

Package ‘GPareto’

May 9, 2019

Type Package

Title Gaussian Processes for Pareto Front Estimation and Optimization

Version 1.1.3

Date 2019-05-09

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Description Gaussian process regression models, a.k.a. Kriging models, are applied to global multi-objective optimization of black-box functions. Multi-objective Expected Improvement and Step-wise Uncertainty Reduction sequential infill criteria are available. A quantification of uncertainty on Pareto fronts is provided using conditional simulations.

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Depends DiceKriging, emoa

Imports Rcpp (>= 0.12.15), methods, rgenoud, pbivnorm, pso, randtoolbox, KrigInv, MASS, DiceDesign, ks, rgl

Suggests knitr

VignetteBuilder knitr

LinkingTo Rcpp

Repository CRAN

URL <http://github.com/mbinois/GPareto>

BugReports <http://github.com/mbinois/GPareto/issues>

RoxygenNote 6.1.1

NeedsCompilation yes

Date/Publication 2019-05-09 20:30:09 UTC

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GPareto-package	<i>Package GPareto</i>
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Description

Multi-objective optimization and quantification of uncertainty on Pareto fronts, using Gaussian process models.

Details

Important functions:

[GParetooptim](#)
[easyGParetooptim](#)
[crit_optimizer](#)
[plotGPareto](#)
[CPF](#)

Note

Part of this work has been conducted within the frame of the ReDice Consortium, gathering industrial (CEA, EDF, IFPEN, IRSN, Renault) and academic (Ecole des Mines de Saint-Etienne, INRIA, and the University of Bern) partners around advanced methods for Computer Experiments. (<http://www.redice-project.org/>).

The authors would like to thank Yves Deville for his precious advices in R programming and packaging, as well as Olivier Roustant and David Ginsbourger for testing and suggestions of improvements for this package. We would also like to thank Tobias Wagner for providing his Matlab codes for the SMS-EGO strategy.

Author(s)

Mickael Binois, Victor Picheny

References

M. Binois, D. Ginsbourger and O. Roustant (2015), Quantifying Uncertainty on Pareto Fronts with Gaussian process conditional simulations, *European Journal of Operational Research*, 243(2), 386-394.

O. Roustant, D. Ginsbourger and Yves Deville (2012), DiceKriging, DiceOptim: Two R Packages for the Analysis of Computer Experiments by Kriging-Based Metamodeling and Optimization, *Journal of Statistical Software*, 51(1), 1-55, <http://www.jstatsoft.org/v51/i01/>.

M. T. Emmerich, A. H. Deutz, J. W. Klinkenberg (2011), Hypervolume-based expected improvement: Monotonicity properties and exact computation, *Evolutionary Computation (CEC)*, 2147-2154.

V. Picheny (2015), Multiobjective optimization using Gaussian process emulators via stepwise uncertainty reduction, *Statistics and Computing*, 25(6), 1265-1280.

T. Wagner, M. Emmerich, A. Deutz, W. Ponweiser (2010), On expected-improvement criteria for model-based multi-objective optimization, *Parallel Problem Solving from Nature*, 718-727, Springer, Berlin.

J. D. Svenson (2011), *Computer Experiments: Multiobjective Optimization and Sensitivity Analysis*, Ohio State University, PhD thesis.

C. Chevalier (2013), *Fast uncertainty reduction strategies relying on Gaussian process models*, University of Bern, PhD thesis.

M. Binois, V. Picheny (2019), GPareto: An R Package for Gaussian-Process-Based Multi-Objective Optimization and Analysis, *Journal of Statistical Software*, 89(8), 1–30.

See Also

[DiceKriging](#), [DiceOptim](#)

Examples

```
## Not run:
#-----
# Example 1 : Surrogate-based multi-objective Optimization with postprocessing
#-----
set.seed(25468)
```

```

d <- 2
fname <- P2

plotParetoGrid(P2) # For comparison

# Optimization
budget <- 25
lower <- rep(0, d)
upper <- rep(1, d)

omEGO <- easyGParetooptim(fn = fname, budget = budget, lower = lower, upper = upper)

# Postprocessing
plotGPareto(omEGO, add= FALSE, UQ_PF = TRUE, UQ_PS = TRUE, UQ_dens = TRUE)

## End(Not run)
#-----
# Example 2 : Surrogate-based multi-objective Optimization including a cheap function
#-----
set.seed(42)
library(DiceDesign)

d <- 2

fname <- P1
n.grid <- 19
test.grid <- expand.grid(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid))
nappr <- 15
design.grid <- maximinESE_LHS(lhsDesign(nappr, d, seed = 42)$design)$design
response.grid <- t(apply(design.grid, 1, fname))

mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])
model <- list(mf1, mf2)

nsteps <- 1
lower <- rep(0, d)
upper <- rep(1, d)

# Optimization with fastfun: hypervolume with discrete search

optimcontrol <- list(method = "discrete", candidate.points = test.grid)
omEGO2 <- GParetooptim(model = model, fn = fname, cheapfn = branin, crit = "SMS",
                      nsteps = nsteps, lower = lower, upper = upper,
                      optimcontrol = optimcontrol)

print(omEGO2$par)
print(omEGO2$values)

## Not run:
plotGPareto(omEGO2)

```

```

#-----
# Example 3 : Surrogate-based multi-objective Optimization (4 objectives)
#-----
set.seed(42)
library(DiceDesign)

d <- 5

fname <- DTLZ3
nappr <- 25
design.grid <- maximinESE_LHS(lhsDesign(nappr, d, seed = 42)$design)$design
response.grid <- t(apply(design.grid, 1, fname, nobj = 4))
mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])
mf3 <- km(~., design = design.grid, response = response.grid[,3])
mf4 <- km(~., design = design.grid, response = response.grid[,4])

# Optimization
nsteps <- 5
lower <- rep(0, d)
upper <- rep(1, d)
omEG03 <- GParetooptim(model = list(mf1, mf2, mf3, mf4), fn = fname, crit = "EMI",
                        nsteps = nsteps, lower = lower, upper = upper, nobj = 4)

print(omEG03$par)
print(omEG03$values)
plotGPareto(omEG03)

#-----
# Example 4 : quantification of uncertainty on Pareto front
#-----
library(DiceDesign)
set.seed(42)

nvar <- 2

# Test function P1
fname <- "P1"

# Initial design
nappr <- 10
design.grid <- maximinESE_LHS(lhsDesign(nappr, nvar, seed = 42)$design)$design
response.grid <- t(apply(design.grid, 1, fname))

PF <- t(nondominated_points(t(response.grid)))

# kriging models : matern5_2 covariance structure, linear trend, no nugget effect
mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])

# Conditional simulations generation with random sampling points
nsim <- 100 # increase for better results
npointssim <- 1000 # increase for better results
Simu_f1 <- matrix(0, nrow = nsim, ncol = npointssim)

```

```

Simu_f2 <- matrix(0, nrow = nsim, ncol = npointssim)
design.sim <- array(0, dim = c(npointssim, nvar, nsim))

for(i in 1:nsim){
  design.sim[, ,i] <- matrix(runif(nvar*npointssim), nrow = npointssim, ncol = nvar)
  Simu_f1[i,] <- simulate(mf1, nsim = 1, newdata = design.sim[, ,i], cond = TRUE,
                        checkNames = FALSE, nugget.sim = 10^-8)
  Simu_f2[i,] <- simulate(mf2, nsim = 1, newdata = design.sim[, ,i], cond = TRUE,
                        checkNames = FALSE, nugget.sim = 10^-8)
}

# Computation of the attainment function and Vorob'ev Expectation
CPF1 <- CPF(Simu_f1, Simu_f2, response.grid, paretoFront = PF)

summary(CPF1)

plot(CPF1)

# Display of the symmetric deviation function
plotSymDevFun(CPF1)

## End(Not run)

```

checkPredict

Prevention of numerical instability for a new observation

Description

Check that the new point is not too close to already known observations to avoid numerical issues. Closeness can be estimated with several distances.

Usage

```
checkPredict(x, model, threshold = 1e-04, distance = "euclidean",
            type = "UK")
```

Arguments

x	a vector representing the input to check, alternatively a matrix with one point per row,
model	list of objects of class <code>km</code> , one for each objective functions,
threshold	optional value for the minimal distance to an existing observation, default to $1e-4$,
distance	selection of the distance between new observations, between "euclidean" (default), "none", "covdist" and "covratio", see details,
type	"SK" or "UK" (default), depending whether uncertainty related to trend estimation has to be taken into account.

Details

If the distance between x and the closest observations in `model` is below `threshold`, x should not be evaluated to avoid numerical instabilities. The distance can simply be the Euclidean distance or the canonical distance associated with the kriging predictive covariance k :

$$d(x, y) = \sqrt{k(x, x) - 2k(x, y) + k(y, y)}.$$

The last solution is the ratio between the prediction variance at x and the variance of the process. none can be used, e.g., if points have been selected already.

Value

TRUE if the point should not be tested.

 CPF

Conditional Pareto Front simulations

Description

Compute (on a regular grid) the empirical attainment function from conditional simulations of Gaussian processes corresponding to two objectives. This is used to estimate the Vorob'ev expectation of the attained set and the Vorob'ev deviation.

Usage

```
CPF(fun1sims, fun2sims, response, paretoFront = NULL, f1lim = NULL,
    f2lim = NULL, refPoint = NULL, n.grid = 100,
    compute.VorobExp = TRUE, compute.VorobDev = TRUE)
```

Arguments

<code>fun1sims</code>	numeric matrix containing the conditional simulations of the first output (one sample in each row),
<code>fun2sims</code>	numeric matrix containing the conditional simulations of the second output (one sample in each row),
<code>response</code>	a matrix containing the value of the two objective functions, one output per row,
<code>paretoFront</code>	optional matrix corresponding to the Pareto front of the observations. It is estimated from <code>response</code> if not provided,
<code>f1lim</code>	optional vector (see details),
<code>f2lim</code>	optional vector (see details),
<code>refPoint</code>	optional vector (see details),
<code>n.grid</code>	integer determining the grid resolution,
<code>compute.VorobExp</code>	optional boolean indicating whether the Vorob'ev Expectation should be computed. Default is TRUE,
<code>compute.VorobDev</code>	optional boolean indicating whether the Vorob'ev deviation should be computed. Default is TRUE.

Details

Works with two objectives. The user can provide locations of grid lines for computation of the attainment function with vectors `f1lim` and `f2lim`, in the form of regularly spaced points. It is possible to provide only `refPoint` as a reference for hypervolume computations. When missing, values are determined from the axis-wise extrema of the simulations.

Value

A list which is given the S3 class "CPF".

- `x`, `y`: locations of grid lines at which the values of the attainment are computed,
- `values`: numeric matrix containing the values of the attainment on the grid,
- `PF`: matrix corresponding to the Pareto front of the observations,
- `responses`: matrix containing the value of the two objective functions, one objective per column,
- `fun1sims`, `fun2sims`: conditional simulations of the first/second output,
- `VE`: Vorob'ev expectation, computed if `compute.VorobExp = TRUE` (default),
- `beta_star`: Vorob'ev threshold, computed if `compute.VorobExp = TRUE` (default),
- `VD`: Vorob'ev deviation, computed if `compute.VorobDev = TRUE` (default),

References

M. Binois, D. Ginsbourger and O. Roustant (2015), Quantifying Uncertainty on Pareto Fronts with Gaussian process conditional simulations, *European Journal of Operational Research*, 243(2), 386-394.

