

Package: pGPx (via r-universe)

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Type Package

Title Pseudo-Realizations for Gaussian Process Excursions

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<https://cs.brown.edu/people/pfelzens/dt/index.html>)

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Description Computes pseudo-realizations from the posterior distribution of a Gaussian Process (GP) with the method described in Azzimonti et al. (2016) <[doi:10.1137/141000749](https://doi.org/10.1137/141000749)>.

The realizations are obtained from simulations of the field at few well chosen points that minimize the expected distance in measure between the true excursion set of the field and the approximate one. Also implements a R interface for (the main function of) Distance Transform of sampled Functions (<<https://cs.brown.edu/people/pfelzens/dt/index.html>>).

URL <https://doi.org/10.1137/141000749>

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Imports Rcpp (>= 0.12.13), DiceKriging, pbivnorm, KrigInv, rgenoud, randtoolbox, pracma, grDevices

Suggests anMC, DiceDesign

LinkingTo Rcpp, RcppArmadillo

RoxygenNote 7.2.3

NeedsCompilation yes

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computeVolumes	<i>Compute Excursion Volume Distribution</i>
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Description

Compute the volume of excursion for each realization, includes a bias.correction for the mean. If the input is the actual GP values, compute also the random sets.

Usage

```
computeVolumes(
  rand.set,
  threshold,
  nsim,
  n.int.points,
  bias.corr = F,
  model = NULL,
  bias.corr.points = NULL
)
```

Arguments

rand.set	a matrix of size n.int.points \times nsim containing the excursion set realizations stored as long vectors. For example the excursion set obtained from the result of simulate_and_interpolate .
threshold	threshold value
nsim	number of simulations of the excursion set
n.int.points	total length of the excursion set discretization. The size of the image is sqrt(n.int.points).

`bias.corr` a boolean, if TRUE it computes the bias correction for the volume distribution.

`model` the `km` model for computing the bias correction. If NULL the bias correction is not computed.

`bias.corr.points` a matrix with d columns with the input points where to compute the bias correction. If NULL it is initialized as the first `n.int.points` of the Sobol' sequence.

Value

A vector of size `nsim` containing the excursion volumes for each realization.

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

Examples

```
### Simulate and interpolate for a 2d example
if (!requireNamespace("DiceKriging", quietly = TRUE)) {
  stop("DiceKriging needed for this example to work. Please install it.",
       call. = FALSE)
}
if (!requireNamespace("DiceDesign", quietly = TRUE)) {
  stop("DiceDesign needed for this example to work. Please install it.",
       call. = FALSE)
}
# Define the function
g=function(x){
  return(-DiceKriging::branin(x))
}
d=2
# Fit OK km model
design<-DiceDesign::maximinESE_LHS(design = DiceDesign::lhsDesign(n=50,
                                                                dimension = 2,
                                                                seed=42)$design)$design

colnames(design)<-c("x1", "x2")
observations<-apply(X = design, MARGIN = 1, FUN = g)
kmModel<-DiceKriging::km(formula = ~1, design = design, response = observations,
                        covtype = "matern3_2", control=list(trace=FALSE))

# Get simulation points
# Here they are not optimized, you can use optim_dist_measure to find optimized points
simu_points <- DiceDesign::maximinSA_LHS(DiceDesign::lhsDesign(n=100,
                                                                dimension = d,
                                                                seed=1)$design)$design

# obtain nsims posterior realization at simu_points
nsims <- 30
nn_data<-expand.grid(seq(0,1,,50),seq(0,1,,50))
```

```

nn_data<-data.frame(nn_data)
colnames(nn_data)<-colnames(kmModel@X)
approx.simu <- simulate_and_interpolate(object=kmModel, nsim = nsims, simupoints = simu_points,
                                       interpolatepoints = as.matrix(nn_data),
                                       nugget.sim = 0, type = "UK")
exVol<- computeVolumes(rand.set = approx.simu,threshold = -10,
                       nsim = nsims,n.int.points = 50^2,bias.corr=TRUE, model=kmModel)

hist(exVol, main="Excursion Volume")

```

compute_contourLength *Compute contour lengths*

Description

Computes the contour lengths for the excursion sets in gpRealizations

Usage

```
compute_contourLength(gpRealizations, threshold, nRealizations, verb = 1)
```

Arguments

gpRealizations	a matrix of size nRealizationsximageSize^2 containing the GP realizations stored as long vectors. For example the object returned by simulate_and_interpolate .
threshold	threshold value
nRealizations	number of simulations of the excursion set
verb	an integer to choose the level of verbosity

Value

A vector of size nRealizations containing the countour lines lenghts.

