as.CRS.list (inla.CRSargs), 57
- inla.as.CRSargs.list (inla.CRSargs), 57
- inla.as.dgTMatrix (inla.as.sparse), 46
- inla.as.list.CRS (inla.CRSargs), 57
- inla.as.list.CRSargs (inla.CRSargs), 57
- inla.as.sparse, 46
- inla.as.wkt.wkt_tree (inla.as.wkt_tree.wkt), 47
- inla.as.wkt_tree.wkt, 47
- inla.barrier, 48
- inla.barrier (inla.barrier.pcmatern), 49
- inla.barrier.pcmatern, 49
- inla.binary.install, 50
- inla.changelog, 51
- inla.changes (inla.changelog), 51
- inla.collect.results, 51
- inla.compare.results, 52
- inla.contour.segment (inla.mesh.segment), 94
- inla.control.bgev.default (control.bgev.default), 11
- inla.control.compute (control.compute), 12
- inla.control.expert (control.expert), 13
- inla.control.family (control.family), 14
- inla.control.fixed (control.fixed), 15
- inla.control.group (control.group), 16
- inla.control.hazard (control.hazard), 17
- inla.control.inla (control.inla), 18
- inla.control.lincomb (control.lincomb), 21
- inla.control.link (control.link), 22
- inla.control.mix (control.mix), 23
- inla.control.mode (control.mode), 24
- inla.control.pardiso.default (control.pardiso.default), 24
- inla.control.predictor (control.predictor), 25
- inla.control.results (control.results), 26
- inla.control.update (control.update), 27
- inla.coxph, 53
- inla.cpo, 54
- inla.CRS, 55
- inla.CRS(), 58, 67, 268, 308
- inla.crs_get_ellipsoid_radius (inla.wkt_is_geocent), 278
- inla.crs_get_lengthunit (inla.wkt_is_geocent), 278
- inla.crs_get_wkt (inla.wkt_is_geocent), 278
- inla.crs_get_wkt(), 57
- inla.crs_is_geocent (inla.wkt_is_geocent), 278
- inla.crs_set_ellipsoid_radius (inla.wkt_is_geocent), 278
- inla.crs_set_lengthunit (inla.wkt_is_geocent), 278

