Package 'pGPx'

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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>. 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>).

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Encoding UTF-8

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

Suggests anMC, DiceDesign

LinkingTo Rcpp, RcppArmadillo

RoxygenNote 7.2.3

NeedsCompilation yes

Repository CRAN

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Contents

`computeVolumes` *Compute Excursion Volume Distribution*

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.pointsxnsim` 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 lenghts*

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)
```

`dtt_fast` *Rcpp implementation of Felzenszwalb distance transfom*

Description

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

Usage

```
dtt_fast(x)
```

Arguments

x	matrix of booleans of size nxm representing a (binary) image

Value

A matrix of size nxm 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<- dtt_fast(xx)

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

DTV *Compute Distance Transform Variability*

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 n.int.pointsxnsim 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).

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$dbar,ncol=50),col=grey.colors(20),main="average distance transform")
image(matrix(Dvar$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*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*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` *Compute expected distance in measure of approximate excursion set*

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 *Gradient of the weights for interpolating simulations*

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])
}
```

`integrand_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

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

`max_distance_measure` *Minimize the distance in measure criterion*

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*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 `batchsize*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: • `"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.

Value

A list containing

- par a matrix batchsize*d containing the optimal points
- value a vector of length batchsize with the values of the criterion at each step

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))

# Run optim_dist_measure, algorithm B to obtain one simulation point
# NOTE: the approximating process resulting from 1 simulation point
# is very rough and it should not be used, see below for a more principled example.
simu_pointsB <- optim_dist_measure(model=kmModel,threshold = -10,
                                   lower = c(0,0),upper = c(1,1),
                                   batchsize = 1,algorithm = "B")

## Not run:
# Get 75 simulation points with algorithm A
batchsize <- 50
simu_pointsA <- optim_dist_measure(model=kmModel,threshold = -10,
                                   lower = c(0,0),upper = c(1,1),
```

```
                                    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(
```

```
  object,
  nsim = 1,
  simupoints = NULL,
  interpolatepoints = NULL,
  nugget.sim = 0,
  type = "UK"
)
```

Arguments

object	km object
nsim	numbero of simulations
simupoints	simulations points, locations where the field was simulated
interpolatepoints	points where posterior simulations are approximated
nugget.sim	nugget to be added to simulations for numerical stability
type	type of kriging model used for approximation (default Universal Kriging)

Value

A matrix `nsim*interpolatepoints` containing the approximate realizations.

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 = 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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