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

Examples

```

### Simulate and interpolate for a 2d example
if (!requireNamespace("DiceKriging", quietly = TRUE)) {
  stop("DiceKriging needed for this example to work. Please install it.",
       call. = FALSE)
}
if (!requireNamespace("DiceDesign", quietly = TRUE)) {
  stop("DiceDesign needed for this example to work. Please install it.",
       call. = FALSE)
}
# Define the function
g=function(x){
  return(-DiceKriging::branin(x))
}
d=2
# Fit OK km model
design<-DiceDesign::maximinESE_LHS(design = DiceDesign::lhsDesign(n=50,
                                                                dimension = 2,
                                                                seed=42)$design)$design

colnames(design)<-c("x1", "x2")
observations<-apply(X = design, MARGIN = 1, FUN = g)
kmModel<-DiceKriging::km(formula = ~1, design = design, response = observations,
                        covtype = "matern3_2", control=list(trace=FALSE))

# Get simulation points
# Here they are not optimized, you can use optim_dist_measure to find optimized points
simu_points <- DiceDesign::maximinSA_LHS(DiceDesign::lhsDesign(n=100,
                                                            dimension = d,
                                                            seed=1)$design)$design

# obtain nsims posterior realization at simu_points
nsims <- 1
nn_data<-expand.grid(seq(0,1,,50),seq(0,1,,50))
nn_data<-data.frame(nn_data)
colnames(nn_data)<-colnames(kmModel@X)
approx.simu <- simulate_and_interpolate(object=kmModel, nsim = nsims, simupoints = simu_points,
                                       interpolatepoints = as.matrix(nn_data),
                                       nugget.sim = 0, type = "UK")
cLLs<- compute_contourLength(gpRealizations = approx.simu, threshold = -10,
                             nRealizations = nsims, verb = 1)

```

dtf_fast

*Rcpp implementation of Felzenszwalb distance transform***Description**

Rcpp wrapper for the distance transform algorithm described in Felzenszwalb and Huttenlocher (2012)

Usage

```
dt_fast(x)
```

Arguments

`x` matrix of booleans of size $n \times m$ representing a (binary) image

Value

A matrix of size $n \times m$ containing the distance transform result. Note that this function does not perform any checks on `x`.

Author(s)

Pedro Felzenszwalb for the header files `dt.h` and `misc.h` that do the work, Dario Azzimonti and Julien Bect for the wrapper.

References

Felzenszwalb, P. F. and Huttenlocher, D. P. (2012). Distance Transforms of Sampled Functions. *Theory of Computing*, 8(19):415-428.

Examples

```
# Create an image with a square
nc = 256
nr = 256
xx = matrix(FALSE, ncol=nc, nrow=nr)
xx[(nr/16):(nr/16*15-1), nc/16] <- rep(TRUE, nr/16*14)
xx[(nr/16):(nr/16*15-1), nc/16*15] <- rep(TRUE, nr/16*14)
xx[nr/16, (nc/16):(nc/16*15-1)] <- rep(TRUE, nc/16*14)
xx[nr/16*15, (nc/16):(nc/16*15-1)] <- rep(TRUE, nc/16*14)
# Compute Distance transform
zz <- dt_fast(xx)

# Plot the results
image(xx, col=grey.colors(20), main="Original image")
image(zz, col=grey.colors(20), main="Distance transform")
```

Description

Compute the expected L^2 distance between the average distance transform and the set realizations. If the input is the actual values of the gaussian process, compute also the random sets.

