

Package ‘rSPDE’

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Type Package

Title Rational Approximations of Fractional Stochastic Partial Differential Equations

Version 2.3.3

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Description Functions that compute rational approximations of fractional elliptic stochastic partial differential equations. The package also contains functions for common statistical usage of these approximations. The main references for rSPDE are Bolin, Simas and Xiong (2023) <doi:10.1080/10618600.2023.2231051> for the covariance-based method and Bolin and Kirchner (2020) <doi:10.1080/10618600.2019.1665537> for the operator-based rational approximation. These can be generated by the citation function in R.

Depends R (>= 3.5.0), Matrix

Imports stats, methods, fmesher, lifecycle, broom

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URL <https://davidbolin.github.io/rSPDE/>

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Additional_repositories <https://inla.r-inla-download.org/R/testing>

BugReports <https://github.com/davidbolin/rSPDE/issues>

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augment.rspde_lme	<i>Augment data with information from a rspde_lme object</i>
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Description

Augment accepts a model object and a dataset and adds information about each observation in the dataset. It includes predicted values in the `.fitted` column, residuals in the `.resid` column, and standard errors for the fitted values in a `.se.fit` column. It also contains the New columns always begin with a `.` prefix to avoid overwriting columns in the original dataset.

Usage

```
## S3 method for class 'rspde_lme'
augment(
  x,
  newdata = NULL,
  loc = NULL,
  mesh = FALSE,
  which_repl = NULL,
  se_fit = FALSE,
  conf_int = FALSE,
  pred_int = FALSE,
  level = 0.95,
  n_samples = 100,
  edge_number = "edge_number",
  distance_on_edge = "distance_on_edge",
  normalized = FALSE,
  ...
)
```

Arguments

`x` A `rspde_lme` object.

newdata	A <code>data.frame</code> or a <code>list</code> containing the covariates, the edge number and the distance on edge for the locations to obtain the prediction. If <code>NULL</code> , the fitted values will be given for the original locations where the model was fitted.
loc	Prediction locations. Can either be a <code>data.frame</code> , a <code>matrix</code> or a character vector, that contains the names of the columns of the coordinates of the locations. For models using <code>metric_graph</code> objects, please use <code>edge_number</code> and <code>distance_on_edge</code> instead.
mesh	Obtain predictions for mesh nodes? The graph must have a mesh, and either <code>only_latent</code> is set to <code>TRUE</code> or the model does not have covariates.
which_repl	Which replicates to obtain the prediction. If <code>NULL</code> predictions will be obtained for all replicates. Default is <code>NULL</code> .
se_fit	Logical indicating whether or not a <code>.se.fit</code> column should be added to the augmented output. If <code>TRUE</code> , it only returns a non-NA value if type of prediction is 'link'.
conf_int	Logical indicating whether or not confidence intervals for the fitted variable should be built.
pred_int	Logical indicating whether or not prediction intervals for future observations should be built.
level	Level of confidence and prediction intervals if they are constructed.
n_samples	Number of samples when computing prediction intervals.
edge_number	Name of the variable that contains the edge number, the default is <code>edge_number</code> .
distance_on_edge	Name of the variable that contains the distance on edge, the default is <code>distance_on_edge</code> .
normalized	Are the distances on edges normalized?
...	Additional arguments.

Value

A `tidyverse::tibble()` with columns:

- `.fitted` Fitted or predicted value.
- `.fittedlwrconf` Lower bound of the confidence interval, if `conf_int = TRUE`
- `.fitteduprconf` Upper bound of the confidence interval, if `conf_int = TRUE`
- `.fittedlwrpred` Lower bound of the prediction interval, if `pred_int = TRUE`
- `.fitteduprpred` Upper bound of the prediction interval, if `pred_int = TRUE`
- `.fixed` Prediction of the fixed effects.
- `.random` Prediction of the random effects.
- `.resid` The ordinary residuals, that is, the difference between observed and fitted values.
- `.se_fit` Standard errors of fitted values, if `se_fit = TRUE`.

See Also

[glance.rspde_lme](#)

`bru_get_mapper.inla_rspde`
rSPDE inlabru mapper

Description

rSPDE inlabru mapper

Usage

```
bru_get_mapper.inla_rspde(model, ...)
ibm_n.bru_mapper_inla_rspde(mapper, ...)
ibm_values.bru_mapper_inla_rspde(mapper, ...)
ibm_jacobian.bru_mapper_inla_rspde(mapper, input, ...)
```

Arguments

<code>model</code>	An <code>inla_rspde</code> for which to construct or extract a mapper
<code>...</code>	Arguments passed on to other methods
<code>mapper</code>	A <code>bru_mapper_inla_rspde</code> object
<code>input</code>	The values for which to produce a mapping matrix

Examples

```
#devel version
if (requireNamespace("INLA", quietly = TRUE) &&
    requireNamespace("inlabru", quietly = TRUE)){
library(INLA)
library(inlabru)

set.seed(123)
m <- 100
loc_2d_mesh <- matrix(runif(m * 2), m, 2)
mesh_2d <- inla.mesh.2d(
  loc = loc_2d_mesh,
  cutoff = 0.05,
  max.edge = c(0.1, 0.5)
)
sigma <- 1
range <- 0.2
nu <- 0.8
kappa <- sqrt(8 * nu) / range
op <- matern.operators(
  mesh = mesh_2d, nu = nu,
  range = range, sigma = sigma, m = 2,
```

```

parameterization = "matern"
)
u <- simulate(op)
A <- inla.spde.make.A(
  mesh = mesh_2d,
  loc = loc_2d_mesh
)
sigma.e <- 0.1
y <- A %*% u + rnorm(m) * sigma.e
y <- as.vector(y)

data_df <- data.frame(y=y, x1 = loc_2d_mesh[,1],
                       x2 = loc_2d_mesh[,2])
coordinates(data_df) <- c("x1", "x2")
rspde_model <- rspde.matern(
  mesh = mesh_2d,
  nu_upper_bound = 2
)
cmp <- y ~ Intercept(1) +
  field(coordinates, model = rspde_model)

rspde_fit <- bru(cmp, data = data_df)
summary(rspde_fit)
}
#devel.tag

```

construct.spde.matern.loglike

Constructor of Matern loglikelihood functions for non-stationary models.

Description

This function evaluates the log-likelihood function for observations of a non-stationary Gaussian process defined as the solution to the SPDE

$$(\kappa(s) - \Delta)^\beta (\tau(s)u(s)) = W.$$

The observations are assumed to be generated as $Y_i = u(s_i) + \epsilon_i$, where ϵ_i are iid mean-zero Gaussian variables. The latent model is approximated using a rational approximation of the fractional SPDE model.

Usage

```
construct.spde.matern.loglike(
  object,
```

```

    Y,
    A,
    sigma.e = NULL,
    mu = 0,
    user_nu = NULL,
    user_m = NULL,
    log_scale = TRUE,
    return_negative_likelihood = TRUE
)

```

Arguments

object	The rational SPDE approximation, computed using matern.operators()
Y	The observations, either a vector or a matrix where the columns correspond to independent replicates of observations.
A	An observation matrix that links the measurement location to the finite element basis.
sigma.e	IF non-null, the standard deviation of the measurement noise will be kept fixed in the returned likelihood.
mu	Expectation vector of the latent field (default = 0).
user_nu	If non-null, the shape parameter will be kept fixed in the returned likelihood.
user_m	If non-null, update the order of the rational approximation, which needs to be a positive integer.
log_scale	Should the parameters be evaluated in log-scale?
return_negative_likelihood	Return minus the likelihood to turn the maximization into a minimization?

Value

The log-likelihood function. The parameters of the returned function are given in the order theta, nu, sigma.e, whenever they are available.

See Also

[matern.operators\(\)](#), [predict.CBrSPDEobj\(\)](#)

Examples

```

# this example illustrates how the function can be used for maximum
# likelihood estimation
# Sample a Gaussian Matern process on R using a rational approximation
set.seed(123)
sigma.e <- 0.1
n.rep <- 10
n.obs <- 100
n.x <- 51
# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = n.x)

```

```

fem <- rSPDE.fem1d(x)
tau <- rep(0.5, n.x)
nu <- 0.8
alpha <- nu + 0.5
kappa <- rep(1, n.x)
# Matern parameterization
# compute rational approximation
op <- spde.matern.operators(
  loc_mesh = x,
  kappa = kappa, tau = tau, alpha = alpha,
  parameterization = "spde", d = 1
)
# Sample the model
u <- simulate(op, n.rep)
# Create some data
obs.loc <- runif(n = n.obs, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
noise <- rnorm(n.obs * n.rep)
dim(noise) <- c(n.obs, n.rep)
Y <- as.matrix(A %*% u + sigma.e * noise)
# define negative likelihood function for optimization using matern.loglike
mlik <- construct.spde.matern.loglike(op, Y, A)
#' #The parameters can now be estimated by minimizing mlik with optim

# Choose some reasonable starting values depending on the size of the domain
theta0 <- log(c( 1 / sqrt(var(c(Y))),sqrt(8), 0.9, 0.01))
# run estimation and display the results
theta <- optim(theta0, mlik)
print(data.frame(
  tau = c(tau[1], exp(theta$par[1])), kappa = c(kappa[1], exp(theta$par[2])),
  nu = c(nu, exp(theta$par[3])), sigma.e = c(sigma.e, exp(theta$par[4])),
  row.names = c("Truth", "Estimates")
))

# SPDE parameterization
# compute rational approximation
op <- spde.matern.operators(
  kappa = kappa, tau = tau, alpha = alpha,
  loc_mesh = x, d = 1,
  parameterization = "spde"
)
# Sample the model
u <- simulate(op, n.rep)
# Create some data
obs.loc <- runif(n = n.obs, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
noise <- rnorm(n.obs * n.rep)
dim(noise) <- c(n.obs, n.rep)
Y <- as.matrix(A %*% u + sigma.e * noise)
# define negative likelihood function for optimization using matern.loglike
mlik <- construct.spde.matern.loglike(op, Y, A)
#' #The parameters can now be estimated by minimizing mlik with optim

```

```

# Choose some reasonable starting values depending on the size of the domain
theta0 <- log(c( 1 / sqrt(var(c(Y))),sqrt(8), 0.9, 0.01))
# run estimation and display the results
theta <- optim(theta0, mlik)
print(data.frame(
  tau = c(tau[1], exp(theta$par[1])), kappa = c(kappa[1], exp(theta$par[2])),
  nu = c(nu, exp(theta$par[3])), sigma.e = c(sigma.e, exp(theta$par[4])),
  row.names = c("Truth", "Estimates")
))

```

cross_validation *Perform cross-validation on a list of fitted models.*

Description

Obtain several scores for a list of fitted models according to a folding scheme.

Usage

```

cross_validation(
  models,
  model_names = NULL,
  scores = c("mse", "crps", "scrps", "dss"),
  cv_type = c("k-fold", "loo", "lpo"),
  k = 5,
  percentage = 20,
  number_folds = 10,
  n_samples = 1000,
  return_scores_folds = FALSE,
  orientation_results = c("negative", "positive"),
  include_best = TRUE,
  train_test_indexes = NULL,
  return_train_test = FALSE,
  parallelize_RP = FALSE,
  n_cores_RP = parallel::detectCores() - 1,
  true_CV = FALSE,
  save_settings = FALSE,
  print = TRUE,
  fit_verbose = FALSE
)

```

Arguments

models	A fitted model obtained from calling the <code>bru()</code> function or a list of models fitted with the <code>bru()</code> function.
--------	---

<code>model_names</code>	A vector containing the names of the models to appear in the returned <code>data.frame</code> . If <code>NULL</code> , the names will be of the form <code>Model 1</code> , <code>Model 2</code> , and so on. By default, it will try to obtain the name from the models list.
<code>scores</code>	A vector containing the scores to be computed. The options are <code>"mse"</code> , <code>"crps"</code> , <code>"scrps"</code> and <code>"dss"</code> . By default, all scores are computed.
<code>cv_type</code>	The type of the folding to be carried out. The options are <code>k-fold</code> for <code>k-fold cross-validation</code> , in which case the parameter <code>k</code> should be provided, <code>loo</code> , for <code>leave-one-out</code> and <code>lpo</code> for <code>leave-percentage-out</code> , in this case, the parameter <code>percentage</code> should be given, and also the <code>number_folds</code> with the number of folds to be done. The default is <code>k-fold</code> .
<code>k</code>	The number of folds to be used in <code>k-fold cross-validation</code> . Will only be used if <code>cv_type</code> is <code>k-fold</code> .
<code>percentage</code>	The percentage (from 1 to 99) of the data to be used to train the model. Will only be used if <code>cv_type</code> is <code>lpo</code> .
<code>number_folds</code>	Number of folds to be done if <code>cv_type</code> is <code>lpo</code> .
<code>n_samples</code>	Number of samples to compute the posterior statistics to be used to compute the scores.
<code>return_scores_folds</code>	If <code>TRUE</code> , the scores for each fold will also be returned.
<code>orientation_results</code>	character vector. The options are <code>"negative"</code> and <code>"positive"</code> . If <code>"negative"</code> , the smaller the scores the better. If <code>"positive"</code> , the larger the scores the better.
<code>include_best</code>	Should a row indicating which model was the best for each score be included?
<code>train_test_indexes</code>	A list containing two entries <code>train</code> , which is a list whose elements are vectors of indexes of the training data, and <code>test</code> , which is a list whose elements are vectors of indexes of the test data. Typically this will be returned list obtained by setting the argument <code>return_train_test</code> to <code>TRUE</code> .
<code>return_train_test</code>	Logical. Should the training and test indexes be returned? If <code>'TRUE'</code> the train and test indexes will the <code>'train_test'</code> element of the returned list.
<code>parallelize_RP</code>	Logical. Should the computation of CRPS and SCRPS be parallelized?
<code>n_cores_RP</code>	Number of cores to be used if <code>parallelize_rp</code> is <code>TRUE</code> .
<code>true_CV</code>	Should a <code>TRUE</code> cross-validation be performed? If <code>TRUE</code> the models will be fitted on the training dataset. If <code>FALSE</code> , the parameters will be kept fixed at the ones obtained in the result object.
<code>save_settings</code>	Logical. If <code>TRUE</code> , the settings used in the cross-validation will also be returned.
<code>print</code>	Should partial results be printed throughout the computation?
<code>fit_verbose</code>	Should INLA's run during cross-validation be verbose?

Value

A `data.frame` with the fitted models and the corresponding scores.

folded.matern.covariance.1d

The 1d folded Matern covariance function

Description

`folded.matern.covariance.1d` evaluates the 1d folded Matern covariance function over an interval $[0, L]$.

Usage

```
folded.matern.covariance.1d(
  h,
  m,
  kappa,
  nu,
  sigma,
  L = 1,
  N = 10,
  boundary = c("neumann", "dirichlet", "periodic")
)
```

Arguments

<code>h, m</code>	Vectors of arguments of the covariance function.
<code>kappa</code>	Range parameter.
<code>nu</code>	Shape parameter.
<code>sigma</code>	Standard deviation.
<code>L</code>	The upper bound of the interval $[0, L]$. By default, <code>L=1</code> .
<code>N</code>	The truncation parameter.
<code>boundary</code>	The boundary condition. The possible conditions are "neumann" (default), "dirichlet" or "periodic".

Details

`folded.matern.covariance.1d` evaluates the 1d folded Matern covariance function over an interval $[0, L]$ under different boundary conditions. For periodic boundary conditions

$$C_{\mathcal{P}}(h, m) = \sum_{k=-\infty}^{\infty} (C(h - m + 2kL),$$

for Neumann boundary conditions

$$C_{\mathcal{N}}(h, m) = \sum_{k=-\infty}^{\infty} (C(h - m + 2kL) + C(h + m + 2kL)),$$

and for Dirichlet boundary conditions:

$$C_{\mathcal{D}}(h, m) = \sum_{k=-\infty}^{\infty} (C(h - m + 2kL) - C(h + m + 2kL)),$$

where $C(\cdot)$ is the Matern covariance function:

$$C(h) = \frac{\sigma^2}{2^{\nu-1}\Gamma(\nu)} (\kappa h)^{\nu} K_{\nu}(\kappa h).$$

We consider the truncation:

$$C_{\mathcal{P},N}(h, m) = \sum_{k=-N}^N C(h - m + 2kL), C_{\mathcal{N},N}(h, m) = \sum_{k=-\infty}^{\infty} (C(h - m + 2kL) + C(h + m + 2kL)),$$

and

$$C_{\mathcal{D},N}(h, m) = \sum_{k=-N}^N (C(h - m + 2kL) - C(h + m + 2kL)).$$

Value

A matrix with the corresponding covariance values.