- inla.CRSargs, 57
- inla.cut (cut), 28
- inla.debug.graph (debug.graph), 29
- inla.delaunay (inla.mesh.create), 85
- inla.delaunay(), 79
- inla.dev.new, 58
- inla.diameter, 59
- inla.dmarginal (marginal), 288
- inla.doc, 33, 41, 60, 72
- inla.emarginal (marginal), 288
- inla.extract.el, 60
- inla.fallback_PROJ6 (inla.has_PROJ6), 64
- inla.fgn (fgn), 36
- inla.fmesher.smorg, 61
- inla.generate.colors, 62
- inla.geobugs2inla (geobugs2inla), 37
- inla.get.inlaEnv, 63
- inla.getOption (inla.option), 235
- inla.graph (read.graph), 327
- inla.graph2matrix (graph2matrix), 38
- inla.group, 63
- inla.has_PROJ6, 64
- inla.hpdmarginal (marginal), 288
- inla.hyperpar, 45, 65, 289
- inla.hyperpar.sample, 66
- inla.hyperpar.sampler
(inla.hyperpar.sample), 66
- inla.identical.CRS, 67
- inla.identical.CRS(), 57
- inla.idx (idx), 39
- inla.inla.doc (inla.doc), 60
- inla.joint.marginal (joint.marginal),
280
- inla.jp.define (jp.define), 281
- inla.knmodels, 68, 71
- inla.knmodels.sample, 69, 70
- inla.ks.plot, 71
- inla.lattice2node (lattice2node), 283
- inla.link (link), 286
- inla.list.models, 32, 72
- inla.load, 73
- inla.make.lincomb (make.lincomb), 287
- inla.make.lincombs (make.lincomb), 287
- inla.marginal (marginal), 288
- inla.matern.cov, 73
- inla.matrix2vector (lattice2node), 283
- inla.mdata, 74
- inla.merge, 75
- inla.mesh (inla.mesh.create), 85
- inla.mesh(), 81, 82, 87–89, 91, 92, 94, 237,
248, 257, 260, 263, 264, 293, 309
- inla.mesh.1d, 77
- inla.mesh.1d(), 77, 78, 81, 82, 87, 88, 92,
248, 260, 262, 264, 266
- inla.mesh.1d.A, 77
- inla.mesh.1d.bary (inla.mesh.1d.A), 77
- inla.mesh.1d.fem (inla.mesh.fem), 88
- inla.mesh.2d, 78
- inla.mesh.2d(), 82, 83, 87, 95, 257, 262,
266, 290, 335
- inla.mesh.assessment, 80
- inla.mesh.basis, 81
- inla.mesh.basis(), 257, 262, 266
- inla.mesh.boundary, 82
- inla.mesh.components, 83, 84
- inla.mesh.create, 85
- inla.mesh.create(), 79, 83, 93–95, 262,
266, 290, 335
- inla.mesh.create.helper(), 83
- inla.mesh.deriv, 87
- inla.mesh.fem, 88
- inla.mesh.interior
(inla.mesh.boundary), 82
- inla.mesh.lattice, 89
- inla.mesh.lattice(), 86, 87, 92, 93
- inla.mesh.map (inla.mesh.map.lim), 90
- inla.mesh.map.lim, 90
- inla.mesh.project, 91
- inla.mesh.project(), 91
- inla.mesh.projector
(inla.mesh.project), 91
- inla.mesh.query, 93
- inla.mesh.query(), 87
- inla.mesh.segment, 94
- inla.mesh.segment(), 9, 10, 31, 79, 83, 86,
87, 93, 235, 285, 286
- inla.mmarginal (marginal), 288
- inla.models, 96
- inla.nmix.fitted
(inla.nmix.lambda.fitted), 232
- inla.nmix.lambda.fitted, 232
- inla.node2lattice (lattice2node), 283
- inla.nonconvex.hull, 234
- inla.nonconvex.hull(), 79
- inla.not_for_PROJ4 (inla.has_PROJ6), 64
- inla.not_for_PROJ6 (inla.has_PROJ6), 64
- inla.option, 235
- inla.options (inla.option), 235
- inla.over_sp_mesh, 237
- inla.pardiso, 238
- inla.pc.alpha (pc.alpha), 294
- inla.pc.ar (pc.ar), 295
- inla.pc.cor0 (pc.cor0), 295
- inla.pc.cor1 (pc.cor1), 296

