

# Package ‘DiceOptim’

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**Version** 2.1.1

**Title** Kriging-Based Optimization for Computer Experiments

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**Description** Efficient Global Optimization (EGO) algorithm as described in “Roustant et al. (2012)” <doi:10.18637/jss.v051.i01> and adaptations for problems with noise (“Picheny and Ginsbourger, 2012”) <doi:10.1016/j.csda.2013.03.018>, parallel infill, and problems with constraints.

**Depends** DiceKriging (>= 1.2), methods

**Imports** randtoolbox, pbivnorm, rgenoud, mnormt, DiceDesign, parallel

**Suggests** KrigInv, GPareto

**License** GPL-2 | GPL-3

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DiceOptim-package

*Kriging-based optimization methods for computer experiments***Description**

Sequential and parallel Kriging-based optimization methods relying on expected improvement criteria.

**Details**

Package: DiceOptim  
 Type: Package  
 Version: 2.0  
 Date: July 2016  
 License: GPL-2 | GPL-3

## Note

This work is a follow-up of DiceOptim 1.0, which was produced within the frame of the DICE (Deep Inside Computer Experiments) Consortium between ARMINES, Renault, EDF, IRSN, ON-ERA and TOTAL S.A.

The authors would like to thank Yves Deville for his precious advice in R programming and packaging, as well as the DICE members for useful feedbacks, and especially Yann Richet (IRSN) for numerous discussions concerning the user-friendliness of this package.

Package `rngenoud`  $\geq 5.3.3$ . is recommended.

Important functions or methods:

<code>EGO.nsteps</code>	Standard Efficient Global Optimization algorithm with a fixed number of iterations (nsteps) —with model updates including re-estimation of covariance hyperparameters
<code>EI</code>	Expected Improvement criterion (single infill point, noise-free, constraint free problems)
<code>max_EI</code>	Maximization of the EI criterion. No need to specify any objective function
<code>qEI.nsteps</code>	EGO algorithm with batch-sequential (parallel) infill strategy
<code>noisy.optimizer</code>	EGO algorithm for noisy objective functions
<code>EGO.cst</code>	EGO algorithm for (non-linear) constrained problems
<code>easyEGO.cst</code>	User-friendly wrapper for <code>EGO.cst</code>

## Author(s)

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with contributions by M. Binois, C. Chevalier, S. Marmin and T. Wagner

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T.J. Santner, B.J. Williams, and W.J. Notz (2003), *The design and analysis of computer experiments*, Springer.

M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

## Examples

```
set.seed(123)

#####
### 2D optimization USING EGO.nsteps and qEGO.nsteps #####
#####

# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- data.frame(apply(design.fact, 1, branin))
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

### EGO, 5 steps #####
library(rgenoud)
```

```

nsteps <- 5
lower <- rep(0,d)
upper <- rep(1,d)
oEGO <- EGO.nsteps(model=fitted.model1, fun=branin, nsteps=nsteps,
lower=lower, upper=upper, control=list(pop.size=20, BFGSburnin=2))
print(oEGO$par)
print(oEGO$value)

# graphics
n.grid <- 15
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, branin)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid, y.grid, z.grid, 40)
title("EGO")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par, pch=19, col="red")
text(oEGO$par[,1], oEGO$par[,2], labels=1:nsteps, pos=3)

### Parallel EGO, 3 steps with batches of 3 #####
nsteps <- 3
lower <- rep(0,d)
upper <- rep(1,d)
npoints <- 3 # The batchsize
oEGO <- qEGO.nsteps(model = fitted.model1, branin, npoints = npoints, nsteps = nsteps,
crit="exact", lower, upper, optimcontrol = NULL)
print(oEGO$par)
print(oEGO$value)

# graphics
contour(x.grid, y.grid, z.grid, 40)
title("qEGO")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par, pch=19, col="red")
text(oEGO$par[,1], oEGO$par[,2], labels=c(tcrossprod(rep(1,npoints),1:nsteps)), pos=3)

#####
### 2D OPTIMIZATION, NOISY OBJECTIVE ###
#####

set.seed(10)
library(DiceDesign)
# Set test problem parameters
doe.size <- 9
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.1

# Build noisy simulator
funnoise <- function(x)

```

```

{      f.new <- test.function(x) + sqrt(noise.var)*rnorm(n=1)
      return(f.new)}

# Generate DOE and response
doe <- as.data.frame(lhsDesign(doe.size, dim)$design)
y.tilde <- funnoise(doe)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Optimisation with noisy.optimizer
optim.param <- list()
optim.param$quantile <- .7
optim.result <- noisy.optimizer(optim.crit="EQI", optim.param=optim.param, model=model,
                               n.ite=5, noise.var=noise.var, funnoise=funnoise, lower=lower, upper=upper,
                               NoiseReEstimate=FALSE, CovReEstimate=FALSE)

print(optim.result$best.x)

#####
### 2D OPTIMIZATION, 2 INEQUALITY CONSTRAINTS                                     ###
#####
set.seed(25468)
library(DiceDesign)

fun <- goldsteinprice
fun1.cst <- function(x){return(-branin(x) + 25)}
fun2.cst <- function(x){return(3/2 - x[1] - 2*x[2] - .5*sin(2*pi*(x[1]^2 - 2*x[2])))}
constraint <- function(x){return(c(fun1.cst(x), fun2.cst(x)))}

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

## Optimization using the Expected Feasible Improvement criterion
res <- easyEG0.cst(fun=fun, constraint=constraint, n.cst=2, lower=lower, upper=upper, budget=10,
                  control=list(method="EFI", inneroptim="genoud", maxit=20))

cat("best design found:", res$par, "\n")
cat("corresponding objective and constraints:", res$value, "\n")

# Objective function in colour, constraint boundaries in red
# Initial DoE: white circles, added points: blue crosses, best solution: red cross

n.grid <- 15
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun)
cst1.grid <- apply(test.grid, 1, fun1.cst)
cst2.grid <- apply(test.grid, 1, fun2.cst)
filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
               matrix(obj.grid, n.grid), main = "Two inequality constraints",
               xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,