Examples

```
x <- seq(from = 0, to = 1, length.out = 101)
plot(x, folded.matern.covariance.1d(rep(0.5, length(x)), x,
kappa = 10, nu = 1 / 5, sigma = 1),
type = "l", ylab = "C(h)", xlab = "h")
)
```

folded.matern.covariance.2d

The 2d folded Matern covariance function

Description

`folded.matern.covariance.2d` evaluates the 2d folded Matern covariance function over an interval $[0, L] \times [0, L]$.

Usage

```
folded.matern.covariance.2d(
  h,
  m,
  kappa,
  nu,
  sigma,
  L = 1,
  N = 10,
  boundary = c("neumann", "dirichlet", "periodic", "R2")
)
```

Arguments

<code>h, m</code>	Vectors with two coordinates.
<code>kappa</code>	Range parameter.
<code>nu</code>	Shape parameter.
<code>sigma</code>	Standard deviation.
<code>L</code>	The upper bound of the square $[0, L] \times [0, L]$. By default, <code>L=1</code> .
<code>N</code>	The truncation parameter.
<code>boundary</code>	The boundary condition. The possible conditions are "neumann" (default), "dirichlet", "periodic" or "R2".

Details

`folded.matern.covariance.2d` evaluates the 1d folded Matern covariance function over an interval $[0, L] \times [0, L]$ under different boundary conditions. For periodic boundary conditions

$$C_{\mathcal{P}}((h_1, h_2), (m_1, m_2)) = \sum_{k_2=-\infty}^{\infty} \sum_{k_1=-\infty}^{\infty} (C(\|(h_1 - m_1 + 2k_1L, h_2 - m_2 + 2k_2L)\|)),$$

for Neumann boundary conditions

$$C_{\mathcal{N}}((h_1, h_2), (m_1, m_2)) = \sum_{k_2=-\infty}^{\infty} \sum_{k_1=-\infty}^{\infty} (C(\|(h_1 - m_1 + 2k_1L, h_2 - m_2 + 2k_2L)\|) + C(\|(h_1 - m_1 + 2k_1L, h_2 + m_2 + 2k_2L)\|))$$

and for Dirichlet boundary conditions:

$$C_{\mathcal{D}}((h_1, h_2), (m_1, m_2)) = \sum_{k_2=-\infty}^{\infty} \sum_{k_1=-\infty}^{\infty} (C(\|(h_1 - m_1 + 2k_1L, h_2 - m_2 + 2k_2L)\|) - C(\|(h_1 - m_1 + 2k_1L, h_2 + m_2 + 2k_2L)\|))$$

where $C(\cdot)$ is the Matern covariance function:

$$C(h) = \frac{\sigma^2}{2^{\nu-1}\Gamma(\nu)} (\kappa h)^{\nu} K_{\nu}(\kappa h).$$

We consider the truncation for k_1, k_2 from $-N$ to N .

Value

The corresponding covariance.

Examples

```
h <- c(0.5, 0.5)
m <- c(0.5, 0.5)
folded.matern.covariance.2d(h, m, kappa = 10, nu = 1 / 5, sigma = 1)
```

`fractional.operators` *Rational approximations of fractional operators*

Description

`fractional.operators` is used for computing an approximation, which can be used for inference and simulation, of the fractional SPDE

$$L^\beta(\tau u(s)) = W.$$

Here L is a differential operator, $\beta > 0$ is the fractional power, τ is a positive scalar or vector that scales the variance of the solution u , and W is white noise.

Usage

```
fractional.operators(L, beta, C, scale.factor, m = 1, tau = 1)
```

Arguments

<code>L</code>	A finite element discretization of the operator L .
<code>beta</code>	The positive fractional power.
<code>C</code>	The mass matrix of the finite element discretization.
<code>scale.factor</code>	A constant c is a lower bound for the the smallest eigenvalue of the non-discretized operator L .
<code>m</code>	The order of the rational approximation, which needs to be a positive integer. The default value is 1. Higer values gives a more accurate approximation, which are more computationally expensive to use for inference. Currently, the largest value of <code>m</code> that is implemented is 4.
<code>tau</code>	The constant or vector that scales the variance of the solution. The default value is 1.

Details

The approximation is based on a rational approximation of the fractional operator, resulting in an approximate model on the form

$$P_l u(s) = P_r W,$$

where $P_j = p_j(L)$ are non-fractional operators defined in terms of polynomials p_j for $j = l, r$. The order of p_r is given by m and the order of p_l is $m + m_\beta$ where m_β is the integer part of β if $\beta > 1$ and $m_\beta = 1$ otherwise.

The discrete approximation can be written as $u = P_r x$ where $x \sim N(0, Q^{-1})$ and $Q = P_l^T C^{-1} P_l$. Note that the matrices P_r and Q may be ill-conditioned for $m > 1$. In this case, the methods in [operator.operations\(\)](#) should be used for operations involving the matrices, since these methods are more numerically stable.

Value

`fractional.operators` returns an object of class "rSPDEobj". This object contains the following quantities:

P1	The operator P_l .
Pr	The operator P_r .
C	The mass lumped mass matrix.
Ci	The inverse of C.
m	The order of the rational approximation.
beta	The fractional power.
type	String indicating the type of approximation.
Q	The matrix <code>t(P1) %*% solve(C, P1)</code> .
type	String indicating the type of approximation.
P1.factors	List with elements that can be used to assemble P_l .
Pr.factors	List with elements that can be used to assemble P_r .

See Also

[matern.operators\(\)](#), [spde.matern.operators\(\)](#), [matern.operators\(\)](#)

Examples

```
# Compute rational approximation of a Gaussian process with a
# Matern covariance function on R
kappa <- 10
sigma <- 1
nu <- 0.8

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation of covariance function at 0.5
```

```

tau <- sqrt(gamma(nu) / (sigma^2 * kappa^(2 * nu) *
(4 * pi)^(1 / 2) * gamma(nu + 1 / 2)))
op <- fractional.operators(
  L = fem$G + kappa^2 * fem$C, beta = (nu + 1 / 2) / 2,
  C = fem$C, scale.factor = kappa^2, tau = tau
)
v <- t(rSPDE.A1d(x, 0.5))
c.approx <- Sigma.mult(op, v)

# plot the result and compare with the true Matern covariance
plot(x, matern.covariance(abs(x - 0.5), kappa, nu, sigma),
  type = "l", ylab = "C(h)",
  xlab = "h", main = "Matern covariance and rational approximations"
)
lines(x, c.approx, col = 2)

```

get.initial.values.rSPDE

Initial values for log-likelihood optimization in rSPDE models with a latent stationary Gaussian Matern model

Description

Auxiliar function to obtain domain-based initial values for log-likelihood optimization in rSPDE models with a latent stationary Gaussian Matern model

Usage

```

get.initial.values.rSPDE(
  mesh = NULL,
  mesh.range = NULL,
  graph.obj = NULL,
  n.spde = 1,
  dim = NULL,
  B.tau = NULL,
  B.kappa = NULL,
  B.sigma = NULL,
  B.range = NULL,
  nu = NULL,
  parameterization = c("matern", "spde"),
  include.nu = TRUE,
  log.scale = TRUE,
  nu.upper.bound = NULL
)

```

Arguments

<code>mesh</code>	An in INLA mesh
<code>mesh.range</code>	The range of the mesh.
<code>graph.obj</code>	A <code>metric_graph</code> object. To be used in case both <code>mesh</code> and <code>mesh.range</code> are <code>NULL</code> .
<code>n.spde</code>	The number of basis functions in the mesh model.
<code>dim</code>	The dimension of the domain.
<code>B.tau</code>	Matrix with specification of log-linear model for τ . Will be used if <code>parameterization</code> = 'spde'.
<code>B.kappa</code>	Matrix with specification of log-linear model for κ . Will be used if <code>parameterization</code> = 'spde'.
<code>B.sigma</code>	Matrix with specification of log-linear model for σ . Will be used if <code>parameterization</code> = 'matern'.
<code>B.range</code>	Matrix with specification of log-linear model for ρ , which is a range-like parameter (it is exactly the range parameter in the stationary case). Will be used if <code>parameterization</code> = 'matern'.
<code>nu</code>	The smoothness parameter.
<code>parameterization</code>	Which parameterization to use? <code>matern</code> uses <code>range</code> , <code>std. deviation</code> and <code>nu</code> (smoothness). <code>spde</code> uses <code>kappa</code> , <code>tau</code> and <code>nu</code> (smoothness). The default is <code>matern</code> .
<code>include.nu</code>	Should we also provide an initial guess for <code>nu</code> ?
<code>log.scale</code>	Should the results be provided in log scale?
<code>nu.upper bound</code>	Should an upper bound for <code>nu</code> be considered?

Value

A vector of the form (`theta_1,theta_2,theta_3`) or where `theta_1` is the initial guess for `tau`, `theta_2` is the initial guess for `kappa` and `theta_3` is the initial guess for `nu`.

`gg_df`

Data frame for result objects from R-INLA fitted models to be used in ggplot2

Description

Data frame for result objects from R-INLA fitted models to be used in `ggplot2`

Usage

```
gg_df(result, ...)
```

Arguments

- `result` a result object for which the data frame is desired
`...` further arguments passed to or from other methods.

Value

A data frame containing the posterior densities.

`gg_df.rspde_result` *Data frame for rspde_result objects to be used in ggplot2*

Description

Returns a ggplot-friendly data-frame with the marginal posterior densities.

Usage

```
## S3 method for class 'rspde_result'
gg_df(
  result,
  parameter = result$params,
  transform = TRUE,
  restrict_x_axis = NULL,
  restrict_quantiles = NULL,
  ...
)
```

Arguments

- `result` An `rspde_result` object.
`parameter` Vector. Which parameters to get the posterior density in the data.frame? The options are std.dev, range, tau, kappa and nu.
`transform` Should the posterior density be given in the original scale?
`restrict_x_axis` Variables to restrict the range of x axis based on quantiles.
`restrict_quantiles` Named list of quantiles to restrict x axis. It should contain the name of the parameter along with a vector with two elements specifying the lower and upper quantiles. The names should be match the ones in `result$params`. For example, if we want to restrict nu to the 0.05 and 0.95 quantiles we do `restrict_quantiles = c(0.05, 0.95)`.
`...` currently not used.

Value

A data frame containing the posterior densities.

glance.rspde_lme *Glance at an rspde_lme object*

Description

Glance accepts a `rspde_lme` object and returns a `tidy::tibble()` with exactly one row of model summaries. The summaries are the square root of the estimated variance of the measurement error, residual degrees of freedom, AIC, BIC, log-likelihood, the type of latent model used in the fit and the total number of observations.

Usage

```
## S3 method for class 'rspde_lme'  
glance(x, ...)
```

Arguments

`x` An `rspde_lme` object.
`...` Currently not used.

Value

A `tidy::tibble()` with exactly one row and columns:

- `nobs` Number of observations used.
- `sigma` the square root of the estimated residual variance
- `logLik` The log-likelihood of the model.
- `AIC` Akaike's Information Criterion for the model.
- `BIC` Bayesian Information Criterion for the model.
- `deviance` Deviance of the model.
- `df.residual` Residual degrees of freedom.
- `model.type` Type of latent model fitted.

See Also

[augment.rspde_lme](#)

`graph_data_rspde`*Data extraction from metric graphs for 'rSPDE' models*

Description

Extracts data from metric graphs to be used by 'INLA' and 'inlabru'.

Usage

```
graph_data_rspde(
  graph_rspde,
  name = "field",
  repl = NULL,
  group = NULL,
  group_col = NULL,
  only_pred = FALSE,
  loc = NULL,
  loc_name = NULL,
  tibble = FALSE,
  drop_na = FALSE,
  drop_all_na = TRUE
)
```

Arguments

<code>graph_rspde</code>	An <code>inla_metric_graph_spde</code> object built with the <code>rspde.metric_graph()</code> function.
<code>name</code>	A character string with the base name of the effect.
<code>repl</code>	Which replicates? If there is no replicates, one can set <code>repl</code> to <code>NULL</code> . If one wants all replicates, then one sets to <code>repl</code> to <code>.all</code> .
<code>group</code>	Which groups? If there is no groups, one can set <code>group</code> to <code>NULL</code> . If one wants all groups, then one sets to <code>group</code> to <code>.all</code> .
<code>group_col</code>	Which "column" of the data contains the group variable?
<code>only_pred</code>	Should only return the <code>data.frame</code> to the prediction data?
<code>loc</code>	Locations. If not given, they will be chosen as the available locations on the metric graph internal dataset.
<code>loc_name</code>	Character with the name of the location variable to be used in 'inlabru' prediction.
<code>tibble</code>	Should the data be returned as a <code>tidyrr::tibble</code> ?
<code>drop_na</code>	Should the rows with at least one NA for one of the columns be removed? DEFAULT is <code>FALSE</code> . This option is turned to <code>FALSE</code> if <code>only_pred</code> is <code>TRUE</code> .
<code>drop_all_na</code>	Should the rows with all variables being NA be removed? DEFAULT is <code>TRUE</code> . This option is turned to <code>FALSE</code> if <code>only_pred</code> is <code>TRUE</code> .

Value

An 'INLA' and 'inlabru' friendly list with the data.

intrinsic.matern.operators

Covariance-based approximations of intrinsic fields

Description

`intrinsic.matern.operators` is used for computing a covariance-based rational SPDE approximation of intrinsic fields on R^d defined through the SPDE

$$(-\Delta)^{\beta/2}(\kappa^2 - \Delta)^{\alpha/2}(\tau u) = \mathcal{W}$$

Usage

```
intrinsic.matern.operators(
  kappa,
  tau,
  alpha,
  beta = 1,
  G = NULL,
  C = NULL,
  d = NULL,
  mesh = NULL,
  graph = NULL,
  loc_mesh = NULL,
  m_alpha = 2,
  m_beta = 2,
  compute_higher_order = FALSE,
  return_block_list = FALSE,
  type_rational_approximation = c("chebfun", "brasil", "chebfunLB"),
  fem_mesh_matrices = NULL,
  scaling = NULL
)
```

Arguments

<code>kappa</code>	range parameter
<code>tau</code>	precision parameter
<code>alpha</code>	Smoothness parameter
<code>beta</code>	Smoothness parameter
<code>G</code>	The stiffness matrix of a finite element discretization of the domain of interest.
<code>C</code>	The mass matrix of a finite element discretization of the domain of interest.
<code>d</code>	The dimension of the domain.

<code>mesh</code>	An inla mesh.
<code>graph</code>	An optional <code>metric_graph</code> object. Replaces <code>d</code> , <code>C</code> and <code>G</code> .
<code>loc_mesh</code>	locations for the mesh for <code>d=1</code> .
<code>m_alpha</code>	The order of the rational approximation for the Matérn part, which needs to be a positive integer. The default value is 2.
<code>m_beta</code>	The order of the rational approximation for the intrinsic part, which needs to be a positive integer. The default value is 2.
<code>compute_higher_order</code>	Logical. Should the higher order finite element matrices be computed?
<code>return_block_list</code>	Logical. For <code>type = "covariance"</code> , should the block parts of the precision matrix be returned separately as a list?
<code>type_rational_approximation</code>	Which type of rational approximation should be used? The current types are " <code>chebfun</code> ", " <code>brasil</code> " or " <code>chebfunLB</code> ".
<code>fem_mesh_matrices</code>	A list containing FEM-related matrices. The list should contain elements <code>c0</code> , <code>g1</code> , <code>g2</code> , <code>g3</code> , etc.
<code>scaling</code>	second lowest eigenvalue of <code>g1</code>

Details

The covariance operator

$$\tau^{-2}(-\Delta)^\beta(\kappa^2 - \Delta)^\alpha$$

is approximated based on rational approximations of the two fractional components. The Laplacians are equipped with homogeneous Neumann boundary conditions and a zero-mean constraint is additionally imposed to obtain a non-intrinsic model.

Value

`intrinsic.matern.operators` returns an object of class "`intrinsicCBrSPDEobj`". This object is a list containing the following quantities:

<code>C</code>	The mass lumped mass matrix.
<code>Ci</code>	The inverse of <code>C</code> .
<code>GCi</code>	The stiffness matrix <code>G</code> times <code>Ci</code>
<code>Gk</code>	The stiffness matrix <code>G</code> along with the higher-order FEM-related matrices <code>G2</code> , <code>G3</code> , etc.
<code>fem_mesh_matrices</code>	A list containing the mass lumped mass matrix, the stiffness matrix and the higher-order FEM-related matrices.
<code>m_alpha</code>	The order of the rational approximation for the Matérn part.
<code>m_beta</code>	The order of the rational approximation for the intrinsic part.
<code>alpha</code>	The fractional power of the Matérn part of the operator.

beta	The fractional power of the intrinsic part of the operator.
type	String indicating the type of approximation.
d	The dimension of the domain.
A	Matrix that sums the components in the approximation to the mesh nodes.
kappa	Range parameter of the covariance function
tau	Scale parameter of the covariance function.
type	String indicating the type of approximation.