- inla.pc.cormat (pc.cormat), 297
- inla.pc.dalphaw (pc.alphaw), 294
- inla.pc.dcor0 (pc.cor0), 295
- inla.pc.dcor1 (pc.cor1), 296
- inla.pc.ddof (pc.ddof), 299
- inla.pc.dgamma (pc.gamma), 300
- inla.pc.dgammacount (pc.gammacount), 301
- inla.pc.dgevtail (pc.gevtail), 302
- inla.pc.dof (pc.ddof), 299
- inla.pc.dprec (pc.prec), 304
- inla.pc.dsn (pc.sn), 305
- inla.pc.gamma (pc.gamma), 300
- inla.pc.gammacount (pc.gammacount), 301
- inla.pc.gevtail (pc.gevtail), 302
- inla.pc.multvar (pc.multvar), 303
- inla.pc.palphaw (pc.alphaw), 294
- inla.pc.pcor0 (pc.cor0), 295
- inla.pc.pcor1 (pc.cor1), 296
- inla.pc.pgamma (pc.gamma), 300
- inla.pc.pgammacount (pc.gammacount), 301
- inla.pc.pgevtail (pc.gevtail), 302
- inla.pc.pprec (pc.prec), 304
- inla.pc.prec (pc.prec), 304
- inla.pc.psn (pc.sn), 305
- inla.pc.qalphaw (pc.alphaw), 294
- inla.pc.qcor0 (pc.cor0), 295
- inla.pc.qcor1 (pc.cor1), 296
- inla.pc.qgamma (pc.gamma), 300
- inla.pc.qgammacount (pc.gammacount), 301
- inla.pc.qgevtail (pc.gevtail), 302
- inla.pc.qprec (pc.prec), 304
- inla.pc.qsn (pc.sn), 305
- inla.pc.ralphaw (pc.alphaw), 294
- inla.pc.rcor0 (pc.cor0), 295
- inla.pc.rcor1 (pc.cor1), 296
- inla.pc.rgamma (pc.gamma), 300
- inla.pc.rgammacount (pc.gammacount), 301
- inla.pc.rgevtail (pc.gevtail), 302
- inla.pc.rprec (pc.prec), 304
- inla.pc.rsn (pc.sn), 305
- inla.pc.sn (pc.sn), 305
- inla.pc.t (pc.ddof), 299
- inla.plot (plot.inla), 306
- inla.pmarginal (marginal), 288
- inla.posterior.sample (inla.sample), 243
- inla.priors.used, 238
- inla.prune, 239
- inla.q (inla.qstat), 239
- inla.qdel (inla.qstat), 239
- inla.qget (inla.qstat), 239
- inla.qinv (qinv), 322
- inla.qlog (inla.qstat), 239
- inla.qmarginal (marginal), 288
- inla.qnuke (inla.qstat), 239
- inla.qreordering (qreordering), 323
- inla.qsample (qsample), 324
- inla.qsample(), 256
- inla.qsolve (qsolve), 326
- inla.qstat, 239
- inla.rbind.data.frames (inla.coxph), 53
- inla.read.graph, 39
- inla.read.graph (read.graph), 327
- inla.remote (inla.ssh.copy.id), 270
- inla.reorderings, 241
- inla.requires_PROJ6 (inla.has_PROJ6), 64
- inla.rerun, 241
- inla.rgeneric.ar1.model
(rgeneric.define), 329
- inla.rgeneric.define (rgeneric.define),
329
- inla.rgeneric.iid.model
(rgeneric.define), 329
- inla.rgeneric.q (rgeneric.define), 329
- inla.rgeneric.wrapper
(rgeneric.define), 329
- inla.rjmarginal (joint.marginal), 280
- inla.rmarginal (marginal), 288
- inla.row.kron, 242
- inla.sample, 243
- inla.scale.model (scale.model), 331
- inla.set.control.bgev.default.default
(control.bgev.default), 11
- inla.set.control.compute.default
(control.compute), 12
- inla.set.control.expert.default
(control.expert), 13
- inla.set.control.family.default
(control.family), 14
- inla.set.control.fixed.default
(control.fixed), 15
- inla.set.control.group.default
(control.group), 16
- inla.set.control.hazard.default
(control.hazard), 17
- inla.set.control.inla.default
(control.inla), 18
- inla.set.control.lincomb.default
(control.lincomb), 21
- inla.set.control.link.default
(control.link), 22
- inla.set.control.mix.default
(control.mix), 23
- inla.set.control.mode.default
(control.mode), 24