```

```

    plot.axes = {axis(1); axis(2);
      contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
        matrix(cst1.grid, n.grid), level = 0, add=TRUE,
        drawlabels=FALSE, lwd=1.5, col = "red")
      contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
        matrix(cst2.grid, n.grid), level = 0, add=TRUE, drawlabels=FALSE,
        lwd=1.5, col = "red")
      points(res$history$X, col = "blue", pch = 4, lwd = 2)
      points(res$par[1], res$par[2], col = "red", pch = 4, lwd = 2, cex=2)
    }
  )

```

---

 AEI

---

*Augmented Expected Improvement*


---

### Description

Evaluation of the Augmented Expected Improvement (AEI) criterion, which is a modification of the classical EI criterion for noisy functions. The AEI consists of the regular EI multiplied by a penalization function that accounts for the diminishing payoff of observation replicates. The current minimum  $y_{\min}$  is chosen as the kriging predictor of the observation with smallest kriging quantile.

### Usage

```
AEI(x, model, new.noise.var = 0, y.min = NULL, type = "UK", envir = NULL)
```

### Arguments

<code>x</code>	the input vector at which one wants to evaluate the criterion
<code>model</code>	a Kriging model of "km" class
<code>new.noise.var</code>	the (scalar) noise variance of the future observation.
<code>y.min</code>	The kriging predictor at the current best point (point with smallest kriging quantile). If not provided, this quantity is evaluated.
<code>type</code>	Kriging type: "SK" or "UK"
<code>envir</code>	environment for saving intermediate calculations and reusing them within AEI.grad

### Value

Augmented Expected Improvement

### Author(s)

Victor Picheny  
David Ginsbourger



## References

D. Huang, T.T. Allen, W.I. Notz, and N. Zeng (2006), Global Optimization of Stochastic Black-Box Systems via Sequential Kriging Meta-Models, *Journal of Global Optimization*, 34, 441-466.

## Examples

```
#####
### AEI SURFACE ASSOCIATED WITH AN ORDINARY KRIGING MODEL ###
### OF THE BRANIN FUNCTION KNOWN AT A 12-POINT LATIN HYPERCUBE DESIGN ###
#####

set.seed(421)

# Set test problem parameters
doe.size <- 12
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {
y.tilde[i] <- test.function(doe[i,]) + sqrt(noise.var)*rnorm(n=1)
}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Compute actual function and criterion on a grid
n.grid <- 12 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
nt <- nrow(design.grid)
crit.grid <- rep(0,1,nt)
func.grid <- rep(0,1,nt)

crit.grid <- apply(design.grid, 1, AEI, model=model, new.noise.var=noise.var)
func.grid <- apply(design.grid, 1, test.function)

# Compute kriging mean and variance on a grid
names(design.grid) <- c("V1","V2")
pred <- predict.km(model, newdata=design.grid, type="UK")
mk.grid <- pred$m
sk.grid <- pred$sd
```

```

# Plot actual function
z.grid <- matrix(func.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Actual function");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot Kriging mean
z.grid <- matrix(mk.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Kriging mean");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot Kriging variance
z.grid <- matrix(sk.grid^2, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Kriging variance");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot AEI criterion
z.grid <- matrix(crit.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("AEI");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

```

---

AEI.grad

*AEI's Gradient*


---

## Description

Analytical gradient of the Augmented Expected Improvement (AEI) criterion.

## Usage

```
AEI.grad(x, model, new.noise.var = 0, y.min = NULL, type = "UK", envir = NULL)
```

## Arguments

<code>x</code>	the input vector at which one wants to evaluate the criterion
<code>model</code>	a Kriging model of "km" class
<code>new.noise.var</code>	the (scalar) noise variance of the new observation.
<code>y.min</code>	The kriging predictor at the current best point (point with smallest kriging quantile). If not provided, this quantity is evaluated.
<code>type</code>	Kriging type: "SK" or "UK"
<code>envir</code>	environment for inheriting intermediate calculations from AEI

**Value**

Gradient of the Augmented Expected Improvement

**Author(s)**

Victor Picheny

David Ginsbourger

**Examples**

```

set.seed(421)

# Set test problem parameters
doe.size <- 12
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {
  y.tilde[i] <- test.function(doe[i,]) + sqrt(noise.var)*rnorm(n=1)
}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Compute actual function and criterion on a grid
n.grid <- 8 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
nt <- nrow(design.grid)

crit.grid <- apply(design.grid, 1, AEI, model=model, new.noise.var=noise.var)
crit.grad <- t(apply(design.grid, 1, AEI.grad, model=model, new.noise.var=noise.var))

z.grid <- matrix(crit.grid, n.grid, n.grid)
contour(x.grid,y.grid, z.grid, 30)
title("AEI and its gradient")
points(model@X[,1],model@X[,2],pch=17,col="blue")

for (i in 1:nt)
{
  x <- design.grid[i,]
  suppressWarnings(arrows(x$Var1,x$Var2, x$Var1+crit.grad[i,1]*.6,x$Var2+crit.grad[i,2]*.6,

```

```
length=0.04,code=2,col="orange",lwd=2))  
}
```

---

AKG

*Approximate Knowledge Gradient (AKG)*

---

### Description

Evaluation of the Approximate Knowledge Gradient (AKG) criterion.