C. Chevalier (2013), *Fast uncertainty reduction strategies relying on Gaussian process models*, University of Bern, PhD thesis.

I. Molchanov (2005), *Theory of random sets*, Springer.

See Also

Methods `coef`, `summary` and `plot` can be used to get the coefficients from a CPF object, to obtain a summary or to display the attainment function (with the Vorob'ev expectation if `compute.VorobExp` is TRUE).

Examples

```
library(DiceDesign)
set.seed(42)

nvar <- 2

fname <- "P1" # Test function

# Initial design
nappr <- 10
```



```

design.grid <- maximinESE_LHS(lhsDesign(nappr, nvar, seed = 42)$design)$design
response.grid <- t(apply(design.grid, 1, fname))

# kriging models: matern5_2 covariance structure, linear trend, no nugget effect
mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])

# Conditional simulations generation with random sampling points
nsim <- 40
npointssim <- 150 # increase for better results
Simu_f1 <- matrix(0, nrow = nsim, ncol = npointssim)
Simu_f2 <- matrix(0, nrow = nsim, ncol = npointssim)
design.sim <- array(0, dim = c(npointssim, nvar, nsim))

for(i in 1:nsim){
  design.sim[, ,i] <- matrix(runif(nvar*npointssim), nrow = npointssim, ncol = nvar)
  Simu_f1[i,] <- simulate(mf1, nsim = 1, newdata = design.sim[, ,i], cond = TRUE,
                        checkNames = FALSE, nugget.sim = 10^-8)
  Simu_f2[i,] <- simulate(mf2, nsim = 1, newdata = design.sim[, ,i], cond = TRUE,
                        checkNames = FALSE, nugget.sim = 10^-8)
}

# Attainment and Voreb'ev expectation + deviation estimation
CPF1 <- CPF(Simu_f1, Simu_f2, response.grid)

# Details about the Vorob'ev threshold and Vorob'ev deviation
summary(CPF1)

# Graphics
plot(CPF1)

```

crit_EHI

Expected Hypervolume Improvement with m objectives

Description

Multi-objective Expected Hypervolume Improvement with respect to the current Pareto front. With two objectives the analytical formula is used, while Sample Average Approximation (SAA) is used with more objectives. To avoid numerical instabilities, the new point is penalized if it is too close to an existing observation.

Usage

```
crit_EHI(x, model, paretoFront = NULL, critcontrol = list(nb.samp = 50,
  seed = 42), type = "UK")
```

Arguments

x	a vector representing the input for which one wishes to calculate EHI, alternatively a matrix with one point per row,
model	list of objects of class <code>km</code> , one for each objective functions,
paretoFront	(optional) matrix corresponding to the Pareto front of size $[n.pareto \times n.obj]$, or any reference set of observations,
critcontrol	optional list with arguments: <ul style="list-style-type: none"> • <code>nb.samp</code> number of random samples from the posterior distribution (with more than two objectives), default to 50, increasing gives more reliable results at the cost of longer computation time; • <code>seed</code> seed used for the random samples (with more than two objectives); • <code>refPoint</code> reference point for Hypervolume Expected Improvement; • <code>extendper</code> if no reference point <code>refPoint</code> is provided, for each objective it is fixed to the maximum over the Pareto front plus <code>extendper</code> times the range, Default value to 0.2, corresponding to 1.1 for a scaled objective with a Pareto front in $[0, 1]^{n.obj}$. Options for the <code>checkPredict</code> function: <code>threshold</code> ($1e-4$) and <code>distance</code> (<code>covdist</code>) are used to avoid numerical issues occurring when adding points too close to the existing ones.
type	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account.

Details

The computation of the analytical formula with two objectives is adapted from the Matlab source code by Michael Emmerich and Andre Deutz, LIACS, Leiden University, 2010 available here : http://liacs.leidenuniv.nl/~csmoda/code/HV_based_expected_improvement.zip.

Value

The Expected Hypervolume Improvement at `x`.

References

J. D. Svenson (2011), *Computer Experiments: Multiobjective Optimization and Sensitivity Analysis*, Ohio State University, PhD thesis.

M. T. Emmerich, A. H. Deutz, J. W. Klinkenberg (2011), Hypervolume-based expected improvement: Monotonicity properties and exact computation, *Evolutionary Computation (CEC)*, 2147-2154.

See Also

`EI` from package `DiceOptim`, `crit_EMI`, `crit_SUR`, `crit_SMS`.

Examples

```

#-----
# Expected Hypervolume Improvement surface associated with the "P1" problem at a 15 points design
#-----
set.seed(25468)
library(DiceDesign)

n_var <- 2
f_name <- "P1"
n.grid <- 26
test.grid <- expand.grid(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid))
n_appr <- 15
design.grid <- round(maximinESE_LHS(lhsDesign(n_appr, n_var, seed = 42)$design)$design, 1)
response.grid <- t(apply(design.grid, 1, f_name))
Front_Pareto <- t(nondominated_points(t(response.grid)))
mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])

EHI_grid <- crit_EMI(x = as.matrix(test.grid), model = list(mf1, mf2),
                    critcontrol = list(refPoint = c(300,0)))

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
               matrix(EHI_grid, n.grid), main = "Expected Hypervolume Improvement",
               xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,
               plot.axes = {axis(1); axis(2);
                           points(design.grid[,1], design.grid[,2], pch = 21, bg = "white")
                           }
               )

```

crit_EMI

*Expected Maximin Improvement with m objectives***Description**

Expected Maximin Improvement with respect to the current Pareto front with Sample Average Approximation. The semi-analytical formula is used in the bi-objective scale if the Pareto front is in $[-2,2]^2$, for numerical stability reasons. To avoid numerical instabilities, the new point is penalized if it is too close to an existing observation.

Usage

```
crit_EMI(x, model, paretoFront = NULL, critcontrol = list(nb.samp = 50,
                                                         seed = 42), type = "UK")
```

Arguments

x a vector representing the input for which one wishes to calculate EMI,
 model list of objects of class `km`, one for each objective functions,

paretoFront	(optional) matrix corresponding to the Pareto front of size $[n.pareto \times n.obj]$, or any reference set of observations,
critcontrol	optional list with arguments (for more than 2 objectives only): <ul style="list-style-type: none"> • nb.samp number of random samples from the posterior distribution, default to 50, increasing gives more reliable results at the cost of longer computation time; • seed seed used for the random samples. Options for the <code>checkPredict</code> function: <code>threshold</code> ($1e-4$) and <code>distance</code> (<code>covdist</code>) are used to avoid numerical issues occurring when adding points too close to the existing ones.
type	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account.

Details

It is recommended to scale objectives, e.g. to $[0, 1]$. If the Pareto front does not belong to $[-2,2]^2$, then SAA is used.

Value

The Expected Maximin Improvement at x .

References

J. D. Svenson & T. J. Santner (2010), Multiobjective Optimization of Expensive Black-Box Functions via Expected Maximin Improvement, Technical Report.

J. D. Svenson (2011), *Computer Experiments: Multiobjective Optimization and Sensitivity Analysis*, Ohio State University, PhD thesis.

See Also

[EI](#) from package `DiceOptim`, [crit_EHI](#), [crit_SUR](#), [crit_SMS](#).

Examples

```
#-----
# Expected Maximin Improvement surface associated with the "P1" problem at a 15 points design
#-----
set.seed(25468)
library(DiceDesign)

n_var <- 2
f_name <- "P1"
n_grid <- 21
test_grid <- expand.grid(seq(0, 1, length.out = n_grid), seq(0, 1, length.out = n_grid))
n_appr <- 15
design_grid <- round(maximinESE_LHS(lhsDesign(n_appr, n_var, seed = 42)$design)$design, 1)
response_grid <- t(apply(design_grid, 1, f_name))
```

```

Front_Pareto <- t(nondominated_points(t(response.grid)))
mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])

EMI_grid <- apply(test.grid, 1, crit_EMI, model = list(mf1, mf2), paretoFront = Front_Pareto,
                 critcontrol = list(nb_samp = 20))

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(EMI_grid, nrow = n.grid), main = "Expected Maximin Improvement",
              xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,
              plot.axes = {axis(1); axis(2);
                          points(design.grid[,1], design.grid[,2], pch = 21, bg = "white")
                          }
              )

```

crit_optimizer

Maximization of multiobjective infill criterion

Description

Given a list of objects of class `km` and a set of tuning parameters (`lower`, `upper` and `critcontrol`), `crit_optimizer` performs the maximization of an infill criterion and delivers the next point to be visited in a multi-objective EGO-like procedure.

The latter maximization relies either on a genetic algorithm using derivatives, `genoud`, particle swarm algorithm `pso`, exhaustive search at pre-specified points or on a user defined method. It is important to remark that the information needed about the objective function reduces here to the vector of response values embedded in the models (no call to the objective functions or simulators (except with `cheapfn`)).

Usage

```
crit_optimizer(crit = "SMS", model, lower, upper, cheapfn = NULL,
              type = "UK", paretoFront = NULL, critcontrol = NULL,
              optimcontrol = NULL)
```

Arguments

<code>crit</code>	sampling criterion. Four choices are available : "SMS", "EHI", "EMI" and "SUR",
<code>model</code>	list of objects of class <code>km</code> , one for each objective functions,
<code>lower</code>	vector of lower bounds for the variables to be optimized over,
<code>upper</code>	vector of upper bounds for the variables to be optimized over,
<code>cheapfn</code>	optional additional fast-to-evaluate objective function (handled next with class <code>fastfun</code>), which does not need a kriging model,
<code>type</code>	"SK" or "UK" (default), depending whether uncertainty related to trend estimation has to be taken into account.

paretoFront	(optional) matrix corresponding to the Pareto front of size <code>[n.pareto x n.obj]</code> , or any reference set of observations,
critcontrol	optional list of control parameters for criterion <code>crit</code> , see details. Options for the <code>checkPredict</code> function: <code>threshold</code> ($1e-4$) and <code>distance</code> (<code>covdist</code>) are used to avoid numerical issues occurring when adding points too close to the existing ones.
optimcontrol	optional list of control parameters for optimization of the selected infill criterion. "method" set the optimization method; one can choose between "discrete", "pso" and "genoud" or a user defined method name (passed to <code>match.fun</code>). For each method, further parameters can be set. For "discrete", one has to provide the argument "candidate.points". For "pso", one can control the maximum number of iterations "maxit" (400) and the population size "s" (default: <code>max(20, floor(10+2*sqrt(length(dim))))</code>) (see <code>psoptim</code>). For "genoud", one can control, among others, "pop.size" (default: <code>[N = 3*2^dim for dim < 6 and N = 32*dim otherwise]</code>), "max.generations" (12), "wait.generations" (2), "BFGSburnin" (2), <code>BFGSmaxit</code> (N) and <code>solution.tolerance</code> ($1e-21$) of function "genoud" (see <code>genoud</code>). Numbers into brackets are the default values. For a user defined method, it must have arguments like the default <code>optim</code> method, i.e. <code>par</code> , <code>fn</code> , <code>lower</code> , <code>upper</code> , ... and possibly <code>control</code> , and return a list with <code>par</code> and <code>value</code> . A <code>trace</code> argument is available, it can be set to 0 to suppress all messages, to 1 (default) for displaying the optimization progresses, and >1 for the highest level of details.