- inla.set.control.pardiso.default.default
(control.pardiso.default), 24
- inla.set.control.predictor.default
(control.predictor), 25
- inla.set.control.results.default
(control.results), 26
- inla.set.control.update.default
(control.update), 27
- inla.setOption(inla.option), 235
- inla.simplify.curve, 247
- inla.simplify.curve(), 95, 234
- inla.smarginal(marginal), 288
- inla.sp2segment(as.inla.mesh.segment),
9
- inla.sp_get_crs, 269
- inla.sp_get_crs(), 57, 279
- inla.spde.make.A, 248
- inla.spde.make.A(), 242, 249–251, 273
- inla.spde.make.block.A, 249
- inla.spde.make.block.A(), 249
- inla.spde.make.index, 250
- inla.spde.make.index(), 249, 273
- inla.spde.models, 251
- inla.spde.models(), 253, 255
- inla.spde.precision, 252
- inla.spde.precision(), 256
- inla.spde.result, 253
- inla.spde.sample, 256
- inla.spde1(inla.spde1.create), 256
- inla.spde1(), 252
- inla.spde1.create, 256
- inla.spde1.models(inla.spde.models),
251
- inla.spde1.precision
(inla.spde.precision), 252
- inla.spde1.result(inla.spde.result),
253
- inla.spde2(inla.spde2.generic), 258
- inla.spde2(), 252, 257, 259
- inla.spde2.generic, 258
- inla.spde2.generic(), 253, 262, 266
- inla.spde2.matern, 260
- inla.spde2.matern(), 248, 254, 255, 257,
259, 263, 266, 293
- inla.spde2.matern.sd.basis, 263
- inla.spde2.models(inla.spde.models),
251
- inla.spde2.models(), 259
- inla.spde2.pcmatern, 264
- inla.spde2.pcmatern(), 262
- inla.spde2.precision
(inla.spde.precision), 252
- inla.spde2.result(inla.spde.result),
253
- inla.spde2.result(), 251
- inla.spde2.theta2phi0
(inla.spde2.matern), 260
- inla.spde2.theta2phi0(), 253
- inla.spde2.theta2phi1
(inla.spde2.matern), 260
- inla.spde2.theta2phi1(), 253
- inla.spde2.theta2phi2
(inla.spde2.matern), 260
- inla.spde2.theta2phi2(), 253
- inla.spTransform, 267
- inla.spy, 328
- inla.spy(graph2matrix), 38
- inla.ssh.copy.id, 270
- inla.stack(inla.stack.remove.unused),
270
- inla.stack.remove.unused, 270
- inla.surv, 37, 275
- inla.tjmargin(joint.marginal), 280
- inla.tmargin(marginal), 288
- inla.update(inla.upgrade), 276
- inla.upgrade, 276
- inla.vector2matrix(lattice2node), 283
- inla.version, 277
- inla.wkt_get_ellipsoid_radius
(inla.wkt_is_geocent), 278
- inla.wkt_get_lengthunit
(inla.wkt_is_geocent), 278
- inla.wkt_is_geocent, 278
- inla.wkt_predef(inla.CRS), 55
- inla.wkt_set_ellipsoid_radius
(inla.wkt_is_geocent), 278
- inla.wkt_set_lengthunit
(inla.wkt_is_geocent), 278
- inla.wkt_tree_get_item
(inla.as.wkt_tree.wkt), 47
- inla.wkt_tree_set_item
(inla.as.wkt_tree.wkt), 47
- inla.wkt_unit_params
(inla.wkt_is_geocent), 278
- inla.write.graph(read.graph), 327
- inla.zmargin(marginal), 288
- is.inla.mdata(inla.mdata), 74
- is.inla.surv(inla.surv), 275
- joint.marginal, 280
- jp(jp.define), 281
- jp.define, 281
- Kidney, 282
- knmodels(inla.knmodels), 68