### Usage

```
AKG(x, model, new.noise.var = 0, type = "UK", envir = NULL)
```

### Arguments

x	the input vector at which one wants to evaluate the criterion
model	a Kriging model of "km" class
new.noise.var	(scalar) noise variance of the future observation. Default value is 0 (noise-free observation).
type	Kriging type: "SK" or "UK"
envir	environment for saving intermediate calculations and reusing them within AKG.grad

### Value

Approximate Knowledge Gradient

### Author(s)

Victor Picheny

David Ginsbourger

### References

Scott, W., Frazier, P., Powell, W. (2011). The correlated knowledge gradient for simulation optimization of continuous parameters using gaussian process regression. *SIAM Journal on Optimization*, 21(3), 996-1026.

## Examples

```
#####
###   AKG SURFACE ASSOCIATED WITH AN ORDINARY KRIGING MODEL       ###
### OF THE BRANIN FUNCTION KNOWN AT A 12-POINT LATIN HYPERCUBE DESIGN ###
#####
set.seed(421)
# Set test problem parameters
doe.size <- 12
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {
  y.tilde[i] <- test.function(doe[i,]) + sqrt(noise.var)*rnorm(n=1)
}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
            covtype="gauss", noise.var=rep(noise.var,1,doe.size),
            lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Compute actual function and criterion on a grid
n.grid <- 12 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
nt <- nrow(design.grid)

crit.grid <- apply(design.grid, 1, AKG, model=model, new.noise.var=noise.var)
func.grid <- apply(design.grid, 1, test.function)

# Compute kriging mean and variance on a grid
names(design.grid) <- c("V1","V2")
pred <- predict.km(model, newdata=design.grid, type="UK")
mk.grid <- pred$m
sk.grid <- pred$sd

# Plot actual function
z.grid <- matrix(func.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
              plot.axes = {title("Actual function");
                           points(model@X[,1],model@X[,2],pch=17,col="blue");
                           axis(1); axis(2)}})

# Plot Kriging mean
z.grid <- matrix(mk.grid, n.grid, n.grid)
```

```

filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
              plot.axes = {title("Kriging mean");
                          points(model@X[,1],model@X[,2],pch=17,col="blue");
                          axis(1); axis(2)})

# Plot Kriging variance
z.grid <- matrix(sk.grid^2, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
              plot.axes = {title("Kriging variance");
                          points(model@X[,1],model@X[,2],pch=17,col="blue");
                          axis(1); axis(2)})

# Plot AKG criterion
z.grid <- matrix(crit.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
              plot.axes = {title("AKG");
                          points(model@X[,1],model@X[,2],pch=17,col="blue");
                          axis(1); axis(2)})

```

---

AKG.grad

*AKG's Gradient*


---

### Description

Gradient of the Approximate Knowledge Gradient (AKG) criterion.

### Usage

```
AKG.grad(x, model, new.noise.var = 0, type = "UK", envir = NULL)
```

### Arguments

x	the input vector at which one wants to evaluate the criterion
model	a Kriging model of "km" class
new.noise.var	(scalar) noise variance of the future observation. Default value is 0 (noise-free observation).
type	Kriging type: "SK" or "UK"
envir	optional: environment for reusing intermediate calculations from AKG

### Value

Gradient of the Approximate Knowledge Gradient

### Author(s)

Victor Picheny

## Examples

```
#####
###   AKG SURFACE AND ITS GRADIENT ASSOCIATED WITH AN ORDINARY   #####
###                                     KRIGING MODEL             #####
### OF THE BRANIN FUNCTION KNOWN AT A 12-POINT LATIN HYPERCUBE DESIGN #####
#####
set.seed(421)

# Set test problem parameters
doe.size <- 12
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {
y.tilde[i] <- test.function(doe[i,]) + sqrt(noise.var)*rnorm(n=1)
}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Compute actual function and criterion on a grid
n.grid <- 9 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
nt <- nrow(design.grid)

crit.grid <- apply(design.grid, 1, AKG, model=model, new.noise.var=noise.var)
crit.grad <- t(apply(design.grid, 1, AKG.grad, model=model, new.noise.var=noise.var))

z.grid <- matrix(crit.grid, n.grid, n.grid)
contour(x.grid,y.grid, z.grid, 30)
title("AKG and its gradient")
points(model@X[,1],model@X[,2],pch=17,col="blue")

for (i in 1:nt)
{
  x <- design.grid[i,]
  suppressWarnings(arrows(x$Var1,x$Var2, x$Var1+crit.grad[i,1]*.2,x$Var2+crit.grad[i,2]*.2,
length=0.04,code=2,col="orange",lwd=2))
}