Usage

```
DTV(rand.set, threshold, nsim, n.int.points)
```

Arguments

rand.set	a matrix of size <code>n.int.points</code> x <code>nsim</code> containing the excursion set realizations stored as long vectors. For example the excursion set obtained from the result of <code>simulate_and_interpolate</code> .
threshold	threshold value
nsim	number of simulations of the excursion set
n.int.points	total length of the excursion set discretization. The size of the image is <code>sqrt(n.int.points)</code> .

Value

A list containing

- `variance`: Value of the distance transform variability. The integral of `dvar` over the spatial domain.
- `dbar`: empirical distance average transform $1/N \sum_{i=1}^N d(x, \Gamma_i)$, a matrix of size `n.int.points` x `n.int.points`
- `dvar`: empirical variance of distance transform $1/N \sum_{i=1}^N (d(x, \Gamma_i) - dbar)^2$, a matrix of size `n.int.points` x `n.int.points`
- `alldt`: distance transforms for all realizations, a matrix of size `n.int.points` x `nsim`
- `naTot`: Total number of infinite distance transform values. These are returned in realizations where there is no excursion.

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

Felzenszwalb, P. F. and Huttenlocher, D. P. (2012). Distance Transforms of Sampled Functions. *Theory of Computing*, 8(19):415-428.

Examples

```
### Simulate and interpolate for a 2d example
if (!requireNamespace("DiceKriging", quietly = TRUE)) {
  stop("DiceKriging needed for this example to work. Please install it.",
       call. = FALSE)
}
if (!requireNamespace("DiceDesign", quietly = TRUE)) {
  stop("DiceDesign needed for this example to work. Please install it.",
       call. = FALSE)
}
```

```

# Define the function
g=function(x){
  return(-DiceKriging::branin(x))
}
d=2
# Fit OK km model
design<-DiceDesign::maximinESE_LHS(design = DiceDesign::lhsDesign(n=50,
                                                                    dimension = 2,
                                                                    seed=42)$design)$design

colnames(design)<-c("x1", "x2")
observations<-apply(X = design, MARGIN = 1, FUN = g)
kmModel<-DiceKriging::km(formula = ~1, design = design, response = observations,
                          covtype = "matern3_2", control=list(trace=FALSE))

# Get simulation points
# Here they are not optimized, you can use optim_dist_measure to find optimized points
simu_points <- DiceDesign::maximinSA_LHS(DiceDesign::lhsDesign(n=100,
                                                                dimension = d,
                                                                seed=1)$design)$design

# obtain nsims posterior realization at simu_points
nsims <- 30
nn_data<-expand.grid(seq(0,1,,50),seq(0,1,,50))
nn_data<-data.frame(nn_data)
colnames(nn_data)<-colnames(kmModel@X)
approx.simu <- simulate_and_interpolate(object=kmModel, nsim = nsims, simupoints = simu_points,
                                       interpolatepoints = as.matrix(nn_data),
                                       nugget.sim = 0, type = "UK")
Dvar<- DTV(rand.set = approx.simu, threshold = -10,
           nsim = nsims, n.int.points = 50^2)

image(matrix(Dvar$dvar, ncol=50), col=grey.colors(20), main="average distance transform")
image(matrix(Dvar$dvar, ncol=50), col=grey.colors(20), main="variance of distance transform")
points(design, pch=17)

```

edm_crit

Distance in measure criterion

Description

Computes the distance in measure criterion.

Usage

```

edm_crit(
  x,
  integration.points,
  integration.weights = NULL,
  intpoints.oldmean,
  intpoints.oldsd,

```



```

    precalc.data,
    model,
    threshold,
    batchsize,
    alpha,
    current.crit
)

```

Arguments

`x` vector of dimension d representing the ith point where to compute the criterion

`integration.points` $p \times d$ matrix of points for numerical integration in the X space.

`integration.weights` Vector of size p corresponding to the weights of these integration points.

`intpoints.oldmean` Vector of size p corresponding to the kriging mean at the integration points.

`intpoints.oldsd` Vector of size p corresponding to the kriging standard deviation at the integration points.