Details

Extension of the function `max_EI` for multi-objective optimization.
Available infill criteria with `crit` are :

- Expected Hypervolume Improvement (EHI) `crit_EHI`,
- SMS criterion (SMS) `crit_SMS`,
- Expected Maximin Improvement (EMI) `crit_EMI`,
- Stepwise Uncertainty Reduction of the excursion volume (SUR) `crit_SUR`

Depending on the selected criterion, parameters such as a reference point for SMS and EHI or arguments for `integration_design_optim` with SUR can be given with `critcontrol`. Also options for `checkPredict` are available. More precisions are given in the corresponding help pages.

Value

A list with components:

- `par`: The best set of parameters found,
- `value`: The value of expected improvement at `par`.

References

W.R. Jr. Mebane and J.S. Sekhon (2011), Genetic optimization using derivatives: The rgenoud package for R, *Journal of Statistical Software*, 42(11), 1-26

D.R. Jones, M. Schonlau, and W.J. Welch (1998), Efficient global optimization of expensive black-box functions, *Journal of Global Optimization*, 13, 455-492.

Examples

```
## Not run:
#-----
# EHI surface associated with the "P1" problem at a 15 points design
#-----

set.seed(25468)
library(DiceDesign)

d <- 2
n.obj <- 2
fname <- "P1"
n.grid <- 51
test.grid <- expand.grid(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid))
nappr <- 15
design.grid <- round(maximinESE_LHS(lhsDesign(nappr, d, seed = 42)$design)$design, 1)
response.grid <- t(apply(design.grid, 1, fname))
paretoFront <- t(nondominated_points(t(response.grid)))
mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])
model <- list(mf1, mf2)

EHI_grid <- apply(test.grid, 1, crit_EHI, model = list(mf1, mf2),
                 critcontrol = list(refPoint = c(300, 0)))

lower <- rep(0, d)
upper <- rep(1, d)

omEGO <- crit_optimizer(crit = "EHI", model = model, lower = lower, upper = upper,
                      optimcontrol = list(method = "genoud", pop.size = 200, BFGSburnin = 2),
                      critcontrol = list(refPoint = c(300, 0)))

print(omEGO)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(EHI_grid, nrow = n.grid), main = "Expected Hypervolume Improvement",
                  xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,
                  plot.axes = {axis(1); axis(2);
                              points(design.grid[, 1], design.grid[, 2], pch = 21, bg = "white");
                              points(omEGO$par, col = "red", pch = 4)
                              }
              )
```

```

#-----
# SMS surface associated with the "P1" problem at a 15 points design
#-----

SMS_grid <- apply(test.grid, 1, crit_SMS, model = model,
                 critcontrol = list(refPoint = c(300, 0)))

lower <- rep(0, d)
upper <- rep(1, d)

omEG02 <- crit_optimizer(crit = "SMS", model = model, lower = lower, upper = upper,
                       optimcontrol = list(method="genoud", pop.size = 200, BFGSburnin = 2),
                       critcontrol = list(refPoint = c(300, 0)))

print(omEG02)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(pmax(0, SMS_grid), nrow = n.grid), main = "SMS Criterion (>0)",
              xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,
              plot.axes = {axis(1); axis(2);
                          points(design.grid[, 1], design.grid[, 2], pch = 21, bg = "white");
                          points(omEG02$par, col = "red", pch = 4)
                          }
              )

#-----
# Maximin Improvement surface associated with the "P1" problem at a 15 points design
#-----

EMI_grid <- apply(test.grid, 1, crit_EMI, model = model,
                 critcontrol = list(nb_samp = 20, type = "EMI"))

lower <- rep(0, d)
upper <- rep(1, d)

omEG03 <- crit_optimizer(crit = "EMI", model = model, lower = lower, upper = upper,
                       optimcontrol = list(method = "genoud", pop.size = 200, BFGSburnin = 2))

print(omEG03)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(EMI_grid, nrow = n.grid), main = "Expected Maximin Improvement",
              xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,
              plot.axes = {axis(1); axis(2);
                          points(design.grid[, 1], design.grid[, 2], pch = 21, bg = "white");
                          points(omEG03$par, col = "red", pch = 4)
                          }
              )

#-----
# crit_SUR surface associated with the "P1" problem at a 15 points design
#-----
library(KrigInv)

```



```

integration.param <- integration_design_optim(lower = c(0, 0), upper = c(1, 1), model = model)
integration.points <- as.matrix(integration.param$integration.points)
integration.weights <- integration.param$integration.weights

precalc.data <- list()
mn.X <- sn.X <- matrix(0, n.obj, nrow(integration.points))

for (i in 1:n.obj){
  p.tst.all <- predict(model[[i]], newdata = integration.points, type = "UK",
                      checkNames = FALSE)
  mn.X[i,] <- p.tst.all$mean
  sn.X[i,] <- p.tst.all$sd
  precalc.data[[i]] <- precomputeUpdateData(model[[i]], integration.points)
}
critcontrol <- list(integration.points = integration.points,
                  integration.weights = integration.weights,
                  mn.X = mn.X, sn.X = sn.X, precalc.data = precalc.data)
EEV_grid <- apply(test.grid, 1, crit_SUR, model=model, paretoFront = paretoFront,
                 critcontrol = critcontrol)
lower <- rep(0, d)
upper <- rep(1, d)

omEG04 <- crit_optimizer(crit = "SUR", model = model, lower = lower, upper = upper,
                       optimcontrol = list(method = "genoud", pop.size = 200, BFGSburnin = 2))
print(omEG04)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
              matrix(pmax(0,EEV_grid), n.grid), main = "EEV criterion", nlevels = 50,
              xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,
              plot.axes = {axis(1); axis(2);
                          points(design.grid[,1], design.grid[,2], pch = 21, bg = "white")
                          points(omEG04$par, col = "red", pch = 4)
                          }
              )

# example using user defined optimizer, here L-BFGS-B from base optim
userOptim <- function(par, fn, lower, upper, control, ...){
  return(optim(par = par, fn = fn, method = "L-BFGS-B", lower = lower, upper = upper,
              control = control, ...))
}
omEG04bis <- crit_optimizer(crit = "SUR", model = model, lower = lower, upper = upper,
                          optimcontrol = list(method = "userOptim"))
print(omEG04bis)

#-----
# crit_SMS surface with problem "P1" with 15 design points, using cheapfn
#-----

# Optimization with fastfun: SMS with discrete search
# Separation of the problem P1 in two objectives:
# the first one to be kriged, the second one with fastobj

```

```

# Definition of the fastfun
f2 <- function(x){
  return(P1(x)[2])
}

SMS_grid_cheap <- apply(test.grid, 1, crit_SMS, model = list(mf1, fastfun(f2, design.grid)),
  paretoFront = paretoFront, critcontrol = list(refPoint = c(300, 0)))

optimcontrol <- list(method = "pso")
model2 <- list(mf1)
omEG05 <- crit_optimizer(crit = "SMS", model = model2, lower = lower, upper = upper,
  cheapfn = f2, critcontrol = list(refPoint = c(300, 0)),
  optimcontrol = list(method = "genoud", pop.size = 200, BFGSburnin = 2))
print(omEG05)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
  matrix(pmax(0, SMS_grid_cheap), nrow = n.grid), nlevels = 50,
  main = "SMS criterion with cheap 2nd objective (>0)", xlab = expression(x[1]),
  ylab = expression(x[2]), color = terrain.colors,
  plot.axes = {axis(1); axis(2);
  points(design.grid[,1], design.grid[,2], pch = 21, bg = "white")
  points(omEG05$par, col = "red", pch = 4)
  }
)

## End(Not run)

```

crit_SMS

Analytical expression of the SMS-EGO criterion with $m > 1$ objectives

Description

Computes a slightly modified infill Criterion of the SMS-EGO. To avoid numerical instabilities, an additional penalty is added to the new point if it is too close to an existing observation.

Usage

```
crit_SMS(x, model, paretoFront = NULL, critcontrol = NULL,
  type = "UK")
```

Arguments

x	a vector representing the input for which one wishes to calculate the criterion,
model	a list of objects of class <code>km</code> (one for each objective),
paretoFront	(optional) matrix corresponding to the Pareto front of size <code>[n.pareto x n.obj]</code> , or any reference set of observations,
critcontrol	list with arguments:

- currentHV current hypervolume;
- refPoint reference point for hypervolume computations;
- extendper if no reference point refPoint is provided, for each objective it is fixed to the maximum over the Pareto front plus extendper times the range. Default value to 0.2, corresponding to 1.1 for a scaled objective with a Pareto front in $[\theta, 1]^n$.obj;
- epsilon optional value to use in additive epsilon dominance;
- gain optional gain factor for sigma.

Options for the `checkPredict` function: `threshold` ($1e-4$) and `distance` (`covdist`) are used to avoid numerical issues occurring when adding points too close to the existing ones.

`type` "SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account.

Value

Value of the criterion.

References

W. Ponweiser, T. Wagner, D. Biermann, M. Vincze (2008), Multiobjective Optimization on a Limited Budget of Evaluations Using Model-Assisted S-Metric Selection, *Parallel Problem Solving from Nature*, pp. 784-794. Springer, Berlin.

T. Wagner, M. Emmerich, A. Deutz, W. Ponweiser (2010), On expected-improvement criteria for model-based multi-objective optimization. *Parallel Problem Solving from Nature*, pp. 718-727. Springer, Berlin.

See Also

[crit_EHI](#), [crit_SUR](#), [crit_EMI](#).

Examples

```
#-----
# SMS-EGO surface associated with the "P1" problem at a 15 points design
#-----
set.seed(25468)
library(DiceDesign)

n_var <- 2
f_name <- "P1"
n_grid <- 26
test_grid <- expand.grid(seq(0, 1, length.out = n_grid), seq(0, 1, length.out = n_grid))
n_appr <- 15
design_grid <- round(maximinESE_LHS(lhsDesign(n_appr, n_var, seed = 42)$design)$design, 1)
response_grid <- t(apply(design_grid, 1, f_name))
PF <- t(nondominated_points(t(response_grid)))
mf1 <- km(~., design = design_grid, response = response_grid[,1])
```

```

mf2 <- km(~., design = design.grid, response = response.grid[,2])

model <- list(mf1, mf2)
critcontrol <- list(refPoint = c(300, 0), currentHV = dominated_hypervolume(t(PF), c(300, 0)))
SMSEGO_grid <- apply(test.grid, 1, crit_SMS, model = model,
                    paretoFront = PF, critcontrol = critcontrol)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
              matrix(pmax(0, SMSEGO_grid), nrow = n.grid), nlevels = 50,
              main = "SMS-EGO criterion (positive part)", xlab = expression(x[1]),
              ylab = expression(x[2]), color = terrain.colors,
              plot.axes = {axis(1); axis(2);
                          points(design.grid[,1],design.grid[,2], pch = 21, bg = "white")
                          }
              )

```

crit_SUR

*Analytical expression of the SUR criterion for two or three objectives.***Description**

Computes the SUR criterion (Expected Excursion Volume Reduction) at point x for 2 or 3 objectives. To avoid numerical instabilities, the new point is penalized if it is too close to an existing observation.