Examples

```
if (requireNamespace("RSpectra", quietly = TRUE)){
  x <- seq(from = 0, to = 10, length.out = 201)
  beta <- 1
  alpha <- 1
  kappa <- 1
  op <- intrinsic.matern.operators(kappa = kappa, tau = 1, alpha = alpha,
                                    beta = beta, loc_mesh = x, d=1)
  # Compute and plot the variogram of the model
  Sigma <- op$A %*% solve(op$Q,t(op$A))
  One <- rep(1, times = ncol(Sigma))
  D <- diag(Sigma)
  Gamma <- 0.5*(One %*% t(D) + D %*% t(One) - 2 * Sigma)
  k <- 100
  plot(x, Gamma[k, ], type = "l")
  lines(x,
        variogram.intrinsic.spde(x[k], x, kappa, alpha, beta, L = 10, d = 1),
        col=2, lty = 2)
}
```

matern.covariance *The Matern covariance function*

Description

`matern.covariance` evaluates the Matern covariance function

$$C(h) = \frac{\sigma^2}{2^{\nu-1}\Gamma(\nu)}(\kappa h)^\nu K_\nu(\kappa h).$$

Usage

```
matern.covariance(h, kappa, nu, sigma)
```

Arguments

h	Distances to evaluate the covariance function at.
kappa	Range parameter.
nu	Shape parameter.
sigma	Standard deviation.

Value

A vector with the values $C(h)$.

Examples

```
x <- seq(from = 0, to = 1, length.out = 101)
plot(x, matern.covariance(abs(x - 0.5), kappa = 10, nu = 1 / 5, sigma = 1),
      type = "l", ylab = "C(h)", xlab = "h"
)
```

Description

`matern.operators` is used for computing a rational SPDE approximation of a stationary Gaussian random fields on R^d with a Matern covariance function

$$C(h) = \frac{\sigma^2}{2^{\nu-1}\Gamma(\nu)}(\kappa h)^\nu K_\nu(\kappa h)$$

Usage

```
matern.operators(
  kappa = NULL,
  tau = NULL,
  alpha = NULL,
  sigma = NULL,
  range = NULL,
  nu = NULL,
  G = NULL,
  C = NULL,
  d = NULL,
  mesh = NULL,
  graph = NULL,
  range_mesh = NULL,
  loc_mesh = NULL,
  m = 1,
  type = c("covariance", "operator"),
  parameterization = c("spde", "matern"),
  compute_higher_order = FALSE,
  return_block_list = FALSE,
  type_rational_approximation = c("chebfun", "brasil", "chebfunLB"),
  fem_mesh_matrices = NULL,
  compute_logdet = FALSE
)
```

Arguments

kappa	Parameter kappa of the SPDE representation. If NULL, the range parameter will be used. If the range is also NULL, a starting value based on the mesh will be supplied.
tau	Parameter tau of the SPDE representation. If both sigma and tau are NULL, a starting value based on the mesh will be supplied.
alpha	Parameter alpha of the SPDE representation. If alpha is NULL, a starting value will be supplied.
sigma	Standard deviation of the covariance function. Used if parameterization is <code>matern</code> . If NULL, tau will be used. If tau is also NULL, a starting value based on the mesh will be supplied.
range	Range parameter of the covariance function. Used if parameterization is <code>matern</code> . If range is NULL, a starting value based on the mesh will be supplied.
nu	Shape parameter of the covariance function. Used if parameterization is <code>matern</code> . If NULL, a starting value will be supplied.
G	The stiffness matrix of a finite element discretization of the domain of interest. Does not need to be given if either mesh or graph is supplied.
C	The mass matrix of a finite element discretization of the domain of interest. Does not need to be given if either mesh or graph is supplied.
d	The dimension of the domain. Does not need to be given if either mesh or graph is provided.
mesh	An optional fmesher mesh. Replaces d, C and G.
graph	An optional <code>metric_graph</code> object. Replaces d, C and G.
range_mesh	The range of the mesh. Will be used to provide starting values for the parameters. Will be used if mesh and graph are NULL, and if one of the parameters (kappa or tau for spde parameterization, or sigma or range for matern parameterization) are not provided.
loc_mesh	The mesh locations used to construct the matrices C and G. This option should be provided if one wants to use the <code>rspde_lme()</code> function and will not provide neither graph nor mesh. Only works for 1d data. Does not work for metric graphs. For metric graphs you should supply the graph using the graph argument.
m	The order of the rational approximation, which needs to be a positive integer. The default value is 1.
type	The type of the rational approximation. The options are "covariance" and "operator". The default is "covariance".
parameterization	Which parameterization to use? <code>matern</code> uses range, std. deviation and nu (smoothness). <code>spde</code> uses kappa, tau and alpha. The default is <code>spde</code> .
compute_higher_order	Logical. Should the higher order finite element matrices be computed?
return_block_list	Logical. For <code>type = "covariance"</code> , should the block parts of the precision matrix be returned separately as a list?

```

type_rational_approximation
    Which type of rational approximation should be used? The current types are
    "chebfun", "brasil" or "chebfunLB".
fem_mesh_matrices
    A list containing FEM-related matrices. The list should contain elements c0, g1,
    g2, g3, etc.
compute_logdet Should log determinants be computed while building the model? (For covariance-
    based models)

```

Details

If type is "covariance", we use the covariance-based rational approximation of the fractional operator. In the SPDE approach, we model u as the solution of the following SPDE:

$$L^{\alpha/2}(\tau u) = \mathcal{W},$$

where $L = -\Delta + \kappa^2 I$ and \mathcal{W} is the standard Gaussian white noise. The covariance operator of u is given by $L^{-\alpha}$. Now, let L_h be a finite-element approximation of L . We can use a rational approximation of order m on $L_h^{-\alpha}$ to obtain the following approximation:

$$L_{h,m}^{-\alpha} = L_h^{-m_\alpha} p(L_h^{-1}) q(L_h^{-1})^{-1},$$

where $m_\alpha = \lfloor \alpha \rfloor$, p and q are polynomials arising from such rational approximation. From this approximation we construct an approximate precision matrix for u .

If type is "operator", the approximation is based on a rational approximation of the fractional operator $(\kappa^2 - \Delta)^\beta$, where $\beta = (\nu + d/2)/2$. This results in an approximate model of the form

$$P_l u(s) = P_r W,$$

where $P_j = p_j(L)$ are non-fractional operators defined in terms of polynomials p_j for $j = l, r$. The order of p_r is given by m and the order of p_l is $m + m_\beta$ where m_β is the integer part of β if $\beta > 1$ and $m_\beta = 1$ otherwise.

The discrete approximation can be written as $u = P_r x$ where $x \sim N(0, Q^{-1})$ and $Q = P_l^T C^{-1} P_l$. Note that the matrices P_r and Q may be ill-conditioned for $m > 1$. In this case, the methods in [operator.operations\(\)](#) should be used for operations involving the matrices, since these methods are more numerically stable.

Value

If type is "covariance", then `matern.operators` returns an object of class "CBrSPDEobj". This object is a list containing the following quantities:

<code>C</code>	The mass lumped mass matrix.
<code>Ci</code>	The inverse of <code>C</code> .
<code>Gci</code>	The stiffness matrix <code>G</code> times <code>Ci</code>
<code>Gk</code>	The stiffness matrix <code>G</code> along with the higher-order FEM-related matrices <code>G2</code> , <code>G3</code> , etc.
<code>fem_mesh_matrices</code>	A list containing the mass lumped mass matrix, the stiffness matrix and the higher-order FEM-related matrices.

m	The order of the rational approximation.
alpha	The fractional power of the precision operator.
type	String indicating the type of approximation.
d	The dimension of the domain.
nu	Shape parameter of the covariance function.
kappa	Range parameter of the covariance function
tau	Scale parameter of the covariance function.
sigma	Standard deviation of the covariance function.
type	String indicating the type of approximation.

If type is "operator", then `matern.operators` returns an object of class "rSPDEobj". This object contains the quantities listed in the output of [fractional.operators\(\)](#), the G matrix, the dimension of the domain, as well as the parameters of the covariance function.

See Also

[fractional.operators\(\)](#), [spde.matern.operators\(\)](#), [matern.operators\(\)](#)

Examples

```
# Compute the covariance-based rational approximation of a
# Gaussian process with a Matern covariance function on R
kappa <- 10
sigma <- 1
nu <- 0.8
range <- sqrt(8*nu)/kappa

# create mass and stiffness matrices for a FEM discretization
nobs <- 101
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation of covariance function at 0.5
op_cov <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2,
  parameterization = "matern"
)

v <- t(rSPDE.A1d(x, 0.5))
# Compute the precision matrix
Q <- op_cov$Q
# A matrix here is the identity matrix
A <- Diagonal(nobs)
# We need to concatenate 3 A's since we are doing a covariance-based rational
# approximation of order 2
Abar <- cbind(A, A, A)
w <- rbind(v, v, v)
# The approximate covariance function:
```

```

c_cov.approx <- (Abar) %*% solve(Q, w)
c.true <- folded.matern.covariance.1d(rep(0.5, length(x)),
  abs(x), kappa, nu, sigma)

# plot the result and compare with the true Matern covariance
plot(x, c.true,
  type = "l", ylab = "C(h)",
  xlab = "h", main = "Matern covariance and rational approximations"
)
lines(x, c_cov.approx, col = 2)

# Compute the operator-based rational approximation of a Gaussian
# process with a Matern covariance function on R
kappa <- 10
sigma <- 1
nu <- 0.8
range <- sqrt(8*nu)/kappa

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation of covariance function at 0.5
op <- matern.operators(
  range = range, sigma = sigma, nu = nu,
  loc_mesh = x, d = 1,
  type = "operator",
  parameterization = "matern"
)

v <- t(rSPDE.A1d(x, 0.5))
c.approx <- Sigma.mult(op, v)
c.true <- folded.matern.covariance.1d(rep(0.5, length(x)),
  abs(x), kappa, nu, sigma)

# plot the result and compare with the true Matern covariance
plot(x, c.true,
  type = "l", ylab = "C(h)",
  xlab = "h", main = "Matern covariance and rational approximation"
)
lines(x, c.approx, col = 2)

```

Description

Functions for multiplying and solving with the P_r and P_l operators as well as the latent precision matrix $Q = P_l C^{-1} P_l$ and covariance matrix $\Sigma = P_r Q^{-1} P_r^T$. These operations are done without first assembling P_r , P_l in order to avoid numerical problems caused by ill-conditioned matrices.

Usage

```

Pr.mult(obj, v, transpose = FALSE)

Pr.solve(obj, v, transpose = FALSE)

Pl.mult(obj, v, transpose = FALSE)

Pl.solve(obj, v, transpose = FALSE)

Q.mult(obj, v)

Q.solve(obj, v)

Qsqrt.mult(obj, v, transpose = FALSE)

Qsqrt.solve(obj, v, transpose = FALSE)

Sigma.mult(obj, v)

Sigma.solve(obj, v)

```

Arguments

obj	rSPDE object
v	vector to apply the operation to
transpose	set to TRUE if the operation should be performed with the transposed object

Details

`Pl.mult`, `Pr.mult`, and `Q.mult` multiplies the vector with the respective object. Changing `mult` to `solve` in the function names multiplies the vector with the inverse of the object. `Qsqrt.mult` and `Qsqrt.solve` performs the operations with the square-root type object $Q_r = C^{-1/2}P_l$ defined so that $Q = Q_r^T Q_r$.

Value

A vector with the values of the operation

precision

Get the precision matrix of CBrSPDEobj objects

Description

Function to get the precision matrix of a CBrSPDEobj object

Usage

```
precision(object, ...)

## S3 method for class 'CBrSPDEobj'
precision(
  object,
  user_nu = NULL,
  user_kappa = NULL,
  user_sigma = NULL,
  user_range = NULL,
  user_tau = NULL,
  user_m = NULL,
  ...
)
```

Arguments

object	The covariance-based rational SPDE approximation, computed using matern.operators()
...	Currently not used.
user_nu	If non-null, update the shape parameter of the covariance function.
user_kappa	If non-null, update the range parameter of the covariance function.
user_sigma	If non-null, update the standard deviation of the covariance function.
user_range	If non-null, update the range parameter of the covariance function.
user_tau	If non-null, update the parameter tau.
user_m	If non-null, update the order of the rational approximation, which needs to be a positive integer.

Value

The precision matrix.

See Also

[simulate.CBrSPDEobj\(\)](#), [matern.operators\(\)](#)

Examples

```
# Compute the covariance-based rational approximation of a
# Gaussian process with a Matern covariance function on R
kappa <- 10
sigma <- 1
nu <- 0.8
range <- 0.2

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)
```

```

# compute rational approximation of covariance function at 0.5
tau <- sqrt(gamma(nu) / (sigma^2 * kappa^(2 * nu) *
(4 * pi)^(1 / 2) * gamma(nu + 1 / 2)))
op_cov <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2,
  parameterization = "matern"
)

# Get the precision matrix:
prec_matrix <- precision(op_cov)

```

precision.inla_rspde *Get the precision matrix of inla_rspde objects*

Description

Function to get the precision matrix of an `inla_rspde` object created with the `rspde.matern()` function.

Usage

```
## S3 method for class 'inla_rspde'
precision(object, theta = NULL, ...)
```

Arguments

<code>object</code>	The <code>inla_rspde</code> object obtained with the <code>rspde.matern()</code> function.
<code>theta</code>	If null, the starting values for theta will be used. Otherwise, it must be supplied as a vector. For stationary models, we have <code>theta = c(log(tau), log(kappa), nu)</code> . For nonstationary models, we have <code>theta = c(theta_1, theta_2, ..., theta_n, nu)</code> .
<code>...</code>	Currently not used.

Value

The precision matrix.

See Also

[precision.CBrSPDEobj\(\)](#), [matern.operators\(\)](#)

<i>predict.CBrSPDEobj</i>	<i>Prediction of a fractional SPDE using the covariance-based rational SPDE approximation</i>
---------------------------	---

Description

The function is used for computing kriging predictions based on data $Y_i = u(s_i) + \epsilon_i$, where ϵ is mean-zero Gaussian measurement noise and $u(s)$ is defined by a fractional SPDE $(\kappa^2 I - \Delta)^{\alpha/2}(\tau u(s)) = W$, where W is Gaussian white noise and $\alpha = \nu + d/2$, where d is the dimension of the domain.

Usage

```
## S3 method for class 'CBrSPDEobj'
predict(
  object,
  A,
  Aprd,
  Y,
  sigma.e,
  mu = 0,
  compute.variances = FALSE,
  posterior_samples = FALSE,
  n_samples = 100,
  only_latent = FALSE,
  ...
)
```

Arguments

object	The covariance-based rational SPDE approximation, computed using matern.operators()
A	A matrix linking the measurement locations to the basis of the FEM approximation of the latent model.
Aprd	A matrix linking the prediction locations to the basis of the FEM approximation of the latent model.
Y	A vector with the observed data, can also be a matrix where the columns are observations of independent replicates of u .
sigma.e	The standard deviation of the Gaussian measurement noise. Put to zero if the model does not have measurement noise.
mu	Expectation vector of the latent field (default = 0).
compute.variances	Set to also TRUE to compute the kriging variances.
posterior_samples	If TRUE, posterior samples will be returned.
n_samples	Number of samples to be returned. Will only be used if sampling is TRUE.

only_latent Should the posterior samples be only given to the latent model?
 ... further arguments passed to or from other methods.

Value

A list with elements

mean The kriging predictor (the posterior mean of $u|Y$).
 variance The posterior variances (if computed).

Examples

```

set.seed(123)
# Sample a Gaussian Matern process on R using a rational approximation
kappa <- 10
sigma <- 1
nu <- 0.8
sigma.e <- 0.3
range <- 0.2

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

tau <- sqrt(gamma(nu) / (sigma^2 * kappa^(2 * nu) *
  (4 * pi)^(1 / 2) * gamma(nu + 1 / 2)))

# Compute the covariance-based rational approximation
op_cov <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2,
  parameterization = "matern"
)

# Sample the model
u <- simulate(op_cov)

# Create some data
obs.loc <- runif(n = 10, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
Y <- as.vector(A %*% u + sigma.e * rnorm(10))

# compute kriging predictions at the FEM grid
A.krig <- rSPDE.A1d(x, x)
u.krig <- predict(op_cov,
  A = A, Aprd = A.krig, Y = Y, sigma.e = sigma.e,
  compute.variances = TRUE
)

plot(obs.loc, Y,
  ylab = "u(x)", xlab = "x", main = "Data and prediction",
  ylim = c(

```

```

    min(u.krig$mean - 2 * sqrt(u.krig$variance)),
    max(u.krig$mean + 2 * sqrt(u.krig$variance))
  )
)
lines(x, u.krig$mean)
lines(x, u.krig$mean + 2 * sqrt(u.krig$variance), col = 2)
lines(x, u.krig$mean - 2 * sqrt(u.krig$variance), col = 2)

```

predict.rSPDEobj*Prediction of a fractional SPDE using a rational SPDE approximation***Description**

The function is used for computing kriging predictions based on data $Y_i = u(s_i) + \epsilon_i$, where ϵ is mean-zero Gaussian measurement noise and $u(s)$ is defined by a fractional SPDE $L^\beta u(s) = W$, where W is Gaussian white noise.