```

---

 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 = "covdist", type = "UK")
```

### Arguments

x	a vector representing the input to check,
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", "covdist" (default) 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 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.

### Value

TRUE if the point should not be tested.

### Author(s)

Mickael Binois



---

critcst\_optimizer      *Maximization of constrained Expected Improvement criteria*

---

## Description

Given objects of class `km` for the objective and constraints, and a set of tuning parameters (`lower`, `upper` and `critcontrol`), `critcst_optimizer` performs the maximization of a constrained Expected Improvement or SUR criterion and delivers the next point to be visited in an EGO-like procedure.

The latter maximization relies either on a genetic algorithm using derivatives, `genoud` or exhaustive search at pre-specified points. It is important to remark that the information needed about the objective and constraint functions reduces here to the vector of response values embedded in the models (no call to the objective/constraint functions or simulators (except possibly for the objective)).

## Usage

```
critcst_optimizer(
  crit = "EFI",
  model.fun,
  model.constraint,
  equality = FALSE,
  lower,
  upper,
  type = "UK",
  critcontrol = NULL,
  optimcontrol = NULL
)
```

## Arguments

<code>crit</code>	sampling criterion. Three choices are available : "EFI", "AL" and "SUR",
<code>model.fun</code>	object of class <code>km</code> corresponding to the objective function, or, if the objective function is fast-to-evaluate, either the objective function to be minimized or a <code>fastfun</code> object, see details and examples below,
<code>model.constraint</code>	either one or a list of models of class <code>km</code> , one for each constraint,
<code>equality</code>	either FALSE if all constraints are inequalities, or a Boolean vector indicating which are equalities,
<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>type</code>	"SK" or "UK" (default), depending whether uncertainty related to trend estimation has to be taken into account.

- critcontrol** optional list of control parameters for criterion `crit`, see details. 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.
- optimcontrol** optional list of control parameters for optimization of the selected infill criterion. "method" set the optimization method; one can choose between "discrete" and "genoud". For each method, further parameters can be set. For "discrete", one has to provide the argument "candidate.points". For "genoud", one can control, among others, "pop.size" (default:  $[N = 3 \cdot 2^{\dim}$  for  $\dim < 6$  and  $N = 32 \cdot \dim$  otherwise]), "max.generations" (12), "wait.generations" (2)), see `genoud`. Numbers into brackets are the default values. @return A list with components:
- `par`: The best set of parameters found,
  - `value`: The value of expected improvement at `par`.

## Details

Extension of the function `max_EI` for constrained optimization.

Available infill criteria with `crit` are :

- Expected Probability of Feasibility (EFI) `crit_EFI`,
- Augmented Lagrangian (AL) `crit_AL`,
- Stepwise Uncertainty Reduction of the excursion volume (SUR) `crit_SUR_cst`.

Depending on the selected criterion, parameters can be given with `critcontrol`. Also options for `checkPredict` are available. More precisions are given in the corresponding help pages.

If the objective function to minimize is inexpensive, i.e. no need of a kriging model, then one can provide it in `model.obj`, which is handled next with class `fastfun` (or directly as a `fastfun` object). See example below.

In the case of equality constraints, it is possible to define them with `equality`. Additionally, one can modify the tolerance on the constraints using the `tolConstraints` component of `critcontrol`: an optional vector giving a tolerance for each of the constraints (equality or inequality). It is highly recommended to use it when there are equality constraints since the default tolerance of 0.05 (resp. 0 for inequality constraints) in such case might not be suited.

## Author(s)

Victor Picheny

Mickael Binois

## 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.

## See Also

[critcst\\_optimizer](#), [crit\\_EFI](#), [crit\\_AL](#), [crit\\_SUR\\_cst](#)

## Examples

```
#-----
# 2D objective function, 2 cases
#-----

set.seed(2546)
library(DiceDesign)

n_var <- 2
fun <- branin

fun1.cst <- function(x){return(goldsteinprice(x)+.5)}
fun2.cst <- function(x){return(3/2 - x[1] - 2*x[2] - .5*sin(2*pi*(x[1]^2 - 2*x[2])))}

# Constraint function with vectorial output
cstfun <- function(x){return(c(fun1.cst(x), fun2.cst(x)))}

n.grid <- 31
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun)
cst1.grid <- apply(test.grid, 1, fun1.cst)
cst2.grid <- apply(test.grid, 1, fun2.cst)

n_appr <- 12
design.grid <- round(maximinESE_LHS(lhsDesign(n_appr, n_var, seed = 2)$design)$design, 1)
obj.init <- apply(design.grid, 1, fun)
cst1.init <- apply(design.grid, 1, fun1.cst)
cst2.init <- apply(design.grid, 1, fun2.cst)
model.fun <- km(~., design = design.grid, response = obj.init)
model.constraint1 <- km(~., design = design.grid, response = cst1.init, lower=c(.2,.2))
model.constraint2 <- km(~., design = design.grid, response = cst2.init, lower=c(.2,.2))
models.cst <- list(model.constraint1, model.constraint2)

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

#-----
# Augmented Lagrangian Improvement, fast objective function, two ineq constraints,
# optimization with genoud
```

```

#-----
critcontrol <- list(lambda=c(.5,2), rho=.5)
optimcontrol <- list(method = "genoud", max.generations=10, pop.size=20)