Usage

```
crit_SUR(x, model, paretoFront = NULL, critcontrol = NULL,
        type = "UK")
```

Arguments

<code>x</code>	a vector representing the input for which one wishes to calculate the criterion,
<code>model</code>	a list of objects of class <code>km</code> (one for each objective),
<code>paretoFront</code>	(optional) matrix corresponding to the Pareto front of size $[n.pareto \times n.obj]$, or any reference set of observations,
<code>critcontrol</code>	list with two possible options.

A) One can use the four following arguments:

- `integration.points`, matrix of integration points of size $[n.integ.pts \times d]$;
- `integration.weights`, vector of integration weights of length $n.integ.pts$;
- `mn.X` and `sn.X`, matrices of kriging means and sd, each of size $[n.obj \times n.integ.pts]$;
- `precalc.data`, list of precalculated data (based on kriging models at integration points) for faster computation.

B) Alternatively, one can define arguments passed to `integration_design_optim`: `SURcontrol` (optional), `lower`, `upper`, `min.prob` (optional). This is slower since arguments of `A`, used in the function, are then recomputed each time (note that this is not the case when called from `GParetooptim` and `crit_optimizer`).

Options for the `checkPredict` function: `threshold` ($1e-4$) and `distance` (`covdist`) are used to avoid numerical issues occurring when adding points too close to the existing ones.

`type` "SK" or "UK" (default), depending whether uncertainty related to trend estimation has to be taken into account.

Value

Value of the criterion.

References

V. Picheny (2014), Multiobjective optimization using Gaussian process emulators via stepwise uncertainty reduction, *Statistics and Computing*.

See Also

`crit_EHI`, `crit_SMS`, `crit_EMI`.

Examples

```
#-----
# crit_SUR surface associated with the "P1" problem at a 15 points design
#-----
set.seed(25468)
library(DiceDesign)
library(KrigInv)

n_var <- 2
n_obj <- 2
f_name <- "P1"
n_grid <- 14
test.grid <- expand.grid(seq(0, 1, length.out = n_grid), seq(0, 1, length.out = n_grid))
n_appr <- 15
design.grid <- round(maximinESE_LHS(lhsDesign(n_appr, n_var, seed = 42)$design)$design, 1)
response.grid <- t(apply(design.grid, 1, f_name))
paretoFront <- t(nondominated_points(t(response.grid)))
mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])

model <- list(mf1, mf2)

integration.param <- integration_design_optim(lower = c(0, 0), upper = c(1, 1), model = model)
integration.points <- as.matrix(integration.param$integration.points)
integration.weights <- integration.param$integration.weights
```

```

precalc.data <- list()
mn.X <- sn.X <- matrix(0, nrow = n.obj, ncol = nrow(integration.points))

for (i in 1:n.obj){
  p.tst.all <- predict(model[[i]], newdata = integration.points, type = "UK", checkNames = FALSE)
  mn.X[i,] <- p.tst.all$mean
  sn.X[i,] <- p.tst.all$sd
  precalc.data[[i]] <- precomputeUpdateData(model[[i]], integration.points)
}

critcontrol <- list(integration.points = integration.points,
                  integration.weights = integration.weights,
                  mn.X = mn.X, sn.X = sn.X, precalc.data = precalc.data)
## Alternatively: critcontrol <- list(lower = rep(0, n_var), upper = rep(1,n_var))

EEV_grid <- apply(test.grid, 1, crit_SUR, model = model, paretoFront = paretoFront,
                 critcontrol = critcontrol)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
              matrix(pmax(0,EEV_grid), nrow = n.grid), main = "EEV criterion",
              xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,
              plot.axes = {axis(1); axis(2);
                          points(design.grid[,1], design.grid[,2], pch = 21, bg = "white")
                          }
              )

```

easyGParetoptim

EGO algorithm for multiobjective optimization

Description

User-friendly wrapper of the function [GParetoptim](#). Generates initial DOEs and kriging models (objects of class [km](#)), and executes `nsteps` iterations of multiobjective EGO methods.

Usage

```

easyGParetoptim(fn, cheapfn = NULL, budget, lower, upper, par = NULL,
               value = NULL, noise.var = NULL, control = list(method = "SMS",
               trace = 1, inneroptim = "pso", maxit = 100, seed = 42), ...)

```

Arguments

<code>fn</code>	the multi-objective function to be minimized (vectorial output), found by a call to match.fun , see details,
<code>cheapfn</code>	optional additional fast-to-evaluate objective function (handled next with class fastfun), which does not need a kriging model, handled by a call to match.fun ,
<code>budget</code>	total number of calls to the objective function,

lower	vector of lower bounds for the variables to be optimized over,
upper	vector of upper bounds for the variables to be optimized over,
par	initial design of experiments. If not provided, par is taken as a maximin LHD with budget/3 points,
value	initial set of objective observations $fn(par)$. Computed if not provided. Not that value may NOT contain any cheapfn value,
noise.var	optional noise variance, for noisy objectives fn. If not NULL, either a scalar (constant noise, identical for all objectives), a vector (constant noise, different for each objective) or a function (type closure) with vectorial output (variable noise, different for each objective). Alternatively, set noise.var="given_by_fn", see details.
control	an optional list of control parameters. See "Details",
...	additional parameters to be given to the objective fn.

Details

Does not require specific knowledge on kriging models (objects of class [km](#)).

The problem considered is of the form: $\min f(x) = f_1(x), \dots, f_p(x)$. The control argument is a list that can supply any of the following optional components:

- method: choice of multiobjective improvement function: "SMS", "EHI", "EMI" or "SUR" (see [crit_SMS](#), [crit_EHI](#), [crit_EMI](#), [crit_SUR](#)),
- trace: if positive, tracing information on the progress of the optimization is produced (1 (default) for general progress, >1 for more details, e.g., warnings from [genoud](#)),
- inneroptim: choice of the inner optimization algorithm: "genoud", "pso" or "random" (see [genoud](#) and [psoptim](#)),
- maxit: maximum number of iterations of the inner loop,
- seed: to fix the random variable generator,
- refPoint: reference point for hypervolume computations (for "SMS" and "EHI" methods),
- extendper: if no reference point refPoint is provided, for each objective it is fixed to the maximum over the Pareto front plus extendper times the range. Default value to 0.2, corresponding to 1.1 for a scaled objective with a Pareto front in $[0, 1]^n$.obj.

If noise.var="given_by_fn", fn must return a list of two vectors, the first being the objective functions and the second the corresponding noise variances. See examples in [GParetoptim](#).

For additional details or other possible arguments, see [GParetoptim](#).

Display of results and various post-processings are available with [plotGPareto](#).

Value

A list with components:

- `par`: all the non-dominated points found,
- `value`: the matrix of objective values at the points given in `par`,
- `history`: a list containing all the points visited by the algorithm (X) and their corresponding objectives (y),
- `model`: a list of objects of class `km`, corresponding to the last kriging models fitted.

Note that in the case of noisy problems, `value` and `history$y.denoised` are denoised values. The original observations are available in the slot `history$y`.

Author(s)

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References

M. T. Emmerich, A. H. Deutz, J. W. Klinkenberg (2011), Hypervolume-based expected improvement: Monotonicity properties and exact computation, *Evolutionary Computation (CEC)*, 2147-2154.

V. Picheny (2015), Multiobjective optimization using Gaussian process emulators via stepwise uncertainty reduction, *Statistics and Computing*, 25(6), 1265-1280.

T. Wagner, M. Emmerich, A. Deutz, W. Ponweiser (2010), On expected-improvement criteria for model-based multi-objective optimization. *Parallel Problem Solving from Nature*, 718-727, Springer, Berlin.

J. D. Svenson (2011), *Computer Experiments: Multiobjective Optimization and Sensitivity Analysis*, Ohio State university, PhD thesis.

M. Binois, V. Picheny (2019), GPareto: An R Package for Gaussian-Process-Based Multi-Objective Optimization and Analysis, *Journal of Statistical Software*, 89(8), 1–30.

Examples

```
#-----
# 2D objective function, 4 cases
#-----
## Not run:
set.seed(25468)
n_var <- 2
fname <- ZDT3
lower <- rep(0, n_var)
upper <- rep(1, n_var)

#-----
```



```

# 1- Expected Hypervolume Improvement optimization, using pso
#-----
res <- easyGParetoptim(fn=fname, lower=lower, upper=upper, budget=15,
                      control=list(method="EHI", inneroptim="pso", maxit=20))
par(mfrow=c(1,2))
plotGPareto(res)
title("Pareto Front")
plot(res$history$X, main="Pareto set", col = "red", pch = 20)
points(res$par, col="blue", pch = 17)

#-----
# 2- SMS Improvement optimization using random search, with initial DOE given
#-----
library(DiceDesign)
design.init <- maximinESE_LHS(lhsDesign(10, n_var, seed = 42)$design)$design
response.init <- t(apply(design.init, 1, fname))
res <- easyGParetoptim(fn=fname, par=design.init, value=response.init, lower=lower, upper=upper,
                      budget=15, control=list(method="SMS", inneroptim="random", maxit=100))
par(mfrow=c(1,2))
plotGPareto(res)
title("Pareto Front")
plot(res$history$X, main="Pareto set", col = "red", pch = 20)
points(res$par, col="blue", pch = 17)

#-----
# 3- Stepwise Uncertainty Reduction optimization, with one fast objective function
#-----
fname <- camelback
cheapfn <- function(x) {
  if (is.null(dim(x))) return(-sum(x))
  else return(-rowSums(x))
}
res <- easyGParetoptim(fn=fname, cheapfn=cheapfn, lower=lower, upper=upper, budget=15,
                      control=list(method="SUR", inneroptim="pso", maxit=20))
par(mfrow=c(1,2))
plotGPareto(res)
title("Pareto Front")
plot(res$history$X, main="Pareto set", col = "red", pch = 20)
points(res$par, col="blue", pch = 17)

#-----
# 4- Expected Hypervolume Improvement optimization, using pso, noisy fn
#-----
noise.var <- c(0.1, 0.2)
funnoise <- function(x) {ZDT3(x) + sqrt(noise.var)*rnorm(n=2)}
res <- easyGParetoptim(fn=funnoise, lower=lower, upper=upper, budget=30, noise.var=noise.var,
                      control=list(method="EHI", inneroptim="pso", maxit=20))
par(mfrow=c(1,2))
plotGPareto(res)
title("Pareto Front")
plot(res$history$X, main="Pareto set", col = "red", pch = 20)
points(res$par, col="blue", pch = 17)

```

```

#-----
# 5- Stepwise Uncertainty Reduction optimization, functional noise
#-----
funnoise <- function(x) {ZDT3(x) + sqrt(abs(0.1*x))*rnorm(n=2)}
noise.var <- function(x) return(abs(0.1*x))

res <- easyGParetooptim(fn=funnoise, lower=lower, upper=upper, budget=30, noise.var=noise.var,
                       control=list(method="SUR", inneroptim="pso", maxit=20))

par(mfrow=c(1,2))
plotGPareto(res)
title("Pareto Front")
plot(res$history$X, main="Pareto set", col = "red", pch = 20)
points(res$par, col="blue", pch = 17)

## End(Not run)

```

fastfun

Fast-to-evaluate function wrapper

Description

Modification of an R function to be used with methods `predict` and `update` (similar to a `km` object). It creates an S4 object which contains the values corresponding to evaluations of other costly observations. It is useful when an objective can be evaluated fast.