Usage

```

## S3 method for class 'rSPDEobj'
predict(
  object,
  A,
  Aprd,
  Y,
  sigma.e,
  compute.variances = FALSE,
  posterior_samples = FALSE,
  n_samples = 100,
  only_latent = FALSE,
  ...
)

```

Arguments

object	The rational SPDE approximation, computed using fractional.operators() , matern.operators() , or spde.matern.operators() .
A	A matrix linking the measurement locations to the basis of the FEM approximation of the latent model.
Aprd	A matrix linking the prediction locations to the basis of the FEM approximation of the latent model.
Y	A vector with the observed data, can also be a matrix where the columns are observations of independent replicates of u .
sigma.e	The standard deviation of the Gaussian measurement noise. Put to zero if the model does not have measurement noise.
compute.variances	Set to also TRUE to compute the kriging variances.

```

posterior_samples
  If TRUE, posterior samples will be returned.
n_samples      Number of samples to be returned. Will only be used if sampling is TRUE.
only_latent    Should the posterior samples be only given to the latent model?
...
  further arguments passed to or from other methods.

```

Value

A list with elements

mean	The kriging predictor (the posterior mean of $u Y$).
variance	The posterior variances (if computed).
samples	A matrix containing the samples if sampling is TRUE.

Examples

```

# Sample a Gaussian Matern process on R using a rational approximation
kappa <- 10
sigma <- 1
nu <- 0.8
sigma.e <- 0.3
range <- sqrt(8*nu)/kappa

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation
op <- matern.operators(
  range = range, sigma = sigma,
  nu = nu, loc_mesh = x, d = 1,
  parameterization = "matern"
)

# Sample the model
u <- simulate(op)

# Create some data
obs.loc <- runif(n = 10, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
Y <- as.vector(A %*% u + sigma.e * rnorm(10))

# compute kriging predictions at the FEM grid
A.krig <- rSPDE.A1d(x, x)
u.krig <- predict(op,
  A = A, Aprd = A.krig, Y = Y, sigma.e = sigma.e,
  compute.variances = TRUE
)

plot(obs.loc, Y,
  ylab = "u(x)", xlab = "x", main = "Data and prediction",

```

```

ylim = c(
  min(u.krig$mean - 2 * sqrt(u.krig$variance)),
  max(u.krig$mean + 2 * sqrt(u.krig$variance))
)
)
lines(x, u.krig$mean)
lines(x, u.krig$mean + 2 * sqrt(u.krig$variance), col = 2)
lines(x, u.krig$mean - 2 * sqrt(u.krig$variance), col = 2)

```

predict.rspde_lme *Prediction of a mixed effects regression model on a metric graph.*

Description

Prediction of a mixed effects regression model on a metric graph.

Usage

```

## S3 method for class 'rspde_lme'
predict(
  object,
  newdata = NULL,
  loc = NULL,
  mesh = FALSE,
  which_repl = NULL,
  compute_variances = FALSE,
  posterior_samples = FALSE,
  n_samples = 100,
  sample_latent = FALSE,
  edge_number = "edge_number",
  distance_on_edge = "distance_on_edge",
  normalized = FALSE,
  return_as_list = FALSE,
  return_original_order = TRUE,
  ...,
  data = deprecated()
)

```

Arguments

- | | |
|---------|---|
| object | The fitted object with the <code>rspde_lme()</code> function |
| newdata | A <code>data.frame</code> or a <code>list</code> containing the covariates, the edge number and the distance on edge for the locations to obtain the prediction. |
| loc | Prediction locations. Can either be a <code>data.frame</code> , a <code>matrix</code> or a character vector, that contains the names of the columns of the coordinates of the locations. For models using <code>metric_graph</code> objects, please use <code>edge_number</code> and <code>distance_on_edge</code> instead. |

<code>mesh</code>	Obtain predictions for mesh nodes? The graph must have a mesh, and either <code>only_latent</code> is set to TRUE or the model does not have covariates.
<code>which_repl</code>	Which replicates to use? If NULL all replicates will be used.
<code>compute_variances</code>	Set to also TRUE to compute the kriging variances.
<code>posterior_samples</code>	If TRUE, posterior samples will be returned.
<code>n_samples</code>	Number of samples to be returned. Will only be used if <code>sampling</code> is TRUE.
<code>sample_latent</code>	Do posterior samples only for the random effects?
<code>edge_number</code>	Name of the variable that contains the edge number, the default is <code>edge_number</code> .
<code>distance_on_edge</code>	Name of the variable that contains the distance on edge, the default is <code>distance_on_edge</code> .
<code>normalized</code>	Are the distances on edges normalized?
<code>return_as_list</code>	Should the means of the predictions and the posterior samples be returned as a list, with each replicate being an element?
<code>return_original_order</code>	Should the results be return in the original (input) order or in the order inside the graph?
<code>...</code>	Not used.
<code>data</code>	[Deprecated] Use <code>newdata</code> instead.

Value

A list with elements `mean`, which contains the means of the predictions, `fe_mean`, which is the prediction for the fixed effects, `re_mean`, which is the prediction for the random effects, `variance` (if `compute_variance` is TRUE), which contains the variances of the predictions, `samples` (if `posterior_samples` is TRUE), which contains the posterior samples.

<code>rational.order</code>	<i>Get the order of rational approximation.</i>
-----------------------------	---

Description

Get the order of rational approximation.

Usage

```
rational.order(object)
```

Arguments

<code>object</code>	A CBrSPDEobj object or an <code>inla_rspde</code> object.
---------------------	---

Value

The order of rational approximation.

rational.order<- *Changing the order of the rational approximation*

Description

Changing the order of the rational approximation

Usage

```
rational.order(x) <- value
```

Arguments

- | | |
|--------------------|--|
| <code>x</code> | A CBrSPDE or an <code>rpsde.inla</code> object |
| <code>value</code> | The order of rational approximation. |

Value

An object of the same class with the new order of rational approximation.

rational.type *Get type of rational approximation.*

Description

Get type of rational approximation.

Usage

```
rational.type(object)
```

Arguments

- | | |
|---------------------|---|
| <code>object</code> | A CBrSPDEobj object or an <code>inla_rspde</code> object. |
|---------------------|---|

Value

The type of rational approximation.

<code>rational.type<-</code>	<i>Changing the type of the rational approximation</i>
---------------------------------	--

Description

Changing the type of the rational approximation

Usage

```
rational.type(x) <- value
```

Arguments

<code>x</code>	A CBrSPDE or an <code>rpsde.inla</code> object
<code>value</code>	The type of rational approximation. The current options are "chebfun", "brasil" and "chebfunLB"

Value

An object of the same class with the new rational approximation.

<code>require.nowarnings</code>	<i>Warnings free loading of add-on packages</i>
---------------------------------	---

Description

Turn off all warnings for `require()`, to allow clean completion of examples that require unavailable Suggested packages.

Usage

```
require.nowarnings(package, lib.loc = NULL, character.only = FALSE)
```

Arguments

<code>package</code>	The name of a package, given as a character string.
<code>lib.loc</code>	a character vector describing the location of R library trees to search through, or <code>NULL</code> . The default value of <code>NULL</code> corresponds to all libraries currently known to <code>.libPaths()</code> . Non-existent library trees are silently ignored.
<code>character.only</code>	a logical indicating whether package can be assumed to be a character string.

Details

`require(package)` acts the same as `require(package, quietly = TRUE)` but with warnings turned off. In particular, no warning or error is given if the package is unavailable. Most cases should use `requireNamespace(package, quietly = TRUE)` instead, which doesn't produce warnings.

Value

`require.nowarnings` returns (invisibly) TRUE if it succeeds, otherwise FALSE

See Also

[require\(\)](#)

Examples

```
## This should produce no output:
if (require.nowarnings(nonexistent)) {
  message("Package loaded successfully")
}
```

rSPDE

Rational approximations of fractional SPDEs.

Description

`rSPDE` is used for approximating fractional elliptic SPDEs

$$L^\beta(\tau u(s)) = W,$$

where L is a differential operator and $\beta > 0$ is a general fractional power.

Details

The approximation is based on a rational approximation of the fractional operator, and allows for computationally efficient inference and simulation.

The main functions for computing rational approximation objects are:

[fractional.operators\(\)](#) works for general rational operators

[matern.operators\(\)](#) works for random fields with stationary Matern covariance functions

[spde.matern.operators\(\)](#) works for random fields with defined as solutions to a possibly non-stationary Matern-type SPDE model.

[rspde.matern\(\)](#) R-INLA implementation of the covariance-based rational approximation for random fields with stationary Matern covariance functions

Basic statistical operations such as likelihood evaluations (see `[rSPDE.loglike]`, `[rSPDE.matern.loglike]`) and kriging predictions (see `[predict.rSPDEobj]`, `[predict.CBrSPDEobj]`) using the rational approximations are also implemented.

For illustration purposes, the package contains a simple FEM implementation for models on R. For spatial models, the FEM implementation in the R-INLA package is recommended.

For a more detailed introduction to the package, see the rSPDE Vignettes.

rSPDE.A1d*Observation matrix for finite element discretization on R*

Description

A finite element discretization on R can be written as $u(s) = \sum_i^n u_i \varphi_i(s)$ where $\varphi_i(s)$ is a piecewise linear "hat function" centered at location x_i . This function computes an $m \times n$ matrix A that links the basis function in the expansion to specified locations $s = (s_1, \dots, s_m)$ in the domain through $A_{ij} = \varphi_j(s_i)$.

Usage

```
rSPDE.A1d(x, loc)
```

Arguments

- | | |
|-----|---|
| x | The locations of the nodes in the FEM discretization. |
| loc | The locations (s_1, \dots, s_m) |

Value

The sparse matrix A .

Author(s)

David Bolin <davidbolin@gmail.com>

See Also

[rSPDE.fem1d\(\)](#)

Examples

```
# create mass and stiffness matrices for a FEM discretization on [0,1]
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# create the observation matrix for some locations in the domain
obs.loc <- runif(n = 10, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
```

`rSPDE.construct.matern.loglike`
Constructor of Matern loglikelihood functions.

Description

This function returns a log-likelihood function for a Gaussian process with a Matern covariance function, that is observed under Gaussian measurement noise: $\bar{Y}_i = u(s_i) + \epsilon_i$, where ϵ_i are iid mean-zero Gaussian variables. The latent model is approximated using the a rational approximation of the fractional SPDE model corresponding to the Gaussian process.

Usage

```
rSPDE.construct.matern.loglike(
  object,
  Y,
  A,
  sigma.e = NULL,
  mu = 0,
  user_nu = NULL,
  user_tau = NULL,
  user_kappa = NULL,
  user_sigma = NULL,
  user_range = NULL,
  parameterization = c("spde", "matern"),
  user_m = NULL,
  log_scale = TRUE,
  return_negative_likelihood = TRUE
)
```

Arguments

<code>object</code>	The rational SPDE approximation, computed using matern.operators()
<code>Y</code>	The observations, either a vector or a matrix where the columns correspond to independent replicates of observations.
<code>A</code>	An observation matrix that links the measurement location to the finite element basis.
<code>sigma.e</code>	IF non-null, the standard deviation of the measurement noise will be kept fixed in the returned likelihood.
<code>mu</code>	Expectation vector of the latent field (default = 0).
<code>user_nu</code>	If non-null, the shape parameter will be kept fixed in the returned likelihood.
<code>user_tau</code>	If non-null, the tau parameter will be kept fixed in the returned likelihood. (Replaces sigma)
<code>user_kappa</code>	If non-null, the range parameter will be kept fixed in the returned likelihood.

user_sigma If non-null, the standard deviation will be kept fixed in the returned likelihood.
 user_range If non-null, the range parameter will be kept fixed in the returned likelihood.
 (Replaces kappa)
 parameterization
 If spde, then one will use the parameters tau and kappa. If matern, then one
 will use the parameters sigma and range.
 user_m If non-null, update the order of the rational approximation, which needs to be a
 positive integer.
 log_scale Should the parameters be evaluated in log-scale?
 return_negative_likelihood
 Return minus the likelihood to turn the maximization into a minimization?

Value

The log-likelihood function. The parameters of the returned function are given in the order sigma, kappa, nu, sigma.e, whenever they are available.

See Also

[matern.operators\(\)](#), [predict.CBrSPDEobj\(\)](#)

Examples

```

# this example illustrates how the function can be used for maximum
# likelihood estimation

set.seed(123)
# Sample a Gaussian Matern process on R using a rational approximation
nu <- 0.8
sigma <- 1
sigma.e <- 0.1
n.rep <- 10
n.obs <- 200
n.x <- 51
range <- 0.2
# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = n.x)
# Compute the covariance-based rational approximation
op_cov <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2,
  parameterization = "matern"
)
# Sample the model
u <- simulate(op_cov, n.rep)
# Create some data
obs.loc <- runif(n = n.obs, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
noise <- rnorm(n.obs * n.rep)
dim(noise) <- c(n.obs, n.rep)

```

```

Y <- as.matrix(A %*% u + sigma.e * noise)

# Define the negative likelihood function for optimization
# using CBrSPDE.matern.loglike
# Matern parameterization
loglike <- rSPDE.construct.matern.loglike(op_cov, Y, A, parameterization = "matern")

# The parameters can now be estimated by minimizing mlik with optim

# Choose some reasonable starting values depending on the size of the domain
theta0 <- c(get.initial.values.rSPDE(mesh.range = 1, dim = 1),
             log(0.1*sd(as.vector(Y))))
# run estimation and display the results
theta <- optim(theta0, loglike,
               method = "L-BFGS-B"
)
print(data.frame(
  sigma = c(sigma, exp(theta$par[1])), range = c(range, exp(theta$par[2])),
  nu = c(nu, exp(theta$par[3])), sigma.e = c(sigma.e, exp(theta$par[4])),
  row.names = c("Truth", "Estimates")
))

```

Description

This function computes mass and stiffness matrices for a FEM approximation on R, assuming Neumann boundary conditions. These matrices are needed when discretizing the operators in rational approximations.

Usage

```
rSPDE.fem1d(x)
```

Arguments

- | | |
|---|--|
| x | Locations of the nodes in the FEM approximation. |
|---|--|

Value

The function returns a list with the following elements

- | | |
|---|-----------------------|
| G | The stiffness matrix. |
| C | The mass matrix. |

Author(s)

David Bolin <davidbolin@gmail.com>

See Also

[rSPDE.A1d\(\)](#)

Examples

```
# create mass and stiffness matrices for a FEM discretization on [0,1]
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)
```

rSPDE.fem2d

Finite element calculations for problems in 2D

Description

This function computes mass and stiffness matrices for a mesh in 2D, assuming Neumann boundary conditions.

Usage

`rSPDE.fem2d(FV, P)`

Arguments

FV	Matrix where each row defines a triangle
P	Locations of the nodes in the mesh.

Value

The function returns a list with the following elements

G	The stiffness matrix with elements $(\nabla\phi_i, \nabla\phi_j)$.
C	The mass matrix with elements (ϕ_i, ϕ_j) .
Cd	The mass lumped matrix with diagonal elements $(\phi_i, 1)$.
Hxx	Matrix with elements $(\partial_x\phi_i, \partial_x\phi_j)$.
Hyx	Matrix with elements $(\partial_y\phi_i, \partial_x\phi_j)$.
Hxy	Matrix with elements $(\partial_x\phi_i, \partial_y\phi_j)$.
Hyx	Matrix with elements $(\partial_y\phi_i, \partial_x\phi_j)$.

Author(s)

David Bolin <davidbolin@gmail.com>

See Also

[rSPDE.fem1d\(\)](#)

Examples

```
P <- rbind(c(0,0), c(1,0), c(1,1), c(0,1))
FV <- rbind(c(1,2,3), c(2,3,4))
fem <- rSPDE.fem2d(FV,P)
```

[rSPDE.loglike](#)

Object-based log-likelihood function for latent Gaussian fractional SPDE model

Description

This function evaluates the log-likelihood function for a fractional SPDE model $L^\beta u(s) = W$ that is observed under Gaussian measurement noise: $Y_i = u(s_i) + \epsilon_i$, where ϵ_i are iid mean-zero Gaussian variables and $x(s) = \mu(s) + u(s)$, where $\mu(s)$ is the expectation vector of the latent field.

Usage

```
rSPDE.loglike(obj, Y, A, sigma.e, mu = 0)
```

Arguments

<code>obj</code>	The rational SPDE approximation, computed using fractional.operators() , matern.operators() , or spde.matern.operators() .
<code>Y</code>	The observations, either a vector or a matrix where the columns correspond to independent replicates of observations.
<code>A</code>	An observation matrix that links the measurement location to the finite element basis.
<code>sigma.e</code>	The standard deviation of the measurement noise.
<code>mu</code>	Expectation vector of the latent field (default = 0).