`precalc.data` list result of [precomputeUpdateData](#) with `model` and `x`.

`model` km model

`threshold` threshold selected for excursion set

`batchsize` number of simulation points

`alpha` value of Vorob'ev threshold

`current.crit` Current value of the criterion

Value

the value of the expected distance in measure criterion at x

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

edm_crit2

Distance in measure criterion

Description

Computes the distance in measure criterion. To be used in optimization routines.

Usage

```
edm_crit2(
  x,
  other.points,
  integration.points,
  integration.weights = NULL,
  intpoints.oldmean,
  intpoints.oldsd,
  precalc.data,
  model,
  threshold,
  batchsize,
  alpha,
  current.crit
)
```

Arguments

x	vector of dimension d representing the point where to compute the criterion
other.points	Vector giving the other batchsize-1 points at which one wants to evaluate the criterion
integration.points	$p \times d$ matrix of points for numerical integration in the X space.
integration.weights	Vector of size p corresponding to the weights of these integration points.
intpoints.oldmean	Vector of size p corresponding to the kriging mean at the integration points.
intpoints.oldsd	Vector of size p corresponding to the kriging standard deviation at the integration points.
precalc.data	list result of precomputeUpdateData with model and x.
model	km model
threshold	threshold selected for excursion set
batchsize	number of simulation points
alpha	value of Vorob'ev threshold
current.crit	Current value of the criterion

Value

the value of the expected distance in measure criterion at x , other . points.

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

expDistMeasure	<i>Compute expected distance in measure of approximate excursion set</i>
----------------	--

Description

Computes expected distance in measure between the excursion set of the approximated process and the true excursion set.

Usage

```
expDistMeasure(
  simupoints,
  model,
  threshold,
  batchsize,
  integration.param = NULL
)
```

Arguments

simupoints	a numeric array of size batchsize*d containing the simulation points.
model	a km model
threshold	threshold value
batchsize	number of simulations points
integration.param	a list containing parameters for the integration of the criterion A, see max_sur_parallel for more details.

Value

A positive value indicating the expected distance in measure.