Usage

```
fastfun(fn, design, response = NULL)
```

Arguments

<code>fn</code>	the evaluator function, found by a call to <code>match.fun</code> ,
<code>design</code>	a data frame representing the design of experiments. The <i>i</i> th row contains the values of the <i>d</i> input variables corresponding to the <i>i</i> th evaluation.
<code>response</code>	optional vector (or 1-column matrix or data frame) containing the values of the 1-dimensional output given by the objective function at the design points.

Value

An object of class `fastfun-class`.

Examples

```

#####
## Example with a fast to evaluate objective
#####
## Not run:
set.seed(25468)
library(DiceDesign)

```

```

d <- 2

fname <- P1
n.grid <- 21
nappr <- 11
design.grid <- maximinESE_LHS(lhsDesign(nappr, d, seed = 42)$design)$design
response.grid <- t(apply(design.grid, 1, fname))
Front_Pareto <- t(nondominated_points(t(response.grid)))

mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])
model <- list(mf1, mf2)

nsteps <- 5
lower <- rep(0, d)
upper <- rep(1, d)

# Optimization reference: SMS with discrete search
optimcontrol <- list(method = "pso")
omEG01 <- GParetoptim(model = model, fn = fname, crit = "SMS", nsteps = nsteps,
                    lower = lower, upper = upper, optimcontrol = optimcontrol)
print(omEG01$par)
print(omEG01$values)
plot(response.grid, xlim = c(0,300), ylim = c(-40,0), pch = 17, col = "blue")
points(omEG01$values, pch = 20, col = "green")

# Optimization with fastfun: SMS with discrete search
# Separation of the problem P1 in two objectives:
# the first one to be kriged, the second one with fastobj
f1 <- function(x){
  if(is.null(dim(x))) x <- matrix(x, nrow = 1)
  b1 <- 15*x[,1] - 5
  b2 <- 15*x[,2]
  return( (b2 - 5.1*(b1/(2*pi))^2 + 5/pi*b1 - 6)^2 + 10*((1 - 1/(8*pi))*cos(b1) + 1))
}

f2 <- function(x){
  if(is.null(dim(x))) x <- matrix(x, nrow = 1)
  b1<-15*x[,1] - 5
  b2<-15*x[,2]
  return(-sqrt((10.5 - b1)*(b1 + 5.5)*(b2 + 0.5))
        - 1/30*(b2 - 5.1*(b1/(2*pi))^2 - 6)^2
        - 1/3*((1 - 1/(8*pi))*cos(b1) + 1))
}

optimcontrol <- list(method = "pso")
model2 <- list(mf1)
omEG02 <- GParetoptim(model = model2, fn = f1, cheapfn = f2, crit = "SMS", nsteps = nsteps,
                    lower = lower, upper = upper, optimcontrol = optimcontrol)
print(omEG02$par)
print(omEG02$values)

```

```
points(omEG02$values, col = "red", pch = 15)

## End(Not run)
```

getDesign *Get design corresponding to an objective target*

Description

Find the design that maximizes the probability of dominating a target given by the user.

Usage

```
getDesign(model, target, lower, upper, optimcontrol = NULL)
```

Arguments

model	list of objects of class <code>km</code> , one for each objective functions,
target	vector corresponding to the desired output in the objective space,
lower	vector of lower bounds for the variables to be optimized over,
upper	vector of upper bounds for the variables to be optimized over,
optimcontrol	optional list of control parameters for optimization of the selected infill criterion. "method" set the optimization method; one can choose between "discrete", "pso" and "genoud". For each method, further parameters can be set. For "discrete", one has to provide the argument "candidate.points". For "pso", one can control the maximum number of iterations "maxit" (400) and the population size "s" (default: $\max(20, \text{floor}(10+2*\sqrt{\text{length}(\text{dim})}))$) (see psoptim). For "genoud", one can control, among others, "pop.size" (default: $[N = 3*2^{\text{dim}}$ for $\text{dim} < 6$ and $N = 32*\text{dim}$ otherwise]), "max.generations" (12), "wait.generations" (2), "BFGSburnin" (2), BFGSmaxit (N) and solution.tolerance ($1e-21$) of function "genoud" (see genoud). Numbers into brackets are the default values.

Value

A list with components:

- par: best design found,
- value: probability that the design dominates the target,
- mean: kriging mean of the objectives at the design,
- sd: prediction standard deviation at the design.

Examples

```
## Not run:

#-----
# Example of interactive optimization
#-----

set.seed(25468)
library(DiceDesign)

d <- 2
n.obj <- 2
fun <- "P1"
n.grid <- 51
test.grid <- expand.grid(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid))
nappr <- 20
design.grid <- round(maximinESE_LHS(lhsDesign(nappr, d, seed = 42)$design)$design, 1)
response.grid <- t(apply(design.grid, 1, fun))
paretoFront <- t(nondominated_points(t(response.grid)))
mf1 <- km(~., design = design.grid, response = response.grid[,1])
mf2 <- km(~., design = design.grid, response = response.grid[,2])
model <- list(mf1, mf2)
lower <- rep(0, d); upper <- rep(1, d)

sol <- GParetoptim(model, fun, crit = "SUR", nsteps = 5, lower = lower, upper = upper)

plotGPareto(sol)

target1 <- c(15, -25)
points(x = target1[1], y = target1[2], col = "black", pch = 13)

nDesign <- getDesign(sol$lastmodel, target = target1, lower = rep(0, d), upper = rep(1, d))
points(t(nDesign$mean), col = "green", pch = 20)

target2 <- c(48, -27)
points(x = target2[1], y = target2[2], col = "black", pch = 13)
nDesign2 <- getDesign(sol$lastmodel, target = target2, lower = rep(0, d), upper = rep(1, d))
points(t(nDesign2$mean), col = "darkgreen", pch = 20)

## End(Not run)
```

GParetoptim

Sequential multi-objective Expected Improvement maximization and model re-estimation, with a number of iterations fixed in advance by the user

Description

Executes `nsteps` iterations of multi-objective EGO methods to objects of class `km`. At each step, kriging models are re-estimated (including covariance parameters re-estimation) based on the initial

design points plus the points visited during all previous iterations; then a new point is obtained by maximizing one of the four multi-objective Expected Improvement criteria available. Handles noiseless and noisy objective functions.

Usage

```
GParetoptim(model, fn, cheapfn = NULL, crit = "SMS", nsteps, lower,
  upper, type = "UK", cov.reestim = TRUE, critcontrol = NULL,
  noise.var = NULL, reinterpolation = NULL,
  optimcontrol = list(method = "genoud", trace = 1), ...)
```

Arguments

model	list of objects of class <code>km</code> , one for each objective functions,
fn	the multi-objective function to be minimized (vectorial output), found by a call to <code>match.fun</code> ,
cheapfn	optional additional fast-to-evaluate objective function (handled next with class <code>fastfun</code>), which does not need a kriging model, handled by a call to <code>match.fun</code> ,
crit	choice of multi-objective improvement function: "SMS", "EHI", "EMI" or "SUR", see details below,
nsteps	an integer representing the desired number of iterations,
lower	vector of lower bounds for the variables to be optimized over,
upper	vector of upper bounds for the variables to be optimized over,
type	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account, see <code>km</code>
cov.reestim	optional boolean specifying if the kriging hyperparameters should be re-estimated at each iteration,
critcontrol	optional list of parameters for criterion <code>crit</code> , see details,
noise.var	noise variance (of the objective functions). Either NULL (noiseless objectives), a scalar (constant noise, identical for all objectives), a vector (constant noise, different for each objective) or a function (type closure) with vectorial output (variable noise, different for each objective). Alternatively, set <code>noise.var="given_by_fn"</code> , see details. If not provided but <code>km</code> models are based on noisy observations, <code>noise.var</code> is taken as the average of <code>model@noise.var</code> .
reinterpolation	Boolean: for noisy problems, indicates whether a reinterpolation model is used, see details,
optimcontrol	an optional list of control parameters for optimization of the selected infill criterion: "method" can be set to "discrete", "pso", "genoud" or a user defined method name (passed to <code>match.fun</code>). For "discrete", a matrix <code>candidate.points</code> must be given. For "pso" and "genoud", specific parameters to the chosen method can also be specified (see <code>genoud</code> and <code>psoptim</code>). A user defined method must have arguments like the default <code>optim</code> method, i.e. <code>par</code> , <code>fn</code> , <code>lower</code> , <code>upper</code> , ... and eventually <code>control</code> . A trace <code>trace</code> argument is available, it can be set to 0 to suppress all messages, to 1 (default) for displaying the optimization progresses, and >1 for the highest level of details.

... additional parameters to be given to the objective fn.

Details

Extension of the function `EGO.nsteps` for multi-objective optimization.
Available infill criteria with `crit` are:

- Expected Hypervolume Improvement (EHI) `crit_EHI`,
- SMS criterion (SMS) `crit_SMS`,
- Expected Maximin Improvement (EMI) `crit_EMI`,
- Stepwise Uncertainty Reduction of the excursion volume (SUR) `crit_SUR`.

Depending on the selected criterion, parameters such as reference point for SMS and EHI or arguments for `integration_design_optim` with SUR can be given with `critcontrol`. Also options for `checkPredict` are available. More precisions are given in the corresponding help pages.

The `reinterpolation=TRUE` setting can be used to handle noisy objective functions. It works with all criteria and is the recommended option. If `reinterpolation=FALSE` and `noise.var!=NULL`, the criteria are used based on a "denoised" Pareto front.

If `noise.var="given_by_fn"`, `fn` must return a list of two vectors, the first being the objective functions and the second the corresponding noise variances (see examples).

Display of results and various post-processings are available with `plotGPareto`.

Value

A list with components:

- `par`: a data frame representing the additional points visited during the algorithm,
- `values`: a data frame representing the response values at the points given in `par`,
- `nsteps`: an integer representing the desired number of iterations (given in argument),
- `lastmodel`: a list of objects of class `km` corresponding to the last kriging models fitted.
- `observations.denoised`: if `noise.var!=NULL`, a matrix representing the mean values of the `km` models at observation points. If a problem occurs during either model updates or criterion maximization, the last working model and corresponding values are returned.

References

M. T. Emmerich, A. H. Deutz, J. W. Klinkenberg (2011), Hypervolume-based expected improvement: Monotonicity properties and exact computation, *Evolutionary Computation (CEC)*, 2147-2154.

V. Picheny (2014), Multiobjective optimization using Gaussian process emulators via stepwise uncertainty reduction, *Statistics and Computing*, 25(6), 1265-1280

T. Wagner, M. Emmerich, A. Deutz, W. Ponweiser (2010), On expected-improvement criteria for model-based multi-objective optimization. *Parallel Problem Solving from Nature*, 718-727,

Springer, Berlin.