Value

The log-likelihood value.

Note

This example below shows how the function can be used to evaluate the likelihood of a latent Matern model.

See Also

[spde.matern.loglike\(\)](#)

Examples

```

# Sample a Gaussian Matern process on R using a rational approximation
kappa <- 10
sigma <- 1
nu <- 0.8
sigma.e <- 0.3
range <- 0.2

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation
op <- matern.operators(
  range = range, sigma = sigma, nu = nu,
  loc_mesh = x, d = 1,
  type = "operator", parameterization = "matern"
)

# Sample the model
u <- simulate(op)

# Create some data
obs.loc <- runif(n = 10, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
Y <- as.vector(A %*% u + sigma.e * rnorm(10))

# compute log-likelihood of the data
lik1 <- rSPDE.loglike(op, Y, A, sigma.e)
cat(lik1)

```

rspde.make.A

Observation/prediction matrices for rSPDE models.

Description

Constructs observation/prediction weight matrices for rSPDE models based on `inla.mesh` or `inla.mesh.1d` objects.

Usage

```

rspde.make.A(
  mesh = NULL,
  loc = NULL,
  A = NULL,
  dim = NULL,
  rspde.order = 2,
  nu = NULL,
  index = NULL,

```

```

    group = NULL,
    repl = 1L,
    n.group = NULL,
    n.repl = NULL
)

```

Arguments

mesh	An <code>inla.mesh</code> , an <code>inla.mesh.1d</code> object or a <code>metric_graph</code> object.
loc	Locations, needed if an INLA mesh is provided
A	The A matrix from the standard SPDE approach, such as the matrix returned by <code>inla.spde.make.A</code> . Should only be provided if <code>mesh</code> is not provided.
dim	the dimension. Should only be provided if <code>mesh</code> is not provided.
rspde.order	The order of the covariance-based rational SPDE approach.
nu	If <code>NULL</code> , then the model will assume that <code>nu</code> will be estimated. If <code>nu</code> is fixed, you should provide the value of <code>nu</code> .
index	For each observation/prediction value, an index into <code>loc</code> . Default is <code>seq_len(nrow(A.loc))</code> .
group	For each observation/prediction value, an index into the group model.
repl	For each observation/prediction value, the replicate index.
n.group	The size of the group model.
n.repl	The total number of replicates.

Value

The `A` matrix for rSPDE models.

Examples

```

#tryCatch version
tryCatch({
  if (requireNamespace("INLA", quietly = TRUE)){
    library(INLA)

    set.seed(123)
    loc <- matrix(runif(100 * 2) * 100, 100, 2)
    mesh <- inla.mesh.2d(
      loc = loc,
      cutoff = 50,
      max.edge = c(50, 500)
    )
    A <- rspde.make.A(mesh, loc = loc, rspde.order = 3)
  }
  #stable.tryCatch
}, error = function(e){print("Could not run the example")})

```

`rspde.make.index` *rSPDE model index vector generation*

Description

Generates a list of named index vectors for an rSPDE model.

Usage

```
rspde.make.index(
  name,
  n.spde = NULL,
  n.group = 1,
  n.repl = 1,
  mesh = NULL,
  rspde.order = 2,
  nu = NULL,
  dim = NULL
)
```

Arguments

<code>name</code>	A character string with the base name of the effect.
<code>n.spde</code>	The number of basis functions in the mesh model.
<code>n.group</code>	The size of the group model.
<code>n.repl</code>	The total number of replicates.
<code>mesh</code>	An <code>inla.mesh</code> , an <code>inla.mesh.1d</code> object or a <code>metric_graph</code> object.
<code>rspde.order</code>	The order of the rational approximation
<code>nu</code>	If <code>NULL</code> , then the model will assume that <code>nu</code> will be estimated. If <code>nu</code> is fixed, you should provide the value of <code>nu</code> .
<code>dim</code>	the dimension of the domain. Should only be provided if <code>mesh</code> is not provided.

Value

A list of named index vectors.

<code>name</code>	Indices into the vector of latent variables
<code>name.group</code>	'group' indices
<code>name.repl</code>	Indices for replicates

Examples

```

#tryCatch version
tryCatch({
  if (requireNamespace("INLA", quietly = TRUE)){
    library(INLA)

    set.seed(123)

    m <- 100
    loc_2d_mesh <- matrix(runif(m * 2), m, 2)
    mesh_2d <- inla.mesh.2d(
      loc = loc_2d_mesh,
      cutoff = 0.05,
      max.edge = c(0.1, 0.5)
    )
    sigma <- 1
    range <- 0.2
    nu <- 0.8
    kappa <- sqrt(8 * nu) / range
    op <- matern.operators(
      mesh = mesh_2d, nu = nu,
      range = range, sigma = sigma, m = 2,
      parameterization = "matern"
    )
    u <- simulate(op)
    A <- inla.spde.make.A(
      mesh = mesh_2d,
      loc = loc_2d_mesh
    )
    sigma.e <- 0.1
    y <- A %*% u + rnorm(m) * sigma.e
    Abar <- rspde.make.A(mesh = mesh_2d, loc = loc_2d_mesh)
    mesh.index <- rspde.make.index(name = "field", mesh = mesh_2d)
    st.dat <- inla.stack(
      data = list(y = as.vector(y)),
      A = Abar,
      effects = mesh.index
    )
    rspde_model <- rspde.matern(
      mesh = mesh_2d,
      nu.upper.bound = 2
    )
    f <- y ~ -1 + f(field, model = rspde_model)
    rspde_fit <- inla(f,
      data = inla.stack.data(st.dat),
      family = "gaussian",
      control.predictor =
        list(A = inla.stack.A(st.dat))
    )
    result <- rspde.result(rspde_fit, "field", rspde_model)
    summary(result)
  }
}

```

```
#stable.tryCatch
}, error = function(e){print("Could not run the example")})
```

rspde.matern

Matern rSPDE model object for INLA

Description

Creates an INLA object for a stationary Matern model with general smoothness parameter.

Usage

```
rspde.matern(
  mesh,
  nu.upper.bound = 4,
  rspde.order = 2,
  nu = NULL,
  B.sigma = matrix(c(0, 1, 0), 1, 3),
  B.range = matrix(c(0, 0, 1), 1, 3),
  parameterization = c("spde", "matern", "matern2"),
  B.tau = matrix(c(0, 1, 0), 1, 3),
  B.kappa = matrix(c(0, 0, 1), 1, 3),
  start.nu = NULL,
  start.theta = NULL,
  prior.nu = NULL,
  theta.prior.mean = NULL,
  theta.prior.prec = 0.1,
  prior.std.dev.nominal = 1,
  prior.range.nominal = NULL,
  prior.kappa.mean = NULL,
  prior.tau.mean = NULL,
  start.lstd.dev = NULL,
  start.lrange = NULL,
  start.ltau = NULL,
  start.lkappa = NULL,
  prior.theta.param = c("theta", "spde"),
  prior.nu.dist = c("beta", "lognormal"),
  nu.prec.inc = 1,
  type.rational.approx = c("chebfun", "brasil", "chebfunLB"),
  debug = FALSE,
  shared_lib = "detect",
  ...
)
```

Arguments

<code>mesh</code>	The mesh to build the model. It can be an <code>inla.mesh</code> or an <code>inla.mesh.1d</code> object. Otherwise, should be a list containing elements <code>d</code> , the dimension, <code>C</code> , the mass matrix, and <code>G</code> , the stiffness matrix.
<code>nu.upper.bound</code>	Upper bound for the smoothness parameter.
<code>rspde.order</code>	The order of the covariance-based rational SPDE approach.
<code>nu</code>	If <code>nu</code> is set to a parameter, <code>nu</code> will be kept fixed and will not be estimated. If <code>nu</code> is <code>NULL</code> , it will be estimated.
<code>B.sigma</code>	Matrix with specification of log-linear model for σ (for 'matern' parameterization) or for σ^2 (for 'matern2' parameterization). Will be used if <code>parameterization = 'matern'</code> or <code>parameterization = 'matern2'</code> .
<code>B.range</code>	Matrix with specification of log-linear model for ρ , which is a range-like parameter (it is exactly the range parameter in the stationary case). Will be used if <code>parameterization = 'matern'</code> or <code>parameterization = 'matern2'</code> .
<code>parameterization</code>	Which parameterization to use? <code>matern</code> uses range, std. deviation and <code>nu</code> (smoothness). <code>spde</code> uses <code>kappa</code> , <code>tau</code> and <code>nu</code> (smoothness). <code>matern2</code> uses range-like (1/ <code>kappa</code>), variance and <code>nu</code> (smoothness). The default is <code>spde</code> .
<code>B.tau</code>	Matrix with specification of log-linear model for τ . Will be used if <code>parameterization = 'spde'</code> .
<code>B.kappa</code>	Matrix with specification of log-linear model for κ . Will be used if <code>parameterization = 'spde'</code> .
<code>start.nu</code>	Starting value for <code>nu</code> .
<code>start.theta</code>	Starting values for the model parameters. In the stationary case, if <code>parameterization='matern'</code> , then <code>theta[1]</code> is the std.dev and <code>theta[2]</code> is the range parameter. If <code>parameterization = 'spde'</code> , then <code>theta[1]</code> is <code>tau</code> and <code>theta[2]</code> is <code>kappa</code> .
<code>prior.nu</code>	a list containing the elements <code>mean</code> and <code>prec</code> for beta distribution, or <code>loglocation</code> and <code>logscale</code> for a truncated lognormal distribution. <code>loglocation</code> stands for the location parameter of the truncated lognormal distribution in the log scale. <code>prec</code> stands for the precision of a beta distribution. <code>logscale</code> stands for the scale of the truncated lognormal distribution on the log scale. Check details below.
<code>theta.prior.mean</code>	A vector for the mean priors of <code>theta</code> .
<code>theta.prior.prec</code>	A precision matrix for the prior of <code>theta</code> .
<code>prior.std.dev.nominal</code>	Prior std. deviation to be used for the priors and for the starting values.
<code>prior.range.nominal</code>	Prior range to be used for the priors and for the starting values.
<code>prior.kappa.mean</code>	Prior kappa to be used for the priors and for the starting values.
<code>prior.tau.mean</code>	Prior tau to be used for the priors and for the starting values.

<code>start.lstd.dev</code>	Starting value for log of std. deviation. Will not be used if <code>start.ltau</code> is non-null. Will be only used in the stationary case and if <code>parameterization = 'matern'</code> .
<code>start.lrange</code>	Starting value for log of range. Will not be used if <code>start.lkappa</code> is non-null. Will be only used in the stationary case and if <code>parameterization = 'matern'</code> .
<code>start.ltau</code>	Starting value for log of tau. Will be only used in the stationary case and if <code>parameterization = 'spde'</code> .
<code>start.lkappa</code>	Starting value for log of kappa. Will be only used in the stationary case and if <code>parameterization = 'spde'</code> .
<code>prior.theta.param</code>	Should the lognormal prior be on theta or on the SPDE parameters (tau and kappa on the stationary case)?
<code>prior.nu.dist</code>	The distribution of the smoothness parameter. The current options are "beta" or "lognormal". The default is "lognormal".
<code>nu.prec.inc</code>	Amount to increase the precision in the beta prior distribution. Check details below.
<code>type.rational.approx</code>	Which type of rational approximation should be used? The current types are "chebfun", "brasil" or "chebfunLB".
<code>debug</code>	INLA debug argument
<code>shared_lib</code>	Which shared lib to use for the cgeneric implementation? If "detect", it will check if the shared lib exists locally, in which case it will use it. Otherwise it will use INLA's shared library. If "INLA", it will use the shared lib from INLA's installation. If 'rSPDE', then it will use the local installation (does not work if your installation is from CRAN). Otherwise, you can directly supply the path of the .so (or .dll) file.
<code>...</code>	Only being used internally.
<code>prior.kappa</code>	a list containing the elements <code>meanlog</code> and <code>sdlog</code> , that is, the mean and standard deviation on the log scale.
<code>prior.tau</code>	a list containing the elements <code>meanlog</code> and <code>sdlog</code> , that is, the mean and standard deviation on the log scale.
<code>prior.range</code>	a list containing the elements <code>meanlog</code> and <code>sdlog</code> , that is, the mean and standard deviation on the log scale. Will not be used if <code>prior.kappa</code> is non-null.
<code>prior.std.dev</code>	a list containing the elements <code>meanlog</code> and <code>sdlog</code> , that is, the mean and standard deviation on the log scale. Will not be used if <code>prior.tau</code> is non-null.

Value

An INLA model.

rSPDE.matern.loglike *Object-based log-likelihood function for latent Gaussian fractional SPDE model using the rational approximations*

Description

This function evaluates the log-likelihood function for a Gaussian process with a Matern covariance function, that is observed under Gaussian measurement noise: $Y_i = u(s_i) + \epsilon_i$, where ϵ_i are iid mean-zero Gaussian variables. The latent model is approximated using the a rational approximation of the fractional SPDE model corresponding to the Gaussian process.

Usage

```
rSPDE.matern.loglike(
  object,
  Y,
  A,
  sigma.e,
  mu = 0,
  user_nu = NULL,
  user_kappa = NULL,
  user_sigma = NULL,
  user_range = NULL,
  user_tau = NULL,
  user_m = NULL
)
```

Arguments

object	The rational SPDE approximation, computed using matern.operators()
Y	The observations, either a vector or a matrix where the columns correspond to independent replicates of observations.
A	An observation matrix that links the measurement location to the finite element basis.
sigma.e	The standard deviation of the measurement noise.
mu	Expectation vector of the latent field (default = 0).
user_nu	If non-null, update the shape parameter of the covariance function.
user_kappa	If non-null, update the range parameter of the covariance function.
user_sigma	If non-null, update the standard deviation of the covariance function.
user_range	If non-null, update the range parameter of the covariance function.
user_tau	If non-null, update the parameter tau.
user_m	If non-null, update the order of the rational approximation, which needs to be a positive integer.

Value

The log-likelihood value.

See Also

[matern.operators\(\)](#), [predict.CBrSPDEobj\(\)](#)

Examples

```
# this example illustrates how the function can be used for maximum
# likelihood estimation

set.seed(123)
# Sample a Gaussian Matern process on R using a rational approximation
nu <- 0.8
kappa <- 5
sigma <- 1
sigma.e <- 0.1
n.rep <- 10
n.obs <- 100
n.x <- 51
range <- 0.2

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = n.x)
fem <- rSPDE.fem1d(x)

tau <- sqrt(gamma(nu) / (sigma^2 * kappa^(2 * nu) *
(4 * pi)^(1 / 2) * gamma(nu + 1 / 2)))

# Compute the covariance-based rational approximation
op_cov <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2,
  parameterization = "matern"
)

# Sample the model
u <- simulate(op_cov, n.rep)

# Create some data
obs.loc <- runif(n = n.obs, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
noise <- rnorm(n.obs * n.rep)
dim(noise) <- c(n.obs, n.rep)
Y <- as.matrix(A %*% u + sigma.e * noise)

# Define the negative likelihood function for optimization
# using CBrSPDE.matern.loglike

# Notice that we are also using sigma instead of tau, so it can be compared
# to matern.loglike()
```

```

mlik_cov <- function(theta, Y, A, op_cov) {
  kappa <- exp(theta[1])
  sigma <- exp(theta[2])
  nu <- exp(theta[3])
  return(-rSPDE.matern.loglike(
    object = op_cov, Y = Y,
    A = A, user_kappa = kappa, user_sigma = sigma,
    user_nu = nu, sigma.e = exp(theta[4]))
  ))
}

# The parameters can now be estimated by minimizing mlik with optim

# Choose some reasonable starting values depending on the size of the domain
theta0 <- log(c(sqrt(8), 1 / sqrt(var(c(Y))), 0.9, 0.01))

# run estimation and display the results
theta <- optim(theta0, mlik_cov,
  Y = Y, A = A, op_cov = op_cov,
  method = "L-BFGS-B"
)

print(data.frame(
  range = c(range, exp(theta$par[1])), sigma = c(sigma, exp(theta$par[2])),
  nu = c(nu, exp(theta$par[3])), sigma.e = c(sigma.e, exp(theta$par[4])),
  row.names = c("Truth", "Estimates")
))

```

rspde.matern.precision

Precision matrix of the covariance-based rational approximation of stationary Gaussian Matern random fields

Description

`rspde.matern.precision` is used for computing the precision matrix of the covariance-based rational SPDE approximation of a stationary Gaussian random fields on R^d with a Matern covariance function

$$C(h) = \frac{\sigma^2}{2(\nu - 1)\Gamma(\nu)} (\kappa h)^\nu K_\nu(\kappa h)$$

Usage

```
rspde.matern.precision(
  kappa,
  nu,
  tau = NULL,
```

```

sigma = NULL,
rspde.order,
dim,
fem_mesh_matrices,
only_fractional = FALSE,
return_block_list = FALSE,
type_rational_approx = "chebfun"
)

```

Arguments

kappa	Range parameter of the covariance function.
nu	Shape parameter of the covariance function.
tau	Scale parameter of the covariance function. If sigma is not provided, tau should be provided.
sigma	Standard deviation of the covariance function. If tau is not provided, sigma should be provided.
rspde.order	The order of the rational approximation
dim	The dimension of the domain
fem_mesh_matrices	A list containing the FEM-related matrices. The list should contain elements c0, g1, g2, g3, etc.
only_fractional	Logical. Should only the fractional-order part of the precision matrix be returned?
return_block_list	Logical. For type = "covariance", should the block parts of the precision matrix be returned separately as a list?
type_rational_approx	Which type of rational approximation should be used? The current types are "chebfun", "brasil" or "chebfunLB".