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

Examples

```
### Compute optimal simulation points in a 2d example
if (!requireNamespace("DiceKriging", quietly = TRUE)) {
  stop("DiceKriging needed for this example to work. Please install it.",
       call. = FALSE)
}
if (!requireNamespace("DiceDesign", quietly = TRUE)) {
  stop("DiceDesign needed for this example to work. Please install it.",
       call. = FALSE)
}
# Define the function
g=function(x){
  return(-DiceKriging::branin(x))
}
d=2
# Fit OK km model
design<-DiceDesign::maximinESE_LHS(design = DiceDesign::lhsDesign(n=20,
                                                                dimension = 2,
                                                                seed=42)$design)$design

colnames(design)<-c("x1", "x2")
observations<-apply(X = design, MARGIN = 1, FUN = g)
kmModel<-DiceKriging::km(formula = ~1, design = design, response = observations,
                        covtype = "matern3_2", control=list(trace=FALSE))

threshold <- -10

# Obtain simulation point sampling from maximin LHS design
batchsize <- 50
set.seed(1)
mmLHS_simu_points <- DiceDesign::maximinSA_LHS(DiceDesign::lhsDesign(n=batchsize,
                                                                    dimension = d,
                                                                    seed=1)$design)$design

# Compute expected distance in measure for approximation obtain from random simulation points
EDM_mmLHS <- rep(NA, batchsize)
integcontrol <- list(distrib="sobol", n.points=1000)
integration.param <- KrigInv::integration_design(integcontrol, d=d,
                                                lower=c(0, 0), upper=c(1, 1),
                                                model=kmModel, T=threshold)

integration.param$alpha <- 0.5
for(i in seq(1, batchsize)){
```

```

EDM_mmLHS[i]<-expDistMeasure( mmLHS_simu_points[1:i,],model = kmModel,
                             threshold = threshold,batchsize = i,
                             integration.param = integration.param )
}

plot(EDM_mmLHS,type='l',main="Expected distance in measure",xlab="batchsize")

## Not run:
# Get optimized simulation points with algorithm B
simu_points <- optim_dist_measure(model=kmModel,threshold = threshold,
                                 lower = c(0,0),upper = c(1,1),
                                 batchsize = batchsize,algorithm = "B")

# plot the criterion value
plot(1:batchsize,simu_points$value,type='l',main="Criterion value")

# Compute expected distance in measure for approximation obtained from optimized simulation points
EDM_optB <- rep(NA,batchsize)
for(i in seq(1,batchsize)){
  EDM_optB[i]<-expDistMeasure( simu_points$par[1:i,],model = kmModel,threshold = threshold,
                              batchsize = i,integration.param = integration.param )
}
plot(EDM_mmLHS,type='l',main="Expected distance in measure",
     xlab="batchsize",ylab="EDM",
     ylim=range(EDM_mmLHS,EDM_optB))
lines(EDM_optB,col=2,lty=2)
legend("topright",c("Maximin LHS", "B"),lty=c(1,2),col=c(1,2))

# Get optimized simulation points with algorithm A
simu_pointsA <- optim_dist_measure(model=kmModel,threshold = threshold,
                                   lower = c(0,0),upper = c(1,1),
                                   batchsize = batchsize,algorithm = "A")

# plot the criterion value
plot(1:batchsize,simu_pointsA$value,type='l',main="Criterion value")

# Compute expected distance in measure for approximation obtained from optimized simulation points
EDM_optA <- rep(NA,batchsize)
for(i in seq(1,batchsize)){
  EDM_optA[i]<-expDistMeasure( simu_pointsA$par[1:i,],model = kmModel,threshold = threshold,
                              batchsize = i,integration.param = integration.param )
}
plot(EDM_mmLHS,type='l',main="Expected distance in measure",
     xlab="batchsize",ylab="EDM",
     ylim=range(EDM_mmLHS,EDM_optB,EDM_optA))
lines(EDM_optB,col=2,lty=2)
lines(EDM_optA,col=3,lty=3)
legend("topright",c("Maximin LHS", "A", "B"),lty=c(1,3,2),col=c(1,3,2))

## End(Not run)

```

grad_kweights	<i>Gradient of the weights for interpolating simulations</i>
---------------	--

Description

Returns a list with the gradients of the posterior mean and the gradient of the (ordinary) kriging weights for simulations points.

Usage

```
grad_kweights(object, simu_points, krig_points, T.mat = NULL, F.mat = NULL)
```

Arguments

object	km object
simu_points	simulations points, locations where the field was simulated.
krig_points	one point where the interpolation is computed.
T.mat	a matrix (n+p)x(n+p) representing the Choleski factorization of the covariance matrix for the initial design and simulation points.
F.mat	a matrix (n+p)x(fdim) representing the evaluation of the model matrix at the initial design and simulation points.

Value

A list containing the gradients of posterior mean and kriging weights for simulation points.