J. D. Svenson (2011), *Computer Experiments: Multiobjective Optimization and Sensitivity Analysis*, Ohio State university, PhD thesis. V. Picheny and D. Ginsbourger (2013), *Noisy kriging-based optimization methods: A unified implementation within the DiceOptim package*, *Computational Statistics & Data Analysis*, 71: 1035-1053.

Examples

```

set.seed(25468)
library(DiceDesign)

#####
# NOISELESS PROBLEMS
#####
d <- 2
fname <- ZDT3
n.grid <- 21
test.grid <- expand.grid(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid))
nappr <- 15
design.grid <- maximinESE_LHS(lhsDesign(nappr, d, seed = 42)$design)$design
response.grid <- t(apply(design.grid, 1, fname))
Front_Pareto <- t(nondominated_points(t(response.grid)))

mf1 <- km(~1, design = design.grid, response = response.grid[, 1], lower=c(.1,.1))
mf2 <- km(~., design = design.grid, response = response.grid[, 2], lower=c(.1,.1))
model <- list(mf1, mf2)

nsteps <- 2
lower <- rep(0, d)
upper <- rep(1, d)

# Optimization 1: EHI with pso
optimcontrol <- list(method = "pso", maxit = 20)
critcontrol <- list(refPoint = c(1, 10))
omEG01 <- GParetoptim(model = model, fn = fname, crit = "EHI", nsteps = nsteps,
                    lower = lower, upper = upper, critcontrol = critcontrol,
                    optimcontrol = optimcontrol)

print(omEG01$par)
print(omEG01$values)

## Not run:
nsteps <- 10
# Optimization 2: SMS with discrete search
optimcontrol <- list(method = "discrete", candidate.points = test.grid)
critcontrol <- list(refPoint = c(1, 10))
omEG02 <- GParetoptim(model = model, fn = fname, crit = "SMS", nsteps = nsteps,
                    lower = lower, upper = upper, critcontrol = critcontrol,
                    optimcontrol = optimcontrol)

print(omEG02$par)
print(omEG02$values)

```



```

# Optimization 3: SUR with genoud
optimcontrol <- list(method = "genoud", pop.size = 20, max.generations = 10)
critcontrol <- list(distrib = "SUR", n.points = 100)
omEG03 <- GParetoptim(model = model, fn = fname, crit = "SUR", nsteps = nsteps,
                    lower = lower, upper = upper, critcontrol = critcontrol,
                    optimcontrol = optimcontrol)

print(omEG03$par)
print(omEG03$values)

# Optimization 4: EMI with pso
optimcontrol <- list(method = "pso", maxit = 20)
critcontrol <- list(nbsamp = 200)
omEG04 <- GParetoptim(model = model, fn = fname, crit = "EMI", nsteps = nsteps,
                    lower = lower, upper = upper, optimcontrol = optimcontrol)

print(omEG04$par)
print(omEG04$values)

# graphics
sol.grid <- apply(expand.grid(seq(0, 1, length.out = 100),
                             seq(0, 1, length.out = 100)), 1, fname)
plot(t(sol.grid), pch = 20, col = rgb(0, 0, 0, 0.05), xlim = c(0, 1),
     ylim = c(-2, 10), xlab = expression(f[1]), ylab = expression(f[2]))
plotGPareto(res = omEG01, add = TRUE,
            control = list(pch = 20, col = "blue", PF.pch = 17,
                          PF.points.col = "blue", PF.line.col = "blue"))
text(omEG01$values[,1], omEG01$values[,2], labels = 1:nsteps, pos = 3, col = "blue")
plotGPareto(res = omEG02, add = TRUE,
            control = list(pch = 20, col = "green", PF.pch = 17,
                          PF.points.col = "green", PF.line.col = "green"))
text(omEG02$values[,1], omEG02$values[,2], labels = 1:nsteps, pos = 3, col = "green")
plotGPareto(res = omEG03, add = TRUE,
            control = list(pch = 20, col = "red", PF.pch = 17,
                          PF.points.col = "red", PF.line.col = "red"))
text(omEG03$values[,1], omEG03$values[,2], labels = 1:nsteps, pos = 3, col = "red")
plotGPareto(res = omEG04, add = TRUE,
            control = list(pch = 20, col = "orange", PF.pch = 17,
                          PF.points.col = "orange", PF.line.col = "orange"))
text(omEG04$values[,1], omEG04$values[,2], labels = 1:nsteps, pos = 3, col = "orange")
points(response.grid[,1], response.grid[,2], col = "black", pch = 20)
legend("topright", c("EHI", "SMS", "SUR", "EMI"), col = c("blue", "green", "red", "orange"),
     pch = rep(17,4))

# Post-processing
plotGPareto(res = omEG01, UQ_PF = TRUE, UQ_PS = TRUE, UQ_dens = TRUE)

#####
# NOISY PROBLEMS
#####
set.seed(25468)
library(DiceDesign)
d <- 2
nsteps <- 3

```

```

lower <- rep(0, d)
upper <- rep(1, d)
optimcontrol <- list(method = "pso", maxit = 20)
critcontrol <- list(refPoint = c(1, 10))

n.grid <- 21
test.grid <- expand.grid(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid))
n.init <- 30
design <- maximinESE_LHS(lhsDesign(n.init, d, seed = 42)$design)$design

fit.models <- function(u) km(~., design = design, response = response[, u],
                           noise.var=design.noise.var[,u])

# Test 1: EHI, constant noise.var
noise.var <- c(0.1, 0.2)
funnoise1 <- function(x) {ZDT3(x) + sqrt(noise.var)*rnorm(n=d)}
response <- t(apply(design, 1, funnoise1))
design.noise.var <- matrix(rep(noise.var, n.init), ncol=d, byrow=TRUE)
model <- lapply(1:d, fit.models)

omEG01 <- GParetoptim(model = model, fn = funnoise1, crit = "EHI", nsteps = nsteps,
                    lower = lower, upper = upper, critcontrol = critcontrol,
                    reinterpolation=TRUE, noise.var=noise.var, optimcontrol = optimcontrol)
plotGPareto(omEG01)

# Test 2: EMI, noise.var given by fn
funnoise2 <- function(x) {list(ZDT3(x) + sqrt(0.05 + abs(0.1*x))*rnorm(n=d), 0.05 + abs(0.1*x))}
temp <- funnoise2(design)
response <- temp[[1]]
design.noise.var <- temp[[2]]
model <- lapply(1:d, fit.models)

omEG02 <- GParetoptim(model = model, fn = funnoise2, crit = "EMI", nsteps = nsteps,
                    lower = lower, upper = upper, critcontrol = critcontrol,
                    reinterpolation=TRUE, noise.var="given_by_fn", optimcontrol = optimcontrol)
plotGPareto(omEG02)

# Test 3: SMS, functional noise.var
funnoise3 <- function(x) {ZDT3(x) + sqrt(0.025 + abs(0.05*x))*rnorm(n=d)}
noise.var <- function(x) return(0.025 + abs(0.05*x))
response <- t(apply(design, 1, funnoise3))
design.noise.var <- t(apply(design, 1, noise.var))
model <- lapply(1:d, fit.models)

omEG03 <- GParetoptim(model = model, fn = funnoise3, crit = "SMS", nsteps = nsteps,
                    lower = lower, upper = upper, critcontrol = critcontrol,
                    reinterpolation=TRUE, noise.var=noise.var, optimcontrol = optimcontrol)
plotGPareto(omEG03)

# Test 4: SUR, fastfun, constant noise.var
noise.var <- 0.1
funnoise4 <- function(x) {ZDT3(x)[1] + sqrt(noise.var)*rnorm(1)}
cheapfn <- function(x) ZDT3(x)[2]

```

```

response <- apply(design, 1, funnoise4)
design.noise.var <- rep(noise.var, n.init)
model <- list(km(~., design = design, response = response, noise.var=design.noise.var))

omEG04 <- GParetoptim(model = model, fn = funnoise4, cheapfn = cheapfn, crit = "SUR",
  nsteps = nsteps, lower = lower, upper = upper, critcontrol = critcontrol,
  reinterpolation=TRUE, noise.var=noise.var, optimcontrol = optimcontrol)
plotGPareto(omEG04)

# Test 5: EMI, fastfun, noise.var given by fn
funnoise5 <- function(x) {
  if (is.null(dim(x))) x <- matrix(x, nrow=1)
  list(apply(x, 1, ZDT3)[1,] + sqrt(abs(0.05*x[,1]))*rnorm(nrow(x)), abs(0.05*x[,1]))
}

cheapfn <- function(x) {
  if (is.null(dim(x))) x <- matrix(x, nrow=1)
  apply(x, 1, ZDT3)[2,]
}

temp <- funnoise5(design)
response <- temp[[1]]
design.noise.var <- temp[[2]]
model <- list(km(~., design = design, response = response, noise.var=design.noise.var))

omEG05 <- GParetoptim(model = model, fn = funnoise5, cheapfn = cheapfn, crit = "EMI",
  nsteps = nsteps, lower = lower, upper = upper, critcontrol = critcontrol,
  reinterpolation=TRUE, noise.var="given_by_fn", optimcontrol = optimcontrol)
plotGPareto(omEG05)

# Test 6: EHI, fastfun, functional noise.var
noise.var <- 0.1
funnoise6 <- function(x) {ZDT3(x)[1] + sqrt(abs(0.1*x[1]))*rnorm(1)}
noise.var <- function(x) return(abs(0.1*x[1]))
cheapfn <- function(x) ZDT3(x)[2]
response <- apply(design, 1, funnoise6)
design.noise.var <- t(apply(design, 1, noise.var))
model <- list(km(~., design = design, response = response, noise.var=design.noise.var))

omEG06 <- GParetoptim(model = model, fn = funnoise6, cheapfn = cheapfn, crit = "EMI",
  nsteps = nsteps, lower = lower, upper = upper, critcontrol = critcontrol,
  reinterpolation=TRUE, noise.var=noise.var, optimcontrol = optimcontrol)
plotGPareto(omEG06)

## End(Not run)

```

integration_design_optim

Function to build integration points (for the SUR criterion)

Description

Modification of the function `integration_design` from the package `KrigInv` to be usable for SUR-based optimization. Handles two or three objectives. Available important sampling schemes: none so far.

Usage

```
integration_design_optim(SURcontrol = NULL, d = NULL, lower, upper,
  model = NULL, min.prob = 0.001)
```

Arguments

<code>SURcontrol</code>	Optional list specifying the procedure to build the integration points and weights. Many options are possible; see 'Details'.
<code>d</code>	The dimension of the input set. If not provided <code>d</code> is set equal to the length of <code>lower</code> .
<code>lower</code>	Vector containing the lower bounds of the design space.
<code>upper</code>	Vector containing the upper bounds of the design space.
<code>model</code>	A list of kriging models of <code>km</code> class.
<code>min.prob</code>	This argument applies only when importance sampling distributions are chosen. For numerical reasons we give a minimum probability for a point to belong to the importance sample. This avoids probabilities equal to zero and importance sampling weights equal to infinity. In an importance sample of <code>M</code> points, the maximum weight becomes $1/\text{min.prob} * 1/M$.