AL_grid <- apply(test.grid, 1, crit_AL, model.fun = fastfun(fun, design.grid),
               model.constraint = models.cst, critcontrol=critcontrol)

cstEG01 <- critcst_optimizer(crit = "AL", model.fun = fun,
                          model.constraint = models.cst, equality = FALSE,
                          lower = lower, upper = upper,
                          optimcontrol = optimcontrol, critcontrol=critcontrol)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(AL_grid, n.grid), main = "AL map and its maximizer (blue)",
              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")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                          matrix(obj.grid, n.grid), nlevels = 10, add=TRUE,drawlabels=TRUE,
                          col = "black")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                          matrix(cst1.grid, n.grid), level = 0, add=TRUE,drawlabels=FALSE,
                          lwd=1.5, col = "red")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                          matrix(cst2.grid, n.grid), level = 0, add=TRUE,drawlabels=FALSE,
                          lwd=1.5, col = "red")
                          points(cstEG01$par, col = "blue", pch = 4, lwd = 2)
                          }
              )

#-----
# SUR, expensive objective function, one equality constraint,
# optimization with genoud, integration on a regular grid
#-----
optimcontrol <- list(method = "genoud", s = 40, maxit = 40)
critcontrol <- list(tolConstraints = .15, integration.points=as.matrix(test.grid))

SUR_grid <- apply(test.grid, 1, crit_SUR_cst, model.fun = model.fun,
               model.constraint = model.constraint1, equality = TRUE, critcontrol = critcontrol)

cstEG02 <- critcst_optimizer(crit = "SUR", model.fun = model.fun,
                          model.constraint = model.constraint1, equality = TRUE,
                          lower = lower, upper = upper,
                          optimcontrol = optimcontrol, critcontrol = critcontrol)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(SUR_grid, n.grid), main = "SUR map and its maximizer (blue)",
              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")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                          matrix(obj.grid, n.grid), nlevels = 10, add=TRUE,
                          drawlabels=TRUE, col = "black")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),

```

```

matrix(cst1.grid, n.grid), level = c(-critcontrol$tolConstraints,
  critcontrol$tolConstraints),
  add=TRUE, drawlabels=FALSE,lwd=1.5, col = "orange")
  points(cstEG02$par, col = "blue", pch = 4, lwd = 2)
}
)

```

crit\_AL

*Expected Augmented Lagrangian Improvement***Description**

Computes the Expected Augmented Lagrangian Improvement at current location, with or without slack variables. Depending on the cases, the computation is either analytical (very fast), based on MC integration (slow) or on the CDF of a weighted sum of non-central chi-square (WNCS) variates (intermediate)

**Usage**

```

crit_AL(
  x,
  model.fun,
  model.constraint,
  equality = FALSE,
  critcontrol = NULL,
  type = "UK"
)

```

**Arguments**

x	either a vector representing the design or the design AND slack variables (see details)
model.fun	object of class <code>km</code> corresponding to the objective function, or, if the objective function is fast-to-evaluate, a <code>fastfun</code> object,
model.constraint	either one or a list of objects of class <code>km</code> , one for each constraint function,
equality	either FALSE if all constraints are for inequalities, or a vector of Booleans indicating which are equalities
critcontrol	optional list with the following arguments: <ul style="list-style-type: none"> <li>• <code>slack</code>: logical. If TRUE, slack variables are used for inequality constraints (see Details)</li> <li>• <code>rho</code>: penalty term (scalar),</li> <li>• <code>lambda</code>: Lagrange multipliers (vector of size the number of constraints),</li> <li>• <code>elit</code>: logical. If TRUE, sets the criterion to zero for all <code>x</code>'s not improving the objective function</li> </ul>

- `n.mc`: number of Monte-Carlo drawings used to evaluate the criterion (see Details)
- `nt`: number of discretization points for the WNCS distribution (see Details)
- `tolConstraints`, an optional vector giving a tolerance ( $> 0$ ) for each of the constraints (equality or inequality). It is highly recommended to use it when there are equality constraints since the default tolerance of 0.05 in such case might not be suited.

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.

### Details

The AL can be used with or without the help of slack variables for the inequality constraints. If `critcontrol$slack=FALSE`: With a single constraint (inequality or equality) and a fast objective, a very fast formula is used to compute the criterion (recommended setting). Otherwise, an MC estimator of the criterion is used, which is much more costly. The argument `critcontrol$n.mc` tunes the precision of the estimator. On both cases `x` must be of size `d`.

If `critcontrol$slack=TRUE`: Slack variables are used to handle the inequality constraints. They can be provided directly through `x`, which should be of size `d+the number of inequality constraints`. The last values of `x` are slack variables scaled to  $[0,1]$ .

If `x` is of size `d`, estimates of optimal slack variable are used.

### Value

The Expected Augmented Lagrangian Improvement at `x`.

### Author(s)

Victor Picheny

Mickael Binois

### References

R.B. Gramacy, G.A. Gray, S. Le Digabel, H.K.H Lee, P. Ranjan, G. Wells, Garth, and S.M. Wild (2014+), Modeling an augmented Lagrangian for improved blackbox constrained optimization, *arXiv preprint arXiv:1403.4890*.

### See Also

EI from package DiceOptim, [crit\\_EFI](#), [crit\\_SUR\\_cst](#).

## Examples