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

Examples

```
#####
### Compute the weights and gradient on a 2d example
if (!requireNamespace("DiceKriging", quietly = TRUE)) {
  stop("DiceKriging needed for this example to work. Please install it.",
       call. = FALSE)
}
if (!requireNamespace("DiceDesign", quietly = TRUE)) {
  stop("DiceDesign needed for this example to work. Please install it.",
       call. = FALSE)
}
# Define the function
```

```

g=function(x){
  return(-DiceKriging::branin(x))
}
d=2
# Fit OK km model
design<-DiceDesign::maximinESE_LHS(design = DiceDesign::lhsDesign(n=50,
  dimension = 2,
  seed=42)$design)$design

colnames(design)<-c("x1", "x2")
observations<-apply(X = design,MARGIN = 1,FUN = g)
kmModel<-DiceKriging::km(formula = ~1,design = design,response = observations,
  covtype = "matern3_2",control=list(trace=FALSE))

# Get simulation points
# Here they are not optimized, you can use optim_dist_measure to find optimized points
set.seed(1)
simu_points <- matrix(runif(100*d),ncol=d)
# obtain nsims posterior realization at simu_points
nsims <- 1
set.seed(2)
some.simu <- DiceKriging::simulate(object=kmModel,nsim=nsims,newdata=simu_points,nugget.sim=1e-6,
  cond=TRUE,checkNames = FALSE)
nn_data<-expand.grid(seq(0,1,,50),seq(0,1,,50))
nn_data<-data.frame(nn_data)
colnames(nn_data)<-colnames(kmModel@X)
obj<-krig_weight_GPsimu(object = kmModel,simu_points = simu_points,krig_points = as.matrix(nn_data))

## Plot the approximate process realization and the gradient vector field
k_scale<-5e-4
image(matrix(obj$krig.mean.init+crossprod(obj$Lambda.end,some.simu[1,]),ncol=50),
  col=grey.colors(20))
contour(matrix(obj$krig.mean.init+crossprod(obj$Lambda.end,some.simu[1,]),ncol=50),
  nlevels = 20,add=TRUE)

for(c_ii in c(1,seq(10,2500,by = 64))){
  pp<-t(as.matrix(nn_data)[c_ii,])
  obj_deriv <- grad_kweights(object = kmModel,simu_points = simu_points,krig_points = pp)
  S_der<-obj_deriv$krig.mean.init + crossprod(obj_deriv$Lambda.end,some.simu[1,])
  points(x = pp[1],y = pp[2],pch=16)
  arrows(x0=pp[1],y0=pp[2],x1 = pp[1]+k_scale*S_der[1,1],y1=pp[2]+k_scale*S_der[2,1])
}

```

integrands_edm_crit *Integrand of the distance in measure criterion*

Description

Computes the integrand of the distance in measure criterion.

Usage

```
integrand_edm_crit(
  x,
  E,
  model,
  Thresh,
  batchsize,
  alpha,
  predE,
  predx = NULL,
  precalc.data = NULL
)
```

Arguments

<code>x</code>	vector of dimension d representing the i th point where to compute the criterion
<code>E</code>	matrix of dimension $d * (i - 1)$ containing the previously optimized simulation points
<code>model</code>	km model
<code>Thresh</code>	threshold selected for excursion set
<code>batchsize</code>	number of simulation points
<code>alpha</code>	value of Vorob'ev threshold
<code>predE</code>	list containing the posterior mean and standard deviation at E
<code>predx</code>	list containing the posterior mean and standard deviation at x
<code>precalc.data</code>	list result of precomputeUpdateData with model and x.

Value

the value of the integrand at x

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

krig_weight_GPsimu *Weights for interpolating simulations*

Description

Returns a list with the posterior mean and the kriging weights for simulations points.

Usage

```
krig_weight_GPsimu(
  object,
  simu_points,
  krig_points,
  T.mat = NULL,
  F.mat = NULL
)
```

Arguments

object	km object.
simu_points	simulations points, locations where the field was simulated.
krig_points	points where the interpolation is computed.
T.mat	a matrix (n+p)x(n+p) representing the Choleski factorization of the covariance matrix for the initial design and simulation points.
F.mat	a matrix (n+p)x(fdim) representing the evaluation of the model matrix at the initial design and simulation points.

Value

A list containing the posterior mean and the (ordinary) kriging weights for simulation points.