Details

The `SURcontrol` argument is a list with possible entries `integration.points`, `integration.weights`, `n.points`, `n.candidates`, `distrib`, `init.distrib` and `init.distrib.spec`. It can be used in one of the three following ways:

- A) If nothing is specified, $100 * d$ points are chosen using the Sobol sequence;
- B) One can directly set the field `integration.points` ($p * d$ matrix) for prespecified integration points. In this case these integration points and the corresponding vector `integration.weights` will be used for all the iterations of the algorithm;
- C) If the field `integration.points` is not set then the integration points are renewed at each iteration. In that case one can control the number of integration points `n.points` (default: $100*d$) and a specific distribution `distrib`. Possible values for `distrib` are: "sobol", "MC" and "SUR" (default: "sobol"):
 - C.1) The choice "sobol" corresponds to integration points chosen with the Sobol sequence in dimension `d` (uniform weight);
 - C.2) The choice "MC" corresponds to points chosen randomly, uniformly on the domain;
 - C.3) The choice "SUR" corresponds to importance sampling distributions (unequal weights). When important sampling procedures are chosen, `n.points` points are chosen using importance sampling among a discrete set of `n.candidates` points (default: `n.points*10`)

which are distributed according to a distribution `init.distrib` (default: "sobol"). Possible values for `init.distrib` are the space filling distributions "sobol" and "MC" or an user defined distribution "spec". The "sobol" and "MC" choices correspond to quasi random and random points in the domain. If the "spec" value is chosen the user must fill in manually the field `init.distrib.spec` to specify himself a `n.candidates * d` matrix of points in dimension `d`.

Value

A list with components:

- `integration.points` `p * d` matrix of `p` points used for the numerical calculation of integrals
- `integration.weights` a vector of size `p` corresponding to the weight of each point. If all the points are equally weighted, `integration.weights` is set to NULL

References

V. Picheny (2014), Multiobjective optimization using Gaussian process emulators via stepwise uncertainty reduction, *Statistics and Computing*.

See Also

[GParetooptim crit_SUR integration_design](#)

nonDomSet

Non-dominated points with respect to a reference

Description

Determines which elements in a set are dominated by reference points

Usage

```
nonDomSet(points, ref)
```

Arguments

<code>points</code>	matrix (one point per row) that are compared to a reference <code>ref</code> (i.e., not between themselves)
<code>ref</code>	matrix (one point per row) of reference (faster if they are already Pareto optimal)

Examples

```
## Not run:
d <- 6
n <- 1000
n2 <- 1000

test <- matrix(runif(d * n), n)
ref <- matrix(runif(d * n), n)
indPF <- which(!is_dominated(t(ref)))

system.time(res <- nonDomSet(test, ref[indPF,,drop = FALSE]))

res2 <- rep(NA, n2)
library(emoa)
t0 <- Sys.time()
for(i in 1:n2){
  res2[i] <- !is_dominated(t(rbind(test[i,, drop = FALSE], ref[indPF,])))
}
print(Sys.time() - t0)

all(res == res2)

## End(Not run)
```

ParetoSetDensity

Estimation of Pareto set density

Description

Estimation of the density of Pareto optimal points in the variable space.

Usage

```
ParetoSetDensity(model, lower, upper, CPS = NULL, nsim = 50,
  simpoints = 1000, ...)
```

Arguments

model	list of objects of class <code>km</code> , one for each objective functions,
lower	vector of lower bounds for the variables,
upper	vector of upper bounds for the variables,
CPS	optional matrix containing points from Conditional Pareto Set Simulations (in the variable space), see details
nsim	optional number of conditional simulations to perform if CPS is not provided,
simpoints	(optional) If CPS is NULL, either a number of simulation points, or a matrix where conditional simulations are to be performed. In the first case, then simulation points are taken as a maximin LHS design using <code>lhsDesign</code> .

... further arguments to be passed to `kde`. In particular, if the input dimension is greater than three, a matrix `eval.points` can be given (else it is taken as the simulation points).

Details

This function estimates the density of Pareto optimal points in the variable space given by the surrogate models. Based on conditional simulations of the objectives at simulation points, Conditional Pareto Set (CPS) simulations are obtained, out of which a density is fitted.

This function relies on the `ks` package for the kernel density estimation.

Value

An object of class `kde` accounting for the estimated density of Pareto optimal points.

Examples

```
## Not run:
#-----
# Example of estimation of the density of Pareto optimal points
#-----
set.seed(42)
n_var <- 2
fname <- P1
lower <- rep(0, n_var)
upper <- rep(1, n_var)

res1 <- easyGParetooptim(fn = fname, lower = lower, upper = upper, budget = 15,
control=list(method = "EHI", inneroptim = "pso", maxit = 20))

estDens <- ParetoSetDensity(res1$model, lower = lower, upper = upper)

# graphics
par(mfrow = c(1,2))
plot(estDens, display = "persp", xlab = "X1", ylab = "X2")
plot(estDens, display = "filled.contour2", main = "Estimated density of Pareto optimal point")
points(res1$model[[1]]@X[,1], res1$model[[2]]@X[,2], col="blue")
points(estDens$x[, 1], estDens$x[, 2], pch = 20, col = rgb(0, 0, 0, 0.15))
par(mfrow = c(1,1))

## End(Not run)
```

plotGPareto

Plot multi-objective optimization results and post-processing

Description

Display results of multi-objective optimization returned by either `GParetooptim` or `easyGParetooptim`, possibly completed with various post-processings of uncertainty quantification.

Usage

```
plotGPareto(res, add = FALSE, UQ_PF = FALSE, UQ_PS = FALSE,
  UQ_dens = FALSE, lower = NULL, upper = NULL, control = list(pch =
  20, col = "red", PF.line.col = "cyan", PF.pch = 17, PF.points.col =
  "blue", VE.line.col = "cyan", nsim = 100, npsim = 1500, gridtype =
  "runif", displaytype = "persp", printVD = TRUE, use.rgl = TRUE, bounds =
  NULL, meshsize3d = 50, theta = -25, phi = 10, add_denoised_PF = TRUE))
```

Arguments

res	list returned by GParetooptim or easyGParetooptim ,
add	logical; if TRUE adds the first graphical output to an already existing plot; if FALSE, (default) starts a new plot,
UQ_PF	logical; for 2 objectives, if TRUE perform a quantification of uncertainty on the Pareto front to display the symmetric deviation function with plotSymDevFun (cannot be added to existing graph),
UQ_PS	logical; if TRUE call plot_uncertainty representing the probability of non-domination in the variable space,
UQ_dens	logical; for 2D problems, if TRUE call ParetoSetDensity to estimate and display the density of Pareto optimal points in the variable space,
lower	optional vector of lower bounds for the variables. Necessary if UQ_PF and/or UQ_PS are TRUE (if not provided, variables are supposed to vary between 0 and 1),
upper	optional vector of upper bounds for the variables. Necessary if UQ_PF and/or UQ_PS are TRUE (if not provided, variables are supposed to vary between 0 and 1),
control	optional list, see details.

Details

By default, `plotGPareto` displays the Pareto front delimiting the non-dominated area with 2 objectives, by a perspective view with 3 objectives and using parallel coordinates with more objectives.

Setting one or several of `UQ_PF`, `UQ_PS` and `UQ_dens` allows to run and display post-processing tools that assess the precision and confidence of the optimization run, either in the objective (`UQ_PF`) or the variable spaces (`UQ_PS`, `UQ_dens`). Note that these options are computationally intensive.

Various parameters can be used for the display of results and/or passed to subsequent function:

- `col`, `pch` correspond the color and plotting character for observations,
- `PF.line.col`, `PF.pch`, `PF.points.col` define the color of the line denoting the current Pareto front, the plotting character and color of non-dominated observations, respectively,
- `nsim`, `npsim` and `gridtype` define the number of conditional simulations performed with [simulate](#) along with the number of simulation points (in case `UQ_PF` and/or `UQ_dens` are TRUE),

- gridtype to define how simulation points are selected; alternatives are 'runif' (default) for uniformly sampled points, 'LHS' for a Latin Hypercube design using [lhsDesign](#) and 'grid2d' for a two dimensional grid,
- f1lim, f2lim can be passed to [CPF](#),
- resolution, option, nintegpoints are to be passed to [plot_uncertainty](#)
- displaytype type of display for UQ_dens, see [plot.kde](#),
- printVD logical, if TRUE and UQ_PF is TRUE as well, print the value of the Vorob'ev deviation,
- use.rgl if TRUE, use rgl for 3D plots, else [persp](#) is used,
- bounds if use.rgl is TRUE, optional 2*nobj matrix of boundaries, see [plotParetoEmp](#)
- meshsize3d mesh size of the perspective view for 3-objective problems,
- theta, phi angles for perspective view of 3-objective problems,
- add_denoised_PF if TRUE, in the noisy case, add the Pareto front from the estimated mean of the observations.

References

M. Binois, D. Ginsbourger and O. Roustant (2015), Quantifying Uncertainty on Pareto Fronts with Gaussian process conditional simulations, *European Journal of Operational Research*, 243(2), 386-394.

A. Inselberg (2009), *Parallel coordinates*, Springer.

Examples

```
## Not run:
#-----
# 2D objective function
#-----
set.seed(25468)
n_var <- 2
fname <- P1
lower <- rep(0, n_var)
upper <- rep(1, n_var)
res <- easyGParetooptim(fn=fname, lower=lower, upper=upper, budget=15,
control=list(method="EHI", inneroptim="pso", maxit=20))

## Pareto front only
plotGPareto(res)

## With post-processing
plotGPareto(res, UQ_PF = TRUE, UQ_PS = TRUE, UQ_dens = TRUE)

## With noise
noise.var <- c(10, 2)
funnoise <- function(x) {P1(x) + sqrt(noise.var)*rnorm(n=2)}
res2 <- easyGParetooptim(fn=funnoise, lower=lower, upper=upper, budget=15, noise.var=noise.var,
control=list(method="EHI", inneroptim="pso", maxit=20))
```

```

plotGPareto(res2, control=list(add_denoised_PF=FALSE)) # noisy observations only
plotGPareto(res2)

#-----
# 3D objective function
#-----
set.seed(1)
n_var <- 3
fname <- DTLZ1
lower <- rep(0, n_var)
upper <- rep(1, n_var)
res3 <- easyGParetooptim(fn=fname, lower=lower, upper=upper, budget=50,
control=list(method="EHI", inneroptim="pso", maxit=20))

## Pareto front only
plotGPareto(res3)

## With noise
noise.var <- c(10, 2, 5)
funnoise <- function(x) {fname(x) + sqrt(noise.var)*rnorm(n=3)}
res4 <- easyGParetooptim(fn=funnoise, lower=lower, upper=upper, budget=100, noise.var=noise.var,
control=list(method="EHI", inneroptim="pso", maxit=20))

plotGPareto(res4, control=list(add_denoised_PF=FALSE)) # noisy observations only
plotGPareto(res4)

## End(Not run)

```

plotParetoEmp

Pareto front visualization

Description

Plot the Pareto front with step functions.

Usage

```

plotParetoEmp(nondominatedPoints, add = TRUE, max = FALSE,
bounds = NULL, alpha = 0.5, ...)