- ks.plot (inla.ks.plot), 71
- ks.test, 71
- lattice2node, 283
- Leuk, 284
- Leukemia (Leuk), 284
- lines.inla.mesh.segment, 285
- link, 286
- list.models (inla.list.models), 72
- make.lincomb, 287
- make.lincombs (make.lincomb), 287
- marginal, 288
- matrix2vector (lattice2node), 283
- merge.inla (inla.merge), 75
- meshbuilder, 80, 290
- meshbuilder(), 80
- mmarginal (marginal), 288
- Munich, 291
- NewEngland (nwEngland), 292
- nmix.lambda.fitted
 - (inla.nmix.lambda.fitted), 232
- node2lattice (lattice2node), 283
- nwEngland, 292
- Oral, 292
- pacf2acf (inla.ar.pacf2phi), 45
- pacf2phi (inla.ar.pacf2phi), 45
- param2.matern.orig, 293
- pc.alphaw, 294
- pc.ar, 295
- pc.cor0, 295
- pc.cor1, 296
- pc.cormat, 297
- pc.dalphaw (pc.alphaw), 294
- pc.dcor0 (pc.cor0), 295
- pc.dcor1 (pc.cor1), 296
- pc.ddof, 299
- pc.dgamma (pc.gamma), 300
- pc.dgammacount (pc.gammacount), 301
- pc.dgevtail (pc.gevtail), 302
- pc.dof (pc.ddof), 299
- pc.dprec (pc.prec), 304
- pc.dsn (pc.sn), 305
- pc.gamma, 300
- pc.gammacount, 301
- pc.gevtail, 302
- pc.multvar, 303
- pc.palphaw (pc.alphaw), 294
- pc.pcor0 (pc.cor0), 295
- pc.pcor1 (pc.cor1), 296
- pc.pgamma (pc.gamma), 300
- pc.pgammacount (pc.gammacount), 301
- pc.pgevtail (pc.gevtail), 302
- pc.pprec (pc.prec), 304
- pc.prec, 304
- pc.psn (pc.sn), 305
- pc.qalphaw (pc.alphaw), 294
- pc.qcor0 (pc.cor0), 295
- pc.qcor1 (pc.cor1), 296
- pc.qgamma (pc.gamma), 300
- pc.qgammacount (pc.gammacount), 301
- pc.qgevtail (pc.gevtail), 302
- pc.qprec (pc.prec), 304
- pc.qsn (pc.sn), 305
- pc.ralphaw (pc.alphaw), 294
- pc.rcor0 (pc.cor0), 295
- pc.rcor1 (pc.cor1), 296
- pc.rgamma (pc.gamma), 300
- pc.rgammacount (pc.gammacount), 301
- pc.rgevtail (pc.gevtail), 302
- pc.rprec (pc.prec), 304
- pc.rsn (pc.sn), 305
- pc.sn, 305
- pc.t (pc.ddof), 299
- phi2acf (inla.ar.pacf2phi), 45
- phi2pacf (inla.ar.pacf2phi), 45
- plot.CRS (plot.inla.CRS), 307
- plot.CRS(), 57
- plot.inla, 306
- plot.inla.CRS, 307
- plot.inla.graph (read.graph), 327
- plot.inla.mesh, 309
- plot.inla.mesh(), 311
- plot.inla.surv (inla.surv), 275
- plot.inla.trimesh, 310
- plot.inla.trimesh(), 310
- pmarginal (marginal), 288
- posterior.sample (inla.sample), 243
- PRborder, 311
- print.inla, 312
- print.inla.graph.summary (read.graph), 327
- print.inla.mdata (inla.mdata), 74
- print.inla.q (inla.qstat), 239
- print.inla.surv (inla.surv), 275
- print.summary.inla (summary.inla), 334
- print.summary.inla.mesh
 - (summary.inla.mesh), 335
- priors.used (inla.priors.used), 238
- PRprec, 312
- prune (inla.prune), 239
- qinv, 322

- qmarginal (marginal), 288
- qreordering, 323
- qsample, 324
- qsolve, 326

- read.graph, 327
- reorderings (inla.reorderings), 241
- rerun (inla.rerun), 241
- rgdal::CRSargs(), 58
- rgeneric (rgeneric.define), 329
- rgeneric.define, 329
- rjmarginal (joint.marginal), 280
- rmarginal (marginal), 288

- Salm, 330
- scale.model, 331
- Scotland, 332
- Seeds, 333
- set.control.bgev.default.default
 (control.bgev.default), 11
- set.control.compute.default
 (control.compute), 12
- set.control.expert.default
 (control.expert), 13
- set.control.family.default
 (control.family), 14
- set.control.fixed.default
 (control.fixed), 15
- set.control.group.default
 (control.group), 16
- set.control.hazard.default
 (control.hazard), 17
- set.control.inla.default
 (control.inla), 18
- set.control.lincomb.default
 (control.lincomb), 21
- set.control.link.default
 (control.link), 22
- set.control.mix.default (control.mix),
 23
- set.control.mode.default
 (control.mode), 24
- set.control.pardiso.default.default
 (control.pardiso.default), 24
- set.control.predictor.default
 (control.predictor), 25
- set.control.results.default
 (control.results), 26
- set.control.update.default
 (control.update), 27
- smarginal (marginal), 288
- sp::CRS(), 57
- sp::over(), 237
- sp::SpatialPolygons(), 237
- SPDEtoy, 333
- spy (graph2matrix), 38
- ssh.copy.id (inla.ssh.copy.id), 270
- summary.inla, 334
- summary.inla.graph (read.graph), 327
- summary.inla.mesh, 335
- summary.inla.mesh(), 335
- summary.inla.q (inla.qstat), 239
- summary.surv.inla (summary.inla), 334
- Surg, 335
- SurvSim, 336

- tjmarginal (joint.marginal), 280
- Tokyo, 336

- vector2matrix (lattice2node), 283
- version (inla.version), 277

- write.graph (read.graph), 327

- Zambia, 337
- zmarginal (marginal), 288