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

Examples

```
#####
### Compute the weights for approximating process on a 1d example
if (!requireNamespace("DiceKriging", quietly = TRUE)) {
  stop("DiceKriging needed for this example to work. Please install it.",
       call. = FALSE)
}
```

```

}
if (!requireNamespace("DiceDesign", quietly = TRUE)) {
  stop("DiceDesign needed for this example to work. Please install it.",
       call. = FALSE)
}
## Create kriging model from GP realization
design<-DiceDesign::maximinESE_LHS(design = DiceDesign::lhsDesign(n=20,
                                                                dimension = 1,
                                                                seed=42)$design)$design

colnames(design)<-c("x1")
gp0 <- DiceKriging::km(formula = ~1, design = design,
                      response = rep(x = 0, times = nrow(design)),
                      covtype = "matern3_2", coef.trend = 0,
                      coef.var = 1, coef.cov = 0.2)

set.seed(1)
observations <- t(DiceKriging::simulate(object = gp0, newdata = design, cond = FALSE))

# Fit OK km model
kmModel<-DiceKriging::km(formula = ~1,design = design,response = observations,
                        covtype = "matern3_2",control=list(trace=FALSE))

# Get simulation points
# Here they are not optimized, you can use optim_dist_measure to find optimized points
set.seed(2)
simu_points <- matrix(runif(20),ncol=1)
# obtain nsims posterior realization at simu_points
nsims <- 10
set.seed(3)
some.simu <- DiceKriging::simulate(object=kmModel,nsim=nsims,newdata=simu_points,nugget.sim=1e-6,
                                cond=TRUE,checkNames = FALSE)

grid<-seq(0,1,,100)
nn_data<-data.frame(grid)
colnames(nn_data)<-colnames(kmModel@X)
pred_nn<-DiceKriging::predict.km(object = kmModel,newdata = nn_data,type = "UK")
obj <- krig_weight_GPsimu(object=kmModel,simu_points=simu_points,krig_points=grid)

# Plot the posterior mean and some approximate process realizations
result <- matrix(nrow=nsims,ncol=length(grid))

plot(nn_data$x1,pred_nn$mean,type='l')
for(i in 1:nsims){
  some.simu.i <- matrix(some.simu[i,],ncol=1)
  result[i,] <- obj$krig.mean.init + crossprod(obj$Lambda.end,some.simu.i)
  points(simu_points,some.simu.i)
  lines(grid,result[i,],col=3)
}

```

Description

Optimizes the distance in measure criterion.

Usage

```
max_distance_measure(
  lower,
  upper,
  optimcontrol = NULL,
  batchsize,
  integration.param,
  T,
  model
)
```

Arguments

lower	a d dimensional vector containing the lower bounds for the optimization
upper	a d dimensional vector containing the upper bounds for the optimization
optimcontrol	the parameters for the optimization, see max_sur_parallel for more details.
batchsize	number of simulations points to find
integration.param	the parameters for the integration of the criterion, see max_sur_parallel for more details.
T	threshold value
model	a km model

Value

A list containing

- par a matrix $batchsize \times d$ containing the optimal points
- value if `optimcontrol$optim.option!=1` and `optimcontrol$method=="genoud"` (default options) a vector of length `batchsize` containing the optimum at each step otherwise the value of the criterion at the optimum.

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

max_integrand_edm *Maximize the integrand distance in measure criterion*

Description

Optimizes the integrand of the distance in measure criterion.

Usage

```
max_integrand_edm(
  lower,
  upper,
  batchsize,
  alpha = 0.5,
  Thresh,
  model,
  verb = 1
)
```

Arguments

lower	a d dimensional vector containing the lower bounds for the optimization
upper	a d dimensional vector containing the upper bounds for the optimization
batchsize	number of simulations points to find
alpha	value of Vorob'ev threshold
Thresh	threshold value
model	a km model
verb	an integer to choose the level of verbosity

Value

A list containing

- par a matrix $\text{batchsize} \times d$ containing the optimal points
- value a vector of length batchsize with the value of the criterion after each optimization
- fcount count of the number of criterion evaluations

References

Azzimonti D. F., Bect J., Chevalier C. and Ginsbourger D. (2016). Quantifying uncertainties on excursion sets under a Gaussian random field prior. *SIAM/ASA Journal on Uncertainty Quantification*, 4(1):850–874.

Azzimonti, D. (2016). Contributions to Bayesian set estimation relying on random field priors. PhD thesis, University of Bern.

optim_dist_measure *Choose simulation points*

Description

Selects batchsize locations where to simulate the field by minimizing the distance in measure criterion or by maximizing the integrand of the distance in measure criterion. Currently it is only a wrapper for the functions `max_distance_measure` and `max_integrand_edm`.