```

#-----
# Expected Augmented Lagrangian Improvement surface with one inequality constraint,
# fast objective
#-----

set.seed(25468)
library(DiceDesign)

n_var <- 2
fun.obj <- goldsteinprice
fun.cst <- function(x){return(-branin(x) + 25)}
n.grid <- 31
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun.obj)
cst.grid <- apply(test.grid, 1, fun.cst)
n.init <- 15
design.grid <- round(maximinESE_LHS(lhsDesign(n.init, n_var, seed = 42)$design)$design, 1)
obj.init <- apply(design.grid, 1, fun.obj)
cst.init <- apply(design.grid, 1, fun.cst)
model.constraint <- km(~., design = design.grid, response = cst.init)
model.fun <- fastfun(fun.obj, design.grid)
AL_grid <- apply(test.grid, 1, crit_AL, model.fun = model.fun,
                model.constraint = model.constraint)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(AL_grid, n.grid), main = "Expected AL 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")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                matrix(obj.grid, n.grid), nlevels = 10,
                                add=TRUE,drawlabels=TRUE, col = "black")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                matrix(cst.grid, n.grid), level = 0, add=TRUE,
                                drawlabels=FALSE,lwd=1.5, col = "red")
                          }
              )

#-----
# Expected AL Improvement surface with one inequality and one equality constraint,
# using slack variables
#-----

set.seed(25468)
library(DiceDesign)

n_var <- 2
fun.obj <- goldsteinprice
fun.cstineq <- function(x){return(3/2 - x[1] - 2*x[2] - .5*sin(2*pi*(x[1]^2 - 2*x[2])))}
fun.csteq <- function(x){return(branin(x) - 25)}
n.grid <- 51

```

```

test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun.obj)
cstineq.grid <- apply(test.grid, 1, fun.cstineq)
csteq.grid <- apply(test.grid, 1, fun.csteq)
n.init <- 25
design.grid <- round(maximinESE_LHS(lhsDesign(n.init, n_var, seed = 42)$design)$design, 1)
obj.init <- apply(design.grid, 1, fun.obj)
cstineq.init <- apply(design.grid, 1, fun.cstineq)
csteq.init <- apply(design.grid, 1, fun.csteq)
model.fun <- km(~., design = design.grid, response = obj.init)
model.constraintineq <- km(~., design = design.grid, response = cstineq.init)
model.constrainedeq <- km(~., design = design.grid, response = csteq.init)

models.cst <- list(model.constraintineq, model.constrainedeq)

AL_grid <- apply(test.grid, 1, crit_AL, model.fun = model.fun, model.constraint = models.cst,
  equality = c(FALSE, TRUE), critcontrol = list(tolConstraints = c(0.05, 3),
  slack=TRUE))

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
  matrix(AL_grid, n.grid), main = "Expected AL 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")
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
    matrix(obj.grid, n.grid), nlevels = 10,
    add=TRUE,drawlabels=TRUE, col = "black")
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
    matrix(cstineq.grid, n.grid), level = 0, add=TRUE,
    drawlabels=FALSE,lwd=1.5, col = "red")
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
    matrix(csteq.grid, n.grid), level = 0, add=TRUE,
    drawlabels=FALSE,lwd=1.5, col = "orange")
  }
)

```

---

crit\_EFI

*Expected Feasible Improvement*


---

### Description

Computes the Expected Feasible Improvement at current location. The current feasible minimum of the observations can be replaced by an arbitrary value (plugin), which is useful in particular in noisy frameworks.

### Usage

```

crit_EFI(
  x,

```



```

    model.fun,
    model.constraint,
    equality = FALSE,
    critcontrol = NULL,
    plugin = NULL,
    type = "UK"
)

```

## Arguments

<code>x</code>	a vector representing the input for which one wishes to calculate EFI,
<code>model.fun</code>	object of class <code>km</code> corresponding to the objective function, or, if the objective function is fast-to-evaluate, a <code>fastfun</code> object,
<code>model.constraint</code>	either one or a list of objects of class <code>km</code> , one for each constraint function,
<code>equality</code>	either FALSE if all constraints are for inequalities, else a vector of boolean indicating which are equalities,
<code>critcontrol</code>	optional list with argument <code>tolConstraints</code> , an optional vector giving a tolerance ( $> 0$ ) for each of the constraints (equality or inequality). It is highly recommended to use it when there are equality constraints since the default tolerance of 0.05 in such case might not be suited.
	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.
<code>plugin</code>	optional scalar: if provided, it replaces the feasible minimum of the current observations. If set to <code>Inf</code> , e.g. when there is no feasible solution, then the criterion is equal to the probability of feasibility,
<code>type</code>	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account.

## Value

The Expected Feasible Improvement at `x`.

## Author(s)

Victor Picheny  
Mickael Binois

## References

- M. Schonlau, W.J. Welch, and D.R. Jones (1998), Global versus local search in constrained optimization of computer models, *Lecture Notes-Monograph Series*, 11-25.
- M.J. Sasena, P. Papalambros, and P.Goovaerts (2002), Exploration of metamodeling sampling criteria for constrained global optimization, *Engineering optimization*, 34, 263-278.

**See Also**

EI from package DiceOptim, [crit\\_AL](#), [crit\\_SUR\\_cst](#).

**Examples**