Value

The precision matrix

Examples

```

set.seed(123)
nobs <- 101
x <- seq(from = 0, to = 1, length.out = nobs)
fem <- rSPDE.fem1d(x)
kappa <- 40
sigma <- 1
d <- 1
nu <- 2.6
tau <- sqrt(gamma(nu) / (kappa^(2 * nu) * (4 * pi)^(d / 2) *
gamma(nu + d / 2)))

```

```

range <- sqrt(8*nu)/kappa
op_cov <- matern.operators(
  loc_mesh = x, nu = nu, range = range, sigma = sigma,
  d = 1, m = 2, compute_higher_order = TRUE,
  parameterization = "matern"
)
v <- t(rSPDE.A1d(x, 0.5))
c.true <- matern.covariance(abs(x - 0.5), kappa, nu, sigma)
Q <- rspde.matern.precision(
  kappa = kappa, nu = nu, tau = tau, rspde.order = 2, d = 1,
  fem_mesh_matrices = op_cov$fem_mesh_matrices
)
A <- Diagonal(nobs)
Abar <- cbind(A, A, A)
w <- rbind(v, v, v)
c.approx_cov <- (Abar) %*% solve(Q, w)

# plot the result and compare with the true Matern covariance
plot(x, matern.covariance(abs(x - 0.5), kappa, nu, sigma),
  type = "l", ylab = "C(h)",
  xlab = "h", main = "Matern covariance and rational approximations"
)
lines(x, c.approx_cov, col = 2)

```

rspde.matern.precision.integer

Precision matrix of stationary Gaussian Matern random fields with integer covariance exponent

Description

`rspde.matern.precision.integer.opt` is used for computing the precision matrix of stationary Gaussian random fields on R^d with a Matern covariance function

$$C(h) = \frac{\sigma^2}{2(\nu - 1)\Gamma(\nu)} (\kappa h)^\nu K_\nu(\kappa h)$$

, where $\alpha = \nu + d/2$ is a natural number.

Usage

```

rspde.matern.precision.integer(
  kappa,
  nu,
  tau = NULL,
  sigma = NULL,
  dim,
  fem_mesh_matrices
)

```

Arguments

kappa	Range parameter of the covariance function.
nu	Shape parameter of the covariance function.
tau	Scale parameter of the covariance function.
sigma	Standard deviation of the covariance function. If tau is not provided, sigma should be provided.
dim	The dimension of the domain
fem_mesh_matrices	A list containing the FEM-related matrices. The list should contain elements c0, g1, g2, g3, etc.

Value

The precision matrix

Examples

```

set.seed(123)
nobs <- 101
x <- seq(from = 0, to = 1, length.out = nobs)
fem <- rSPDE.fem1d(x)
kappa <- 40
sigma <- 1
d <- 1
nu <- 0.5
tau <- sqrt(gamma(nu) / (kappa^(2 * nu) *
(4 * pi)^(d / 2) * gamma(nu + d / 2)))
range <- sqrt(8*nu)/kappa
op_cov <- matern.operators(
  loc_mesh = x, nu = nu, range = range, sigma = sigma,
  d = 1, m = 2, parameterization = "matern"
)
v <- t(rSPDE.A1d(x, 0.5))
c.true <- matern.covariance(abs(x - 0.5), kappa, nu, sigma)
Q <- rspde.matern.precision.integer(
  kappa = kappa, nu = nu, tau = tau, d = 1,
  fem_mesh_matrices = op_cov$fem_mesh_matrices
)
A <- Diagonal(nobs)
c.approx_cov <- A %*% solve(Q, v)

# plot the result and compare with the true Matern covariance
plot(x, matern.covariance(abs(x - 0.5), kappa, nu, sigma),
  type = "l", ylab = "C(h)",
  xlab = "h", main = "Matern covariance and rational approximations"
)
lines(x, c.approx_cov, col = 2)

```

rspde.matern.precision.integer.opt

Optimized precision matrix of stationary Gaussian Matern random fields with integer covariance exponent

Description

rspde.matern.precision.integer.opt is used for computing the optimized version of the precision matrix of stationary Gaussian random fields on R^d with a Matern covariance function

$$C(h) = \frac{\sigma^2}{2^{\nu-1}\Gamma(\nu)}(\kappa h)^\nu K_\nu(\kappa h),$$

where $\alpha = \nu + d/2$ is a natural number.

Usage

```
rspde.matern.precision.integer.opt(
  kappa,
  nu,
  tau,
  d,
  fem_matrices,
  graph = NULL
)
```

Arguments

<code>kappa</code>	Range parameter of the covariance function.
<code>nu</code>	Shape parameter of the covariance function.
<code>tau</code>	Scale parameter of the covariance function.
<code>d</code>	The dimension of the domain
<code>fem_matrices</code>	A list containing the FEM-related matrices. The list should contain elements C, G, G_2, G_3, etc.
<code>graph</code>	The sparsity graph of the matrices. If NULL, only a vector of the elements will be returned, if non-NUL, a sparse matrix will be returned.

Value

The precision matrix

rspde.matern.precision.opt

Optimized precision matrix of the covariance-based rational approximation

Description

`rspde.matern.precision` is used for computing the optimized version of the precision matrix of the covariance-based rational SPDE approximation of a stationary Gaussian random fields on R^d with a Matern covariance function

$$C(h) = \frac{\sigma^2}{2^{\nu-1}\Gamma(\nu)}(\kappa h)^\nu K_\nu(\kappa h).$$

Usage

```
rspde.matern.precision.opt(
  kappa,
  nu,
  tau,
  rspde.order,
  dim,
  fem_matrices,
  graph = NULL,
  sharp,
  type_rational_approx
)
```

Arguments

<code>kappa</code>	Range parameter of the covariance function.
<code>nu</code>	Shape parameter of the covariance function.
<code>tau</code>	Scale parameter of the covariance function.
<code>rspde.order</code>	The order of the rational approximation
<code>dim</code>	The dimension of the domain
<code>fem_matrices</code>	A list containing the FEM-related matrices. The list should contain elements C, G, G_2, G_3, etc.
<code>graph</code>	The sparsity graph of the matrices. If <code>NULL</code> , only a vector of the elements will be returned, if non- <code>NULL</code> , a sparse matrix will be returned.
<code>sharp</code>	The sparsity graph should have the correct sparsity (costs more to perform a sparsity analysis) or an upper bound for the sparsity?
<code>type_rational_approx</code>	Which type of rational approximation should be used? The current types are "chebfun", "brasil" or "chebfunLB".

Value

The precision matrix

rspde.mesh.project *Calculate a lattice projection to/from an `inla.mesh` for rSPDE objects*

Description

Calculate a lattice projection to/from an `inla.mesh` for rSPDE objects

Usage

```
rspde.mesh.project(...)

rspde.mesh.projector(
  mesh,
  nu = NULL,
  rspde.order = 2,
  loc = NULL,
  lattice = NULL,
  xlim = NULL,
  ylim = NULL,
  dims = c(100, 100),
  projection = NULL,
  ...
)

## S3 method for class 'inla.mesh'
rspde.mesh.project(
  mesh,
  loc = NULL,
  field = NULL,
  rspde.order = 2,
  nu = NULL,
  ...
)

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

## S3 method for class 'inla.mesh.1d'
rspde.mesh.project(mesh, loc, field = NULL, rspde.order = 2, nu = NULL, ...)
```

Arguments

... Additional parameters.

<code>mesh</code>	An <code>inla.mesh</code> or <code>inla.mesh.1d</code> object.
<code>nu</code>	The smoothness parameter. If <code>NULL</code> , it will be assumed that <code>nu</code> was estimated.
<code>rspde.order</code>	The order of the rational approximation.
<code>loc</code>	Projection locations. Can be a matrix or a <code>SpatialPoints</code> or a <code>SpatialPointsDataFrame</code> object.
<code>lattice</code>	An <code>inla.mesh.lattice</code> object.
<code>xlim</code>	X-axis limits for a lattice. For R2 meshes, defaults to covering the domain.
<code>ylim</code>	Y-axis limits for a lattice. For R2 meshes, defaults to covering the domain.
<code>dims</code>	Lattice dimensions.
<code>projection</code>	One of <code>c("default", "longlat", "longsinlat", "mollweide")</code> .
<code>field</code>	Basis function weights, one per mesh basis function, describing the function to be evaluated at the projection locations.
<code>projector</code>	A <code>rspde.mesh.projector</code> object.

Details

This function is built upon the `inla.mesh.project` and `inla.mesh.projector` functions from INLA.

Value

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

`rspde.metric_graph` *Matern rSPDE model object for metric graphs in INLA*

Description

Creates an INLA object for a stationary Matern model on a metric graph with general smoothness parameter.

Usage

```
rspde.metric_graph(
  graph_obj,
  h = NULL,
  nu.upper.bound = 2,
  rspde.order = 2,
  nu = NULL,
  debug = FALSE,
  B.sigma = matrix(c(0, 1, 0), 1, 3),
  B.range = matrix(c(0, 0, 1), 1, 3),
  parameterization = c("matern", "spde"),
```

```

B.tau = matrix(c(0, 1, 0), 1, 3),
B.kappa = matrix(c(0, 0, 1), 1, 3),
start.nu = NULL,
start.theta = NULL,
prior.nu = NULL,
theta.prior.mean = NULL,
theta.prior.prec = 0.1,
prior.std.dev.nominal = 1,
prior.range.nominal = NULL,
prior.kappa.mean = NULL,
prior.tau.mean = NULL,
start.lstd.dev = NULL,
start.lrange = NULL,
start.ltau = NULL,
start.lkappa = NULL,
prior.theta.param = c("theta", "spde"),
prior.nu.dist = c("lognormal", "beta"),
nu.prec.inc = 1,
type.rational.approx = c("chebfun", "brasil", "chebfunLB"),
shared_lib = "INLA"
)

```

Arguments

<code>graph_obj</code>	The graph object to build the model. Needs to be of class <code>metric_graph</code> . It should have a built mesh. If the mesh is not built, one will be built using <code>h=0.01</code> as default.
<code>h</code>	The width of the mesh in case the mesh was not built.
<code>nu.upper bound</code>	Upper bound for the smoothness parameter.
<code>rspde.order</code>	The order of the covariance-based rational SPDE approach.
<code>nu</code>	If <code>nu</code> is set to a parameter, <code>nu</code> will be kept fixed and will not be estimated. If <code>nu</code> is <code>NULL</code> , it will be estimated.
<code>debug</code>	INLA debug argument
<code>B.sigma</code>	Matrix with specification of log-linear model for σ . Will be used if <code>parameterization = 'matern'</code> .
<code>B.range</code>	Matrix with specification of log-linear model for ρ , which is a range-like parameter (it is exactly the range parameter in the stationary case). Will be used if <code>parameterization = 'matern'</code> .
<code>parameterization</code>	Which parameterization to use? <code>matern</code> uses <code>range</code> , <code>std.deviation</code> and <code>nu</code> (smoothness). <code>spde</code> uses <code>kappa</code> , <code>tau</code> and <code>nu</code> (smoothness). The default is <code>matern</code> .
<code>B.tau</code>	Matrix with specification of log-linear model for τ . Will be used if <code>parameterization = 'spde'</code> .
<code>B.kappa</code>	Matrix with specification of log-linear model for κ . Will be used if <code>parameterization = 'spde'</code> .
<code>start.nu</code>	Starting value for <code>nu</code> .

<code>start.theta</code>	Starting values for the model parameters. In the stationary case, if <code>parameterization='matern'</code> , then <code>theta[1]</code> is the std.dev and <code>theta[2]</code> is the range parameter. If <code>parameterization = 'spde'</code> , then <code>theta[1]</code> is tau and <code>theta[2]</code> is kappa.
<code>prior.nu</code>	a list containing the elements <code>mean</code> and <code>prec</code> for beta distribution, or <code>loglocation</code> and <code>logscale</code> for a truncated lognormal distribution. <code>loglocation</code> stands for the location parameter of the truncated lognormal distribution in the log scale. <code>prec</code> stands for the precision of a beta distribution. <code>logscale</code> stands for the scale of the truncated lognormal distribution on the log scale. Check details below.
<code>theta.prior.mean</code>	A vector for the mean priors of theta.
<code>theta.prior.prec</code>	A precision matrix for the prior of theta.
<code>prior.std.dev.nominal</code>	Prior std. deviation to be used for the priors and for the starting values.
<code>prior.range.nominal</code>	Prior range to be used for the priors and for the starting values.
<code>prior.kappa.mean</code>	Prior kappa to be used for the priors and for the starting values.
<code>prior.tau.mean</code>	Prior tau to be used for the priors and for the starting values.
<code>start.lstd.dev</code>	Starting value for log of std. deviation. Will not be used if <code>start.ltau</code> is non-null. Will be only used in the stationary case and if <code>parameterization = 'matern'</code> .
<code>start.lrange</code>	Starting value for log of range. Will not be used if <code>start.lkappa</code> is non-null. Will be only used in the stationary case and if <code>parameterization = 'matern'</code> .
<code>start.ltau</code>	Starting value for log of tau. Will be only used in the stationary case and if <code>parameterization = 'spde'</code> .
<code>start.lkappa</code>	Starting value for log of kappa. Will be only used in the stationary case and if <code>parameterization = 'spde'</code> .
<code>prior.theta.param</code>	Should the lognormal prior be on theta or on the SPDE parameters (tau and kappa on the stationary case)?
<code>prior.nu.dist</code>	The distribution of the smoothness parameter. The current options are "beta" or "lognormal". The default is "lognormal".
<code>nu.prec.inc</code>	Amount to increase the precision in the beta prior distribution. Check details below.
<code>type.rational.approx</code>	Which type of rational approximation should be used? The current types are "chebfun", "brasil" or "chebfunLB".
<code>shared_lib</code>	Which shared lib to use for the cgeneric implementation? If "INLA", it will use the shared lib from INLA's installation. If 'rSPDE', then it will use the local installation (does not work if your installation is from CRAN). Otherwise, you can directly supply the path of the .so (or .dll) file.
<code>prior.kappa</code>	a list containing the elements <code>meanlog</code> and <code>sdlog</code> , that is, the mean and standard deviation on the log scale.

<code>prior.tau</code>	a list containing the elements <code>meanlog</code> and <code>sdlog</code> , that is, the mean and standard deviation on the log scale.
<code>prior.range</code>	a list containing the elements <code>meanlog</code> and <code>sdlog</code> , that is, the mean and standard deviation on the log scale. Will not be used if <code>prior.kappa</code> is non-null.
<code>prior.std.dev</code>	a list containing the elements <code>meanlog</code> and <code>sdlog</code> , that is, the mean and standard deviation on the log scale. Will not be used if <code>prior.tau</code> is non-null.

Value

An INLA model.

`rspde.result`*rSPDE result extraction from INLA estimation results***Description**

Extract field and parameter values and distributions for an `rspde` effect from an `inla` result object.

Usage

```
rspde.result(
  inla,
  name,
  rspde,
  compute.summary = TRUE,
  parameterization = "detect",
  n_samples = 5000,
  n_density = 1024
)
```

Arguments

<code>inla</code>	An <code>inla</code> object obtained from a call to <code>inla()</code> .
<code>name</code>	A character string with the name of the rSPDE effect in the <code>inla</code> formula.
<code>rspde</code>	The <code>inla_rspde</code> object used for the effect in the <code>inla</code> formula.
<code>compute.summary</code>	Should the summary be computed?
<code>parameterization</code>	If 'detect', the parameterization from the model will be used. Otherwise, the options are 'spde', 'matern' and 'matern2'.
<code>n_samples</code>	The number of samples to be used if <code>parameterization</code> is different from the one used to fit the model.
<code>n_density</code>	The number of equally spaced points to estimate the density.