```

Arguments

nondominatedPoints	points considered to plot the Pareto front with segments, matrix with one point per row,
add	optional boolean indicating whether a new graphic should be drawn,
max	optional boolean indicating whether to display a Pareto front in a maximization context,
bounds	for 3D, optional 2*nobj matrix of boundaries

alpha for 3D, optional value in [0,1] for transparency
 ... additional values to be passed to the `lines` function.

Examples

```
#-----
# Simple example
#-----

x <- c(0.2, 0.4, 0.6, 0.8)
y <- c(0.8, 0.7, 0.5, 0.1)

plot(x, y, col = "green", pch = 20)

plotParetoEmp(cbind(x, y), col = "green")
## Alternative
plotParetoEmp(cbind(x, y), col = "red", add = FALSE)

## With maximization
plotParetoEmp(cbind(x, y), col = "blue", max = TRUE)

## 3D plots
library(rgl)
set.seed(5)
X <- matrix(runif(60), ncol=3)
Xnd <- t(nondominated_points(t(X)))
plot3d(X)
plot3d(Xnd, col="red", size=8, add=TRUE)
plot3d(x=min(Xnd[,1]), y=min(Xnd[,2]), z=min(Xnd[,3]), col="green", size=8, add=TRUE)
X.range <- diff(apply(X,2,range))
bounds <- rbind(apply(X,2,min)-0.1*X.range,apply(X,2,max)+0.1*X.range)
plotParetoEmp(nondominatedPoints = Xnd, add=TRUE, bounds=bounds, alpha=0.5)
```

plotParetoGrid

Visualisation of Pareto front and set

Description

Plot the Pareto front and set for 2 variables 2 objectives test problems with evaluations on a grid.

Usage

```
plotParetoGrid(fname = "ZDT1", xlim = c(0, 1), ylim = c(0, 1),
  n.grid = 100)
```

Arguments

fname	name of the function considered,
xlim, ylim	numeric vectors of length 2, giving the x and y coordinates ranges, default is $[0,1] \times [0,1]$,
n.grid	number of divisions of the grid in each dimension.

Examples

```
#-----
# Examples with test functions
#-----

plotParetoGrid("ZDT3", n.grid = 21)

plotParetoGrid("P1", n.grid = 21)

plotParetoGrid("MOP2", xlim = c(0, 1), ylim = c(0, 1), n.grid = 21)
```

plotSymDevFun	<i>Display the Symmetric Deviation Function</i>
---------------	---

Description

Display the Symmetric Deviation Function for an object of class CPF.

Usage

```
plotSymDevFun(CPF, n.grid = 100)
```

Arguments

CPF	CPF object, see CPF ,
n.grid	number of divisions of the grid in each dimension.

Details

Display observations in red and the corresponding Pareto front by a step-line. The blue line is the estimation of the location of the Pareto front of the kriging models, named Vorob'ev expectation. In grayscale is the intensity of the deviation (symmetrical difference) from the Vorob'ev expectation for couples of conditional simulations.

References

M. Binois, D. Ginsbourger and O. Roustant (2015), Quantifying Uncertainty on Pareto Fronts with Gaussian process conditional simulations, *European Journal of Operational Research*, 243(2), 386-394.

Examples

```

library(DiceDesign)
set.seed(42)

nvar <- 2

# Test function
fname = "P1"

# Initial design
nappr <- 10
design.grid <- maximinESE_LHS(lhsDesign(nappr, nvar, seed = 42)$design)$design
response.grid <- t(apply(design.grid, 1, fname))

ParetoFront <- t(nondominated_points(t(response.grid)))

# kriging models : matern5_2 covariance structure, linear trend, no nugget effect
mf1 <- km(~., design = design.grid, response = response.grid[, 1])
mf2 <- km(~., design = design.grid, response = response.grid[, 2])

# Conditional simulations generation with random sampling points
nsim <- 10 # increase for better results
npointssim <- 80 # increase for better results
Simu_f1 = matrix(0, nrow = nsim, ncol = npointssim)
Simu_f2 = matrix(0, nrow = nsim, ncol = npointssim)
design.sim = array(0,dim = c(npointssim, nvar, nsim))

for(i in 1:nsim){
  design.sim[, , i] <- matrix(runif(nvar*npointssim), npointssim, nvar)
  Simu_f1[i,] = simulate(mf1, nsim = 1, newdata = design.sim[, , i], cond = TRUE,
                        checkNames = FALSE, nugget.sim = 10^-8)
  Simu_f2[i,] = simulate(mf2, nsim = 1, newdata = design.sim[, , i], cond=TRUE,
                        checkNames = FALSE, nugget.sim = 10^-8)
}

# Attainment, Voreb'ev expectation and deviation estimation
CPF1 <- CPF(Simu_f1, Simu_f2, response.grid, ParetoFront)

# Symmetric deviation function
plotSymDevFun(CPF1)

```

plotSymDifRNP

Symmetrical difference of RNP sets

Description

Plot the symmetrical difference between two Random Non-Dominated Point (RNP) sets.

Usage

```
plotSymDifRNP(set1, set2, xlim, ylim, fill = "black", add = "FALSE",
  ...)
```

Arguments

set1, set2	RNP sets considered,
xlim, ylim	numeric vectors of length 2, giving the x and y coordinates ranges for plotting,
fill	optional color of the symmetric difference area,
add	logical; if TRUE add to an already existing plot; if FALSE (default) start a new plot taking xlim, ylim as limits.
...	additional parameters for the <code>plot</code> and <code>polygon</code> graphic functions

Examples

```
#-----
# Simple example
#-----
set1 <- rbind(c(0.2, 0.35, 0.5, 0.8),
             c(0.8, 0.6, 0.55, 0.3))

set2 <- rbind(c(0.3, 0.4),
             c(0.7, 0.4))

plotSymDifRNP(set1, set2, xlim = c(0, 1), ylim = c(0, 1), fill = "grey")
points(t(set1), col = "red", pch = 20)
points(t(set2), col = "blue", pch = 20)
```

plot_uncertainty	<i>Plot uncertainty</i>
------------------	-------------------------

Description

Displays the probability of non-domination in the variable space. In dimension larger than two, projections in 2D subspaces are displayed.

Usage

```
plot_uncertainty(model, paretoFront = NULL, type = "pn", lower, upper,
  resolution = 51, option = "mean", nintegpoints = 400)
```

Arguments

model	list of objects of class <code>km</code> , one for each objective functions,
paretoFront	(optional) matrix corresponding to the Pareto front of size $[n.pareto \times n.obj]$,
type	type of uncertainty, for now only the probability of improvement over the Pareto front,
lower	vector of lower bounds for the variables,
upper	vector of upper bounds for the variables,
resolution	grid size (the total number of points is $resolution^d$),
option	optional argument (string) for $n > 2$ variables to define the projection type. The 3 possible values are "mean" (default), "max" and "min",
nintegpoints	number of integration points for computation of mean, max and min values.

Details

Function inspired by the function `print_uncertainty` and `print_uncertainty_nd` from the package `KrigInv`. Non-dominated observations are represented with green diamonds, dominated ones by yellow triangles.

Examples

```
## Not run:
#-----
# 2D, bi-objective function
#-----
set.seed(25468)
n_var <- 2
fname <- P1
lower <- rep(0, n_var)
upper <- rep(1, n_var)
res1 <- easyGParetooptim(fn=fname, lower=lower, upper=upper, budget=15,
control=list(method="EHI", inneroptim="pso", maxit=20))

plot_uncertainty(res1$model, lower = lower, upper = upper)

#-----
# 4D, bi-objective function
#-----
set.seed(25468)
n_var <- 4
fname <- DTLZ2
lower <- rep(0, n_var)
upper <- rep(1, n_var)
res <- easyGParetooptim(fn=fname, lower=lower, upper=upper, budget = 40,
control=list(method="EHI", inneroptim="pso", maxit=40))

plot_uncertainty(res$model, lower = lower, upper = upper, resolution = 31)

## End(Not run)
```

predict_kms	<i>Predict function for list of km models.</i>
-------------	--

Description

Predict function for list of [km](#) models.

Usage

```
predict_kms(model, newdata, type, se.compute = TRUE,  
            cov.compute = FALSE, light.return = FALSE, bias.correct = FALSE,  
            checkNames = TRUE, ...)
```

Arguments

model list of [km](#) models
newdata, type, se.compute, cov.compute, light.return, bias.correct, checkNames, ...
 see [predict.km](#)

Details

So far only `light.return = TRUE` and `cov.compute = FALSE` handled.

ZDT1	<i>Test functions of x</i>
------	---

Description

Multi-objective test functions.

Usage

ZDT1(x)

ZDT2(x)

ZDT3(x)

ZDT4(x)

ZDT6(x)

P1(x)

P2(x)

MOP2(x)

MOP3(x)

DTLZ1(x, nobj = 3)

DTLZ2(x, nobj = 3)

DTLZ3(x, nobj = 3)

DTLZ7(x, nobj = 3)

Arguments

x matrix specifying the location where the function is to be evaluated, one point per row,
 nobj optional argument to select the number of objective for the DTLZ test functions.

Details

These functions are coming from different benchmarks: the ZDT test problems from an article of E. Zitzler et al., P1 from the thesis of J. Parr and P2 from an article of Poloni et al. . MOP2 and MOP3 are from Van Veldhuizen and DTLZ functions are from Deb et al. .

Domains (sometimes rescaled to $[0, 1]$):

- ZDT1-6: $[0, 1]^d$
- P1, P2: $[0, 1]^2$
- MOP2: $[0, 1]^d$
- MOP3: $[-3, 3]$, tri-objective, 2 variables
- DTLZ1-3, 7: $[0, 1]^d$, m-objective problems, with at least $d > m$ variables.

Value

Matrix of values corresponding to the objective functions, the number of columns is the number of objectives.

References

- J. M. Parr (2012), *Improvement Criteria for Constraint Handling and Multiobjective Optimization*, University of Southampton, PhD thesis.
- C. Poloni, A. Giurgevich, L. Onesti, V. Pediroda (2000), Hybridization of a multi-objective genetic algorithm, a neural network and a classical optimizer for a complex design problem in fluid dynamics, *Computer Methods in Applied Mechanics and Engineering*, 186(2), 403-420.
- E. Zitzler, K. Deb, and L. Thiele (2000), Comparison of multiobjective evolutionary algorithms: Empirical results, *Evol. Comput.*, 8(2), 173-195.

K. Deb, L. Thiele, M. Laumanns and E. Zitzler (2002), Scalable Test Problems for Evolutionary Multiobjective Optimization, *IEEE Transactions on Evolutionary Computation*, 6(2), 182-197.

D. A. Van Veldhuizen, G. B. Lamont (1999), Multiobjective evolutionary algorithm test suites, *In Proceedings of the 1999 ACM symposium on Applied computing*, 351-357.

Examples

```
# -----  
# 2-objectives test problems  
# -----  
  
plotParetoGrid("ZDT1", n.grid = 21)  
  
plotParetoGrid("ZDT2", n.grid = 21)  
  
plotParetoGrid("ZDT3", n.grid = 21)  
  
plotParetoGrid("ZDT4", n.grid = 21)  
  
plotParetoGrid("ZDT6", n.grid = 21)  
  
plotParetoGrid("P1", n.grid = 21)  
  
plotParetoGrid("P2", n.grid = 21)  
  
plotParetoGrid("MOP2", n.grid = 21)
```

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