```

#-----
# Expected Feasible Improvement surface with one inequality constraint
#-----

set.seed(25468)
library(DiceDesign)

n_var <- 2
fun.obj <- goldsteinprice
fun.cst <- function(x){return(-branin(x) + 25)}
n.grid <- 51
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun.obj)
cst.grid <- apply(test.grid, 1, fun.cst)
n.init <- 15
design.grid <- round(maximinESE_LHS(lhsDesign(n.init, n_var, seed = 42)$design)$design, 1)
obj.init <- apply(design.grid, 1, fun.obj)
cst.init <- apply(design.grid, 1, fun.cst)
model.fun <- km(~., design = design.grid, response = obj.init)
model.constraint <- km(~., design = design.grid, response = cst.init)

EFI_grid <- apply(test.grid, 1, crit_EFI, model.fun = model.fun,
                 model.constraint = model.constraint)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(EFI_grid, n.grid), main = "Expected Feasible 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")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                  matrix(obj.grid, n.grid), nlevels = 10,
                                  add=TRUE,drawlabels=TRUE, col = "black")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                  matrix(cst.grid, n.grid), level = 0, add=TRUE,
                                  drawlabels=FALSE,lwd=1.5, col = "red")
                          }
              )

#-----
# Expected Feasible Improvement surface with one inequality and one equality constraint
#-----

set.seed(25468)
library(DiceDesign)

n_var <- 2
fun.obj <- goldsteinprice

```

```

fun.cstineq <- function(x){return(3/2 - x[1] - 2*x[2] - .5*sin(2*pi*(x[1]^2 - 2*x[2])))}
fun.csteq <- function(x){return(branin(x) - 25)}
n.grid <- 51
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun.obj)
cstineq.grid <- apply(test.grid, 1, fun.cstineq)
csteq.grid <- apply(test.grid, 1, fun.csteq)
n.init <- 25
design.grid <- round(maximinESE_LHS(lhsDesign(n.init, n_var, seed = 42)$design)$design, 1)
obj.init <- apply(design.grid, 1, fun.obj)
cstineq.init <- apply(design.grid, 1, fun.cstineq)
csteq.init <- apply(design.grid, 1, fun.csteq)
model.fun <- km(~., design = design.grid, response = obj.init)
model.constraintineq <- km(~., design = design.grid, response = cstineq.init)
model.constrainteq <- km(~., design = design.grid, response = csteq.init)

models.cst <- list(model.constraintineq, model.constrainteq)

EFI_grid <- apply(test.grid, 1, crit_EFI, model.fun = model.fun, model.constraint = models.cst,
  equality = c(FALSE, TRUE), critcontrol = list(tolConstraints = c(0.05, 3)))

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
  matrix(EFI_grid, n.grid), main = "Expected Feasible 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")
    contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
      matrix(obj.grid, n.grid), nlevels = 10,
      add=TRUE,drawlabels=TRUE, col = "black")
    contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
      matrix(cstineq.grid, n.grid), level = 0, add=TRUE,
      drawlabels=FALSE,lwd=1.5, col = "red")
    contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
      matrix(csteq.grid, n.grid), level = 0, add=TRUE,
      drawlabels=FALSE,lwd=1.5, col = "orange")
  }
)

```

crit\_SUR\_cst

*Stepwise Uncertainty Reduction criterion***Description**

Computes the Stepwise Uncertainty Reduction (SUR) criterion at current location

**Usage**

```
crit_SUR_cst(
  x,
```

```

model.fun,
model.constraint,
equality = FALSE,
critcontrol = NULL,
type = "UK"
)

```

### Arguments

x	a vector representing the input for which one wishes to calculate SUR,
model.fun	object of class <code>km</code> corresponding to the objective function, or, if the objective function is fast-to-evaluate, a <code>fastfun</code> object,
model.constraint	either one or a list of objects of class <code>km</code> , one for each constraint function,
equality	either FALSE if all constraints are for inequalities, else a vector of boolean indicating which are equalities
critcontrol	optional list with arguments: <ul style="list-style-type: none"> <li>• <code>tolConstraints</code> optional vector giving a tolerance (<math>&gt; 0</math>) for each of the constraints (equality or inequality). It is highly recommended to use it when there are equality constraints since the default tolerance of 0.05 in such case might not be suited;</li> <li>• <code>integration.points</code> and <code>integration.weights</code>: optional matrix and vector of integration points;</li> <li>• <code>precalc.data.cst</code>, <code>precalc.data.obj</code>, <code>mn.X.cst</code>, <code>sn.X.cst</code>, <code>mn.X.obj</code>, <code>sn.X.obj</code>: useful quantities for the fast evaluation of the criterion.</li> <li>• Options for the <code>checkPredict</code> function: <code>threshold</code> (<math>1e-4</math>) and <code>distance</code> (<code>covdist</code>) are used to avoid numerical issues occurring when adding points too close to the existing ones.</li> </ul>
type	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account.

### Value

The Stepwise Uncertainty Reduction criterion at  $x$ .

### Author(s)

Victor Picheny  
Mickael Binois

### References

V. Picheny (2014), A stepwise uncertainty reduction approach to constrained global optimization, *Proceedings of the 17th International Conference on Artificial Intelligence and Statistics*, JMLR W&CP 33, 787-795.

**See Also**

EI from package DiceOptim, [crit\\_EFI](#), [crit\\_AL](#).

**Examples**