Usage

```
optim_dist_measure(
  model,
  threshold,
  lower,
  upper,
  batchsize,
  algorithm = "B",
  alpha = 0.5,
  verb = 1,
  optimcontrol = NULL,
  integration.param = NULL
)
```

Arguments

model	a km model
threshold	threshold value
lower	a d dimensional vector containing the lower bounds for the optimization
upper	a d dimensional vector containing the upper bounds for the optimization
batchsize	number of simulations points to find
algorithm	type of algorithm used to obtain simulation points: <ul style="list-style-type: none"> • "A" minimize the full integral criterion; • "B" maximize the integrand of the criterion.
alpha	value of Vorob'ev threshold
verb	an integer to choose the level of verbosity
optimcontrol	a list containing optional parameters for the optimization, see max_sur_parallel for more details.
integration.param	a list containing parameters for the integration of the criterion A, see max_sur_parallel for more details.


```

                                batchsize = batchsize,algorithm = "A")

# Get 75 simulation points with algorithm B
batchsize <- 75
simu_pointsB <- optim_dist_measure(model=kmModel,threshold = -10,
                                lower = c(0,0),upper = c(1,1),
                                batchsize = batchsize,algorithm = "B")

# plot the criterion value
critValA <-c(simu_pointsA$value,rep(NA,25))
par(mar = c(5,5,2,5))
plot(1:batchsize,critValA,type='l',main="Criterion value",ylab="Algorithm A",xlab="batchsize")
par(new=T)
plot(1:batchsize,simu_pointsB$value, axes=F, xlab=NA, ylab=NA,col=2,lty=2,type='l')
axis(side = 4)
mtext(side = 4, line = 3, 'Algorithm B')
legend("topright",c("Algorithm A","Algorithm B"),lty=c(1,2),col=c(1,2))
par(mar= c(5, 4, 4, 2) + 0.1)

# obtain nsims posterior realization at simu_points
nsims <- 1
nn_data<-expand.grid(seq(0,1,,50),seq(0,1,,50))
nn_data<-data.frame(nn_data)
colnames(nn_data)<-colnames(kmModel@X)
approx.simu <- simulate_and_interpolate(object=kmModel, nsim = 1, simupoints = simu_pointsA$par,
                                      interpolatepoints = as.matrix(nn_data),
                                      nugget.sim = 0, type = "UK")

## Plot the approximate process realization
image(matrix(approx.simu[1,],ncol=50),
      col=grey.colors(20))
contour(matrix(approx.simu[1,],ncol=50),
        nlevels = 20,add=TRUE)
points(simu_pointsA$par,pch=17)
points(simu_pointsB$par,pch=1,col=2)

## End(Not run)

```

```
simulate_and_interpolate
```

Simulate and interpolate

Description

Generates nsims approximate posterior field realizations at interpolatepoints. The approximate realizations are computed by simulating the field only at simupoints simulation points.

Usage

```
simulate_and_interpolate(
```



```
seed=42)$design)$design
colnames(design)<-c("x1","x2")
observations<-apply(X = design,MARGIN = 1,FUN = g)
kmModel<-DiceKriging::km(formula = ~1,design = design,response = observations,
  covtype = "matern3_2",control=list(trace=FALSE))
# Get simulation points
# Here they are not optimized, you can use optim_dist_measure to find optimized points
simu_points <- DiceDesign::maximinSA_LHS(DiceDesign::lhsDesign(n=100,
  dimension = d,
  seed=1)$design)$design

# obtain nsims posterior realization at simu_points
nsims <- 1
nn_data<-expand.grid(seq(0,1,,50),seq(0,1,,50))
nn_data<-data.frame(nn_data)
colnames(nn_data)<-colnames(kmModel@X)
approx.simu <- simulate_and_interpolate(object=kmModel, nsim = 1, simupoints = simu_points,
  interpolatepoints = as.matrix(nn_data),
  nugget.sim = 0, type = "UK")

## Plot the approximate process realization
image(matrix(approx.simu[1,],ncol=50),
  col=grey.colors(20))
contour(matrix(approx.simu[1,],ncol=50),
  nlevels = 20,add=TRUE)
```

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