Value

If the model was fitted with `matern` parameterization (the default), it returns a list containing:

```

marginals.range      Marginal densities for the range parameter
marginals.log.range   Marginal densities for log(range)
marginals.std.dev     Marginal densities for std. deviation
marginals.log.std.dev Marginal densities for log(std. deviation)
marginals.values       Marginal densities for the field values
summary.log.range     Summary statistics for log(range)
summary.log.std.dev    Summary statistics for log(std. deviation)
summary.values        Summary statistics for the field values

If compute.summary is TRUE, then the list will also contain

summary.kappa    Summary statistics for kappa
summary.tau       Summary statistics for tau

```

If the model was fitted with the `spde` parameterization, it returns a list containing:

```

marginals.kappa      Marginal densities for kappa
marginals.log.kappa    Marginal densities for log(kappa)
marginals.log.tau      Marginal densities for log(tau)
marginals.tau        Marginal densities for tau
marginals.values       Marginal densities for the field values
summary.log.kappa     Summary statistics for log(kappa)
summary.log.tau       Summary statistics for log(tau)
summary.values        Summary statistics for the field values

```

If `compute.summary` is TRUE, then the list will also contain

```

summary.kappa    Summary statistics for kappa
summary.tau       Summary statistics for tau

```

For both cases, if `nu` was estimated, then the list will also contain

```

marginals.nu      Marginal densities for nu

If nu was estimated and a beta prior was used, then the list will also contain

marginals.logit.nu
                      Marginal densities for logit(nu)
summary.logit.nu
                      Marginal densities for logit(nu)

If nu was estimated and a truncated lognormal prior was used, then the list will also contain

marginals.log.nu
                      Marginal densities for log(nu)
summary.log.nu Marginal densities for log(nu)

If nu was estimated and compute.summary is TRUE, then the list will also contain

summary.nu       Summary statistics for nu

```

Examples

```

#tryCatch version
tryCatch({
  if (requireNamespace("INLA", quietly = TRUE)){
    library(INLA)

    set.seed(123)

    m <- 100
    loc_2d_mesh <- matrix(runif(m * 2), m, 2)
    mesh_2d <- inla.mesh.2d(
      loc = loc_2d_mesh,
      cutoff = 0.05,
      max.edge = c(0.1, 0.5)
    )
    sigma <- 1
    range <- 0.2
    nu <- 0.8
    kappa <- sqrt(8 * nu) / range
    op <- matern.operators(
      mesh = mesh_2d, nu = nu,
      range = range, sigma = sigma, m = 2,
      parameterization = "matern"
    )
    u <- simulate(op)
    A <- inla.spde.make.A(
      mesh = mesh_2d,
      loc = loc_2d_mesh
    )
    sigma.e <- 0.1
    y <- A %*% u + rnorm(m) * sigma.e
    Abar <- rspde.make.A(mesh = mesh_2d, loc = loc_2d_mesh)
    mesh.index <- rspde.make.index(name = "field", mesh = mesh_2d)
  }
}

```

```

st.dat <- inla.stack(
  data = list(y = as.vector(y)),
  A = Abar,
  effects = mesh.index
)
rspde_model <- rspde.matern(
  mesh = mesh_2d,
  nu.upper.bound = 2
)
f <- y ~ -1 + f(field, model = rspde_model)
rspde_fit <- inla(f,
  data = inla.stack.data(st.dat),
  family = "gaussian",
  control.predictor =
    list(A = inla.stack.A(st.dat)))
)
result <- rspde.result(rspde_fit, "field", rspde_model)
summary(result)
}
#stable.tryCatch
}, error = function(e){print("Could not run the example")})

```

rspde_lme

rSPDE linear mixed effects models

Description

Fitting linear mixed effects model with latent Whittle-Matern models.

Usage

```

rspde_lme(
  formula,
  loc,
  data,
  model = NULL,
  repl = NULL,
  which_repl = NULL,
  optim_method = "L-BFGS-B",
  use_data_from_graph = TRUE,
  starting_values_latent = NULL,
  start_sigma_e = NULL,
  start_alpha = NULL,
  alpha = NULL,
  start_nu = NULL,
  nu = NULL,
  nu_upper_bound = 4,
  rspde_order = NULL,

```

```

parallel = FALSE,
n_cores = parallel::detectCores() - 1,
optim_controls = list(),
improve_hessian = FALSE,
hessian_args = list()
)

```

Arguments

<code>formula</code>	Formula object describing the relation between the response variables and the fixed effects. If the response variable is a matrix, each column of the matrix will be treated as a replicate.
<code>loc</code>	A vector with the names of the columns in <code>data</code> that contain the observation locations, or a <code>matrix</code> or a <code>data.frame</code> containing the observation locations. If the model is of class <code>metric_graph</code> , the locations must be either a <code>matrix</code> or a <code>data.frame</code> with two columns, or a character vector with the names of the two columns. The first column being the number of the edge, and the second column being the normalized position on the edge. If the model is a 2d model, <code>loc</code> must be either a <code>matrix</code> or <code>data.frame</code> with two columns or a character vector with the name of the two columns that contain the location, the first entry corresponding to the <code>x</code> entry and the second corresponding to the <code>y</code> entry.
<code>data</code>	A <code>data.frame</code> containing the data to be used.
<code>model</code>	Either an object generated by <code>matern.operators()</code> or <code>spde.matern.operators()</code> . If <code>NULL</code> , a simple linear regression will be performed.
<code>repl</code>	Vector indicating the replicate of each observation. If <code>NULL</code> it will assume there is only one replicate.
<code>which_repl</code>	Which replicates to use? If <code>NULL</code> all replicates will be used.
<code>optim_method</code>	The method to be used with <code>optim</code> function.
<code>use_data_from_graph</code>	Logical. Only for models generated from graphs from <code>metric_graph</code> class. In this case, should the data, the locations and the replicates be obtained from the graph object?
<code>starting_values_latent</code>	A vector containing the starting values for the latent model. If the latent model was generated by <code>matern.operators()</code> , then the starting values should be provided as a vector of the form <code>c(tau,kappa)</code> . If the model was generated by <code>spde.matern.operators()</code> , then the starting values should be provided as a vector containing the nonstationary parameters.
<code>start_sigma_e</code>	Starting value for the standard deviation of the measurement error.
<code>start_alpha</code>	Starting value for the smoothness parameter. Will be used if <code>start_nu</code> is not given.
<code>alpha</code>	If <code>NULL</code> , the smoothness parameter will be estimated, otherwise the smoothness parameter will be kept fixed at the provided value. Will be used if <code>nu</code> is not given.
<code>start_nu</code>	Starting value for the smoothness parameter.

nu	If NULL, the smoothness parameter will be estimated, otherwise the smoothness parameter will be kept fixed at the provided value.
nu_upper_bound	A parameter that limits the maximum value that nu can assume.
rspde_order	The order of the rational approximation to be used while fitting the model. If not given, the order from the model object will be used.
parallel	logical. Indicating whether to use optimParallel or not.
n_cores	Number of cores to be used if parallel is true.
optim_controls	Additional controls to be passed to optim or optimParallel.
improve_hessian	Should a more precise estimate of the hessian be obtained? Turning on might increase the overall time.
hessian_args	List of controls to be used if improve_hessian is TRUE. The list can contain the arguments to be passed to the method.args argument in the numDeriv::hessian function. See the help of the hessian function in numDeriv package for details. Observe that it only accepts the "Richardson" method for now, the method "complex" is not supported.

Value

A list containing the fitted model.

simulate.CBrSPDEobj	<i>Simulation of a fractional SPDE using the covariance-based rational SPDE approximation</i>
---------------------	---

Description

The function samples a Gaussian random field based using the covariance-based rational SPDE approximation.

Usage

```
## S3 method for class 'CBrSPDEobj'
simulate(
  object,
  nsim = 1,
  seed = NULL,
  user_nu = NULL,
  user_kappa = NULL,
  user_sigma = NULL,
  user_range = NULL,
  user_tau = NULL,
  user_theta = NULL,
  user_m = NULL,
  ...
)
```

Arguments

<code>object</code>	The covariance-based rational SPDE approximation, computed using matern.operators()
<code>nsim</code>	The number of simulations.
<code>seed</code>	An object specifying if and how the random number generator should be initialized ('seeded').
<code>user_nu</code>	If non-null, update the shape parameter of the covariance function.
<code>user_kappa</code>	If non-null, update the range parameter of the covariance function.
<code>user_sigma</code>	If non-null, update the standard deviation of the covariance function.
<code>user_range</code>	If non-null, update the range parameter of the covariance function.
<code>user_tau</code>	If non-null, update the parameter tau.
<code>user_theta</code>	For non-stationary models. If non-null, update the vector of parameters.
<code>user_m</code>	If non-null, update the order of the rational approximation, which needs to be a positive integer.
<code>...</code>	Currently not used.

Value

A matrix with the `n` samples as columns.

Examples

```
# Sample a Gaussian Matern process on R using a rational approximation
kappa <- 10
sigma <- 1
nu <- 0.8
range <- sqrt(8*nu)/kappa

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation of covariance function at 0.5
tau <- sqrt(gamma(nu) / (sigma^2 * kappa^(2 * nu) *
(4 * pi)^(1 / 2) * gamma(nu + 1 / 2)))
op_cov <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2,
  parameterization = "matern"
)

# Sample the model and plot the result
Y <- simulate(op_cov)
plot(x, Y, type = "l", ylab = "u(x)", xlab = "x")
```

<code>simulate.rSPDEobj</code>	<i>Simulation of a fractional SPDE using a rational SPDE approximation</i>
--------------------------------	--

Description

The function samples a Gaussian random field based on a pre-computed rational SPDE approximation.

Usage

```
## S3 method for class 'rSPDEobj'
simulate(object, nsim = 1, seed = NULL, ...)
```

Arguments

<code>object</code>	The rational SPDE approximation, computed using fractional.operators() , matern.operators() , or spde.matern.operators() .
<code>nsim</code>	The number of simulations.
<code>seed</code>	an object specifying if and how the random number generator should be initialized ('seeded').
<code>...</code>	Currently not used.

Value

A matrix with the `n` samples as columns.

See Also

[simulate.CBrSPDEobj\(\)](#)

Examples

```
# Sample a Gaussian Matern process on R using a rational approximation
kappa <- 10
sigma <- 1
nu <- 0.8
range <- sqrt(8*nu)/kappa

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation
op <- matern.operators(
  range = range, sigma = sigma,
  nu = nu, loc_mesh = x, d = 1,
  parameterization = "matern"
)
```

```
# Sample the model and plot the result
Y <- simulate(op)
plot(x, Y, type = "l", ylab = "u(x)", xlab = "x")
```

spde.make.A

Observation/prediction matrices for rSPDE models with integer smoothness.

Description

Constructs observation/prediction weight matrices for rSPDE models with integer smoothness based on `inla.mesh` or `inla.mesh.1d` objects.

Usage

```
spde.make.A(
  mesh = NULL,
  loc = NULL,
  A = NULL,
  index = NULL,
  group = NULL,
  repl = 1L,
  n.group = NULL,
  n.repl = NULL
)
```

Arguments

<code>mesh</code>	An <code>inla.mesh</code> , an <code>inla.mesh.1d</code> object or a <code>metric_graph</code> object.
<code>loc</code>	Locations, needed if an INLA mesh is provided
<code>A</code>	The <code>A</code> matrix from the standard SPDE approach, such as the matrix returned by <code>inla.spde.make.A</code> . Should only be provided if <code>mesh</code> is not provided.
<code>index</code>	For each observation/prediction value, an index into <code>loc</code> . Default is <code>seq_len(nrow(A.loc))</code> .
<code>group</code>	For each observation/prediction value, an index into the group model.
<code>repl</code>	For each observation/prediction value, the replicate index.
<code>n.group</code>	The size of the group model.
<code>n.repl</code>	The total number of replicates.

Value

The `A` matrix for rSPDE models.

Examples

```
#tryCatch version
tryCatch({
  if (requireNamespace("fmesher", quietly = TRUE)){
    library(fmesher)

    set.seed(123)
    loc <- matrix(runif(100 * 2) * 100, 100, 2)
    mesh <- fm_mesh_2d(
      loc = loc,
      cutoff = 50,
      max.edge = c(50, 500)
    )
    A <- spde.make.A(mesh, loc = loc)
  }
  #stable.tryCatch
}, error = function(e){print("Could not run the example")})
```

spde.matern.loglike *Parameter-based log-likelihood for a latent Gaussian Matern SPDE model using a rational SPDE approximation*

Description

This function evaluates the log-likelihood function for observations of a Gaussian process defined as the solution to the SPDE

$$(\kappa(s) - \Delta)^\beta (\tau(s)u(s)) = W.$$

Usage

```
spde.matern.loglike(
  object,
  Y,
  A,
  sigma.e,
  mu = 0,
  user_nu = NULL,
  user_kappa = NULL,
  user_tau = NULL,
  user_theta = NULL,
  user_m = NULL
)
```

Arguments

object	The rational SPDE approximation, computed using spde.matern.operators()
--------	---

Y	The observations, either a vector or a matrix where the columns correspond to independent replicates of observations.
A	An observation matrix that links the measurement location to the finite element basis.
sigma.e	IF non-null, the standard deviation of the measurement noise will be kept fixed in the returned likelihood.
mu	Expectation vector of the latent field (default = 0).
user_nu	If non-null, the shape parameter will be kept fixed in the returned likelihood.
user_kappa	If non-null, updates the range parameter.
user_tau	If non-null, updates the parameter tau.
user_theta	If non-null, updates the parameter theta (that connects tau and kappa to the model matrices in object).
user_m	If non-null, update the order of the rational approximation, which needs to be a positive integer.

Details

The observations are assumed to be generated as $Y_i = u(s_i) + \epsilon_i$, where ϵ_i are iid mean-zero Gaussian variables. The latent model is approximated using a rational approximation of the fractional SPDE model.

Value

The log-likelihood value.

See Also

[rSPDE.loglike\(\)](#).

Examples

```
# this example illustrates how the function can be used for maximum
# likelihood estimation
# Sample a Gaussian Matern process on R using a rational approximation
sigma.e <- 0.1
n.rep <- 10
n.obs <- 100
n.x <- 51
# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = n.x)
fem <- rSPDE.fem1d(x)
tau <- rep(0.5, n.x)
nu <- 0.8
alpha <- nu + 1/2
kappa <- rep(1, n.x)
# compute rational approximation
op <- spde.matern.operators(
  kappa = kappa, tau = tau, alpha = alpha,
```

```

parameterization = "spde", d = 1,
loc_mesh = x
)
# Sample the model
u <- simulate(op, n.rep)
# Create some data
obs.loc <- runif(n = n.obs, min = 0, max = 1)
A <- rSPDE.A1d(x, obs.loc)
noise <- rnorm(n.obs * n.rep)
dim(noise) <- c(n.obs, n.rep)
Y <- as.matrix(A %*% u + sigma.e * noise)
# define negative likelihood function for optimization using matern.loglike
mlik <- function(theta) {
  return(-spde.matern.loglike(op, Y, A, sigma.e = exp(theta[4]),
                               user_nu = exp(theta[3]),
                               user_kappa = exp(theta[2]),
                               user_tau = exp(theta[1])))
}
#' #The parameters can now be estimated by minimizing mlik with optim

# Choose some reasonable starting values depending on the size of the domain
theta0 <- log(c(1 / sqrt(var(c(Y))), sqrt(8), 0.9, 0.01))
# run estimation and display the results
theta <- optim(theta0, mlik)
print(data.frame(
  tau = c(tau[1], exp(theta$par[1])), kappa = c(kappa[1], exp(theta$par[2])),
  nu = c(nu, exp(theta$par[3])), sigma.e = c(sigma.e, exp(theta$par[4])),
  row.names = c("Truth", "Estimates")
))

```

spde.matern.operators *Rational approximations of non-stationary Gaussian SPDE Matern random fields*

Description

`spde.matern.operators` is used for computing a rational SPDE approximation of a Gaussian random fields on R^d defined as a solution to the SPDE

$$(\kappa(s) - \Delta)^\beta (\tau(s)u(s)) = W.$$

Usage

```

spde.matern.operators(
  kappa = NULL,
  tau = NULL,
  theta = NULL,
  B.tau = matrix(c(0, 1, 0), 1, 3),
  B.kappa = matrix(c(0, 0, 1), 1, 3),

```

```

B.sigma = matrix(c(0, 1, 0), 1, 3),
B.range = matrix(c(0, 0, 1), 1, 3),
alpha = NULL,
nu = NULL,
parameterization = c("spde", "matern"),
G = NULL,
C = NULL,
d = NULL,
graph = NULL,
mesh = NULL,
range_mesh = NULL,
loc_mesh = NULL,
m = 1,
type = c("covariance", "operator"),
type_rational_approximation = c("chebfun", "brasil", "chebfunLB")
)

```

Arguments

<code>kappa</code>	Vector with the, possibly spatially varying, range parameter evaluated at the locations of the mesh used for the finite element discretization of the SPDE.
<code>tau</code>	Vector with the, possibly spatially varying, precision parameter evaluated at the locations of the mesh used for the finite element discretization of the SPDE.
<code>theta</code>	Theta parameter that connects <code>B.tau</code> and <code>B.kappa</code> to <code>tau</code> and <code>kappa</code> through a log-linear regression, in case the parameterization is <code>spde</code> , and that connects <code>B.sigma</code> and <code>B.range</code> to <code>tau</code> and <code>kappa</code> in case the parameterization is <code>matern</code> .
<code>B.tau</code>	Matrix with specification of log-linear model for τ . Will be used if <code>parameterization = 'spde'</code> .
<code>B.kappa</code>	Matrix with specification of log-linear model for κ . Will be used if <code>parameterization = 'spde'</code> .
<code>B.sigma</code>	Matrix with specification of log-linear model for σ . Will be used if <code>parameterization = 'matern'</code> .
<code>B.range</code>	Matrix with specification of log-linear model for ρ , which is a range-like parameter (it is exactly the range parameter in the stationary case). Will be used if <code>parameterization = 'matern'</code> .
<code>alpha</code>	smoothness parameter. Will be used if the parameterization is <code>'spde'</code> .
<code>nu</code>	Shape parameter of the covariance function. Will be used if the parameterization is <code>'matern'</code> .
<code>parameterization</code>	Which parameterization to use? <code>matern</code> uses <code>range</code> , <code>std.</code> deviation and <code>nu</code> (smoothness). <code>spde</code> uses <code>kappa</code> , <code>tau</code> and <code>nu</code> (smoothness). The default is <code>matern</code> .
<code>G</code>	The stiffness matrix of a finite element discretization of the domain of interest.
<code>C</code>	The mass matrix of a finite element discretization of the domain of interest.
<code>d</code>	The dimension of the domain. Does not need to be given if <code>mesh</code> is used.
<code>graph</code>	An optional <code>metric_graph</code> object. Replaces <code>d</code> , <code>C</code> and <code>G</code> .