```
#-----
# Stepwise Uncertainty Reduction criterion surface with one inequality constraint
#-----

set.seed(25468)
library(DiceDesign)

n_var <- 2
fun.obj <- goldsteinprice
fun.cst <- function(x){return(-branin(x) + 25)}
n.grid <- 21
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun.obj)
cst.grid <- apply(test.grid, 1, fun.cst)

n_appr <- 15
design.grid <- round(maximinESE_LHS(lhsDesign(n_appr, n_var, seed = 42)$design)$design, 1)
obj.init <- apply(design.grid, 1, fun.obj)
cst.init <- apply(design.grid, 1, fun.cst)
model.fun <- km(~., design = design.grid, response = obj.init)
model.constraint <- km(~., design = design.grid, response = cst.init)

integration.param <- integration_design_cst(integcontrol =list(integration.points = test.grid),
                                           lower = rep(0, n_var), upper = rep(1, n_var))

SUR_grid <- apply(test.grid, 1, crit_SUR_cst, model.fun = model.fun,
                 model.constraint = model.constraint, critcontrol=integration.param)

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
               matrix(SUR_grid, n.grid), main = "SUR 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")
                           contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                   matrix(obj.grid, n.grid), nlevels = 10,
                                   add=TRUE,drawlabels=TRUE, col = "black")
                           contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                   matrix(cst.grid, n.grid), level = 0, add=TRUE,
                                   drawlabels=FALSE,lwd=1.5, col = "red")
                           }
               )

#-----
# SUR with one inequality and one equality constraint
#-----

set.seed(25468)
```

```

library(DiceDesign)

n_var <- 2
fun.obj <- goldsteinprice
fun.cstineq <- function(x){return(3/2 - x[1] - 2*x[2] - .5*sin(2*pi*(x[1]^2 - 2*x[2])))}
fun.csteq <- function(x){return(branin(x) - 25)}
n.grid <- 21
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun.obj)
cstineq.grid <- apply(test.grid, 1, fun.cstineq)
csteq.grid <- apply(test.grid, 1, fun.csteq)
n_appr <- 25
design.grid <- round(maximinESE_LHS(lhsDesign(n_appr, n_var, seed = 42)$design)$design, 1)
obj.init <- apply(design.grid, 1, fun.obj)
cstineq.init <- apply(design.grid, 1, fun.cstineq)
csteq.init <- apply(design.grid, 1, fun.csteq)
model.fun <- km(~., design = design.grid, response = obj.init)
model.constraintineq <- km(~., design = design.grid, response = cstineq.init)
model.constrainteq <- km(~., design = design.grid, response = csteq.init)

models.cst <- list(model.constraintineq, model.constrainteq)

SUR_grid <- apply(test.grid, 1, crit_SUR_cst, model.fun = model.fun, model.constraint = models.cst,
  equality = c(FALSE, TRUE), critcontrol = list(tolConstraints = c(0.05, 3),
  integration.points=integration.param$integration.points))

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
  matrix(SUR_grid, n.grid), main = "SUR 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")
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
  matrix(obj.grid, n.grid), nlevels = 10,
  add=TRUE,drawlabels=TRUE, col = "black")
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
  matrix(cstineq.grid, n.grid), level = 0, add=TRUE,
  drawlabels=FALSE,lwd=1.5, col = "red")
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
  matrix(csteq.grid, n.grid), level = 0, add=TRUE,
  drawlabels=FALSE,lwd=1.5, col = "orange")
  }
)

```

**Description**

User-friendly wrapper of the functions `fastEGO.nsteps` and `TREGO.nsteps`. Generates initial DOEs and kriging models (objects of class `km`), and executes `nsteps` iterations of either EGO or TREGO.

**Usage**

```
easyEGO(
  fun,
  budget,
  lower,
  upper,
  X = NULL,
  y = NULL,
  control = list(trace = 1, seed = 42),
  n.cores = 1,
  ...
)
```

**Arguments**

<code>fun</code>	scalar function to be minimized,
<code>budget</code>	total number of calls to the objective and constraint 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>X</code>	initial design of experiments. If not provided, <code>X</code> is taken as a maximin LHD with <code>budget/3</code> points
<code>y</code>	initial set of objective observations $f(X)$ . Computed if not provided.
<code>control</code>	an optional list of control parameters. See "Details".
<code>n.cores</code>	number of cores for parallel computation
<code>...</code>	additional parameters to be given to <code>fun</code>

**Details**

Does not require knowledge on kriging models (objects of class `km`)

The control argument is a list that can supply any of the following components:

- `trace`: between -1 and 3
- `seed`: to fix the seed of the run
- `cov.reestim`: Boolean, if TRUE (default) the covariance parameters are re-estimated at each iteration
- `model.trend`: trend for the GP model
- `lb`, `ub`: lower and upper bounds for the GP covariance ranges
- `nugget`: optional nugget effect

- `covtype`: covariance of the GP model (default "matern5\_2")
- `optim.method`: optimisation of the GP hyperparameters (default "BFGS")
- `multistart`: number of restarts of BFGS
- `gpmean.trick`, `gpmean.freq`: Boolean and integer, resp., for the `gpmean` trick
- `scaling`: Boolean, activates input scaling
- `warping`: Boolean, activates output warping
- `TR`: Boolean, activates TREGO instead of EGO
- `trcontrol`: list of parameters of the trust region, see [TREGO.nsteps](#)
- `always.sample`: Boolean, activates force observation even if it leads to poor conditioning

### Value

A list with components:

- `par`: the