<code>mesh</code>	An optional inla mesh. <code>d</code> , <code>C</code> and <code>G</code> must be given if <code>mesh</code> is not given.
<code>range_mesh</code>	The range of the mesh. Will be used to provide starting values for the parameters. Will be used if <code>mesh</code> and <code>graph</code> are <code>NULL</code> , and if one of the parameters (<code>kappa</code> or <code>tau</code> for <code>spde</code> parameterization, or <code>sigma</code> or <code>range</code> for <code>matern</code> parameterization) are not provided.
<code>loc_mesh</code>	The mesh locations used to construct the matrices <code>C</code> and <code>G</code> . This option should be provided if one wants to use the <code>rspde_lme()</code> function and will not provide neither <code>graph</code> nor <code>mesh</code> . Only works for 1d data. Does not work for metric graphs. For metric graphs you should supply the graph using the <code>graph</code> argument.
<code>m</code>	The order of the rational approximation, which needs to be a positive integer. The default value is 1.
<code>type</code>	The type of the rational approximation. The options are "covariance" and "operator". The default is "covariance".
<code>type_rational_approximation</code>	Which type of rational approximation should be used? The current types are "chebfun", "brasil" or "chebfunLB".

Details

The approximation is based on a rational approximation of the fractional operator $(\kappa(s)^2 - \Delta)^\beta$, where $\beta = (\nu + d/2)/2$. This results in an approximate model on the form

$$P_l u(s) = P_r W,$$

where $P_j = p_j(L)$ are non-fractional operators defined in terms of polynomials p_j for $j = l, r$. The order of p_r is given by `m` and the order of p_l is $m + m_\beta$ where m_β is the integer part of β if $\beta > 1$ and $m_\beta = 1$ otherwise.

The discrete approximation can be written as $u = P_r x$ where $x \sim N(0, Q^{-1})$ and $Q = P_l^T C^{-1} P_l$. Note that the matrices P_r and Q may be ill-conditioned for $m > 1$. In this case, the methods in `operator.operations()` should be used for operations involving the matrices, since these methods are more numerically stable.

Value

`spde.matern.operators` returns an object of class "rSPDEobj". This object contains the quantities listed in the output of `fractional.operators()` as well as the smoothness parameter ν .

See Also

`fractional.operators()`, `spde.matern.operators()`, `matern.operators()`

Examples

```
# Sample non-stationary Matern field on R
tau <- 1
nu <- 0.8

# create mass and stiffness matrices for a FEM discretization
```

```

x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# define a non-stationary range parameter
kappa <- seq(from = 2, to = 20, length.out = length(x))
alpha <- nu + 1/2
# compute rational approximation
op <- spde.matern.operators(
  kappa = kappa, tau = tau, alpha = alpha,
  G = fem$G, C = fem$C, d = 1
)

# sample the field
u <- simulate(op)

# plot the sample
plot(x, u, type = "l", ylab = "u(s)", xlab = "s")

```

summary.CBrSPDEobj *Summarise CBrSPDE objects*

Description

Summary method for class "CBrSPDEobj"

Usage

```

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

## S3 method for class 'summary.CBrSPDEobj'
print(x, ...)

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

```

Arguments

- object an object of class "CBrSPDEobj", usually, a result of a call to [matern.operators\(\)](#).
- ... further arguments passed to or from other methods.
- x an object of class "summary.CBrSPDEobj", usually, a result of a call to [summary.CBrSPDEobj\(\)](#).

Examples

```

# Compute the covariance-based rational approximation of a
# Gaussian process with a Matern covariance function on R
kappa <- 10

```

```

sigma <- 1
nu <- 0.8
range <- sqrt(8*nu)/kappa

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation of covariance function at 0.5
tau <- sqrt(gamma(nu) / (sigma^2 * kappa^(2 * nu) *
(4 * pi)^(1 / 2) * gamma(nu + 1 / 2)))
op_cov <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2,
  parameterization = "matern"
)
op_cov

```

summary.rSPDEobj *Summarise rSPDE objects*

Description

Summary method for class "rSPDEobj"

Usage

```

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

## S3 method for class 'summary.rSPDEobj'
print(x, ...)

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

```

Arguments

- | | |
|--------|--|
| object | an object of class "rSPDEobj", usually, a result of a call to fractional.operators() , matern.operators() , or spde.matern.operators() . |
| ... | further arguments passed to or from other methods. |
| x | an object of class "summary.rSPDEobj", usually, a result of a call to summary.rSPDEobj() . |

`summary.rspde_lme` *Summary Method for rspde_lme Objects.*

Description

Function providing a summary of results related to mixed effects regression models with Whittle-Matern latent models.

Usage

```
## S3 method for class 'rspde_lme'
summary(object, all_times = FALSE, ...)
```

Arguments

<code>object</code>	an object of class "rspde_lme" containing results from the fitted model.
<code>all_times</code>	Show all computed times.
<code>...</code>	not used.

Value

An object of class `summary_rspde_lme` containing several informations of a `rspde_lme` object.

`summary.rspde_result` *Summary for posteriors of field parameters for an inla_rspde model from a rspde_result object*

Description

Summary for posteriors of rSPDE field parameters in their original scales.

Usage

```
## S3 method for class 'rspde_result'
summary(object, digits = 6, ...)
```

Arguments

<code>object</code>	A <code>rspde_result</code> object.
<code>digits</code>	integer, used for number formatting with <code>signif()</code>
<code>...</code>	Currently not used.

Value

Returns a `data.frame` containing the summary.

Examples

```

#tryCatch version
tryCatch({
  if (requireNamespace("INLA", quietly = TRUE)){
    library(INLA)

    set.seed(123)

    m <- 100
    loc_2d_mesh <- matrix(runif(m * 2), m, 2)
    mesh_2d <- inla.mesh.2d(
      loc = loc_2d_mesh,
      cutoff = 0.05,
      max.edge = c(0.1, 0.5)
    )
    sigma <- 1
    range <- 0.2
    nu <- 0.8
    kappa <- sqrt(8 * nu) / range
    op <- matern.operators(
      mesh = mesh_2d, nu = nu,
      range = range, sigma = sigma, m = 2,
      parameterization = "matern"
    )
    u <- simulate(op)
    A <- inla.spde.make.A(
      mesh = mesh_2d,
      loc = loc_2d_mesh
    )
    sigma.e <- 0.1
    y <- A %*% u + rnorm(m) * sigma.e
    Abar <- rspde.make.A(mesh = mesh_2d, loc = loc_2d_mesh)
    mesh.index <- rspde.make.index(name = "field", mesh = mesh_2d)
    st.dat <- inla.stack(
      data = list(y = as.vector(y)),
      A = Abar,
      effects = mesh.index
    )
    rspde_model <- rspde.matern(
      mesh = mesh_2d,
      nu.upper.bound = 2
    )
    f <- y ~ -1 + f(field, model = rspde_model)
    rspde_fit <- inla(f,
      data = inla.stack.data(st.dat),
      family = "gaussian",
      control.predictor =
        list(A = inla.stack.A(st.dat))
    )
    result <- rspde.result(rspde_fit, "field", rspde_model)
    summary(result)
  }
}

```

```
#stable.tryCatch
}, error = function(e){print("Could not run the example")})
```

update.CBrSPDEobj *Update parameters of CBrSPDEobj objects*

Description

Function to change the parameters of a CBrSPDEobj object

Usage

```
## S3 method for class 'CBrSPDEobj'
update(
  object,
  user_nu = NULL,
  user_alpha = NULL,
  user_kappa = NULL,
  user_tau = NULL,
  user_sigma = NULL,
  user_range = NULL,
  user_theta = NULL,
  user_m = NULL,
  mesh = NULL,
  loc_mesh = NULL,
  graph = NULL,
  range_mesh = NULL,
  compute_higher_order = object$higher_order,
  parameterization = NULL,
  type_rational_approximation = object$type_rational_approximation,
  return_block_list = object$return_block_list,
  ...
)
```

Arguments

<code>object</code>	The covariance-based rational SPDE approximation, computed using matern.operators()
<code>user_nu</code>	If non-null, update the shape parameter of the covariance function. Will be used if parameterization is 'matern'.
<code>user_alpha</code>	If non-null, update the fractional SPDE order parameter. Will be used if parameterization is 'spde'.
<code>user_kappa</code>	If non-null, update the parameter kappa of the SPDE. Will be used if parameterization is 'spde'.
<code>user_tau</code>	If non-null, update the parameter tau of the SPDE. Will be used if parameterization is 'spde'.

user_sigma	If non-null, update the standard deviation of the covariance function. Will be used if parameterization is 'matern'.
user_range	If non-null, update the range parameter of the covariance function. Will be used if parameterization is 'matern'.
user_theta	For non-stationary models. If non-null, update the vector of parameters.
user_m	If non-null, update the order of the rational approximation, which needs to be a positive integer.
mesh	An optional inla mesh. Replaces d, C and G.
loc_mesh	The mesh locations used to construct the matrices C and G. This option should be provided if one wants to use the <code>rspde_lme()</code> function and will not provide neither graph nor mesh. Only works for 1d data. Does not work for metric graphs. For metric graphs you should supply the graph using the graph argument.
graph	An optional <code>metric_graph</code> object. Replaces d, C and G.
range_mesh	The range of the mesh. Will be used to provide starting values for the parameters. Will be used if mesh and graph are NULL, and if one of the parameters (kappa or tau for spde parameterization, or sigma or range for matern parameterization) are not provided.
compute_higher_order	Logical. Should the higher order finite element matrices be computed?
parameterization	If non-null, update the parameterization. Only works for stationary models.
type_rational_approximation	Which type of rational approximation should be used? The current types are "chebfun", "brasil" or "chebfunLB".
return_block_list	Logical. For type = "covariance", should the block parts of the precision matrix be returned separately as a list?
...	Currently not used.

Value

It returns an object of class "CBrSPDEobj". This object contains the same quantities listed in the output of `matern.operators()`.

See Also

`simulate.CBrSPDEobj()`, `matern.operators()`

Examples

```
# Compute the covariance-based rational approximation of a
# Gaussian process with a Matern covariance function on R
kappa <- 10
sigma <- 1
nu <- 0.8
```

```

range <- sqrt(8*nu)/kappa

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation of covariance function at 0.5
op_cov <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2,
  parameterization = "matern"
)
op_cov

# Update the range parameter of the model:
op_cov <- update(op_cov, user_kappa = 20)
op_cov

```

update.rSPDEobj*Update parameters of rSPDEobj objects***Description**

Function to change the parameters of a rSPDEobj object

Usage

```

## S3 method for class 'rSPDEobj'
update(
  object,
  user_nu = NULL,
  user_alpha = NULL,
  user_kappa = NULL,
  user_sigma = NULL,
  user_range = NULL,
  user_tau = NULL,
  user_theta = NULL,
  user_m = NULL,
  mesh = NULL,
  loc_mesh = NULL,
  graph = NULL,
  range_mesh = NULL,
  parameterization = NULL,
  ...
)

```

Arguments

object	The operator-based rational SPDE approximation, computed using matern.operators() with type="operator"
user_nu	If non-null, update the shape parameter of the covariance function.
user_alpha	If non-null, update the fractional order.
user_kappa	If non-null, update the range parameter of the covariance function.
user_sigma	If non-null, update the standard deviation of the covariance function.
user_range	If non-null, update the range parameter of the covariance function.
user_tau	If non-null, update the parameter tau.
user_theta	If non-null, update the parameter theta, that connects tau and kappa to the model matrices.
user_m	If non-null, update the order of the rational approximation, which needs to be a positive integer.
mesh	An optional inla mesh. Replaces d, C and G.
loc_mesh	The mesh locations used to construct the matrices C and G. This option should be provided if one wants to use the rspde_lme() function and will not provide neither graph nor mesh. Only works for 1d data. Does not work for metric graphs. For metric graphs you should supply the graph using the graph argument.
graph	An optional metric_graph object. Replaces d, C and G.
range_mesh	The range of the mesh. Will be used to provide starting values for the parameters. Will be used if mesh and graph are NULL, and if one of the parameters (kappa or tau for spde parameterization, or sigma or range for matern parameterization) are not provided.
parameterization	If non-null, update the parameterization. Only works for stationary models.
...	Currently not used.

Value

It returns an object of class "rSPDEobj". This object contains the same quantities listed in the output of [matern.operators\(\)](#).

See Also

[simulate.rSPDEobj\(\)](#), [matern.operators\(\)](#)

Examples

```
# Compute the operator-based rational approximation of a
# Gaussian process with a Matern covariance function on R
kappa <- 10
sigma <- 1
nu <- 0.8
range <- sqrt(8*nu)/kappa
```

```

# create mass and stiffness matrices for a FEM discretization
x <- seq(from = 0, to = 1, length.out = 101)
fem <- rSPDE.fem1d(x)

# compute rational approximation of covariance function at 0.5
op <- matern.operators(
  loc_mesh = x, nu = nu,
  range = range, sigma = sigma, d = 1, m = 2, type = "operator",
  parameterization = "matern"
)
op

# Update the range parameter of the model:
op <- update(op, user_kappa = 20)
op

```

variogram.intrinsic.spde*Variogram of intrinsic SPDE model***Description**

Variogram $\gamma(s_0, s)$ of intrinsic SPDE model

$$(-\Delta)^{\beta/2}(\kappa^2 - \Delta)^{\alpha/2}(\tau u) = \mathcal{W}$$

with Neumann boundary conditions and a mean-zero constraint on a square $[0, L]^d$ for $d = 1$ or $d = 2$.

Usage

```

variogram.intrinsic.spde(
  s0 = NULL,
  s = NULL,
  kappa = NULL,
  alpha = NULL,
  beta = NULL,
  tau = 1,
  L = NULL,
  N = 100,
  d = NULL
)

```

Arguments

<code>s0</code>	The location where the variogram should be evaluated, either a double for 1d or a vector for 2d
-----------------	---

s	A vector (in 1d) or matrix (in 2d) with all locations where the variogram is computed
kappa	Range parameter.
alpha	Smoothness parameter.
beta	Smoothness parameter.
tau	Precision parameter.
L	The side length of the square domain.
N	The number of terms in the Karhunen-Loeve expansion.
d	The dimension (1 or 2).

Details

The variogram is computed based on a Karhunen-Loeve expansion of the covariance function.

See Also

[intrinsic.matern.operators\(\)](#)

Examples

```
if (requireNamespace("RSpectra", quietly = TRUE)){
  x <- seq(from = 0, to = 10, length.out = 201)
  beta <- 1
  alpha <- 1
  kappa <- 1
  op <- intrinsic.matern.operators(kappa = kappa, tau = 1, alpha = alpha,
                                    beta = beta, loc_mesh = x, d=1)
  # Compute and plot the variogram of the model
  Sigma <- op$A %*% solve(op$Q,t(op$A))
  One <- rep(1, times = ncol(Sigma))
  D <- diag(Sigma)
  Gamma <- 0.5*(One %*% t(D) + D %*% t(One) - 2 * Sigma)
  k <- 100
  plot(x, Gamma[k, ], type = "l")
  lines(x,
        variogram.intrinsic.spde(x[k], x, kappa, alpha, beta, L = 10, d = 1),
        col=2, lty = 2)
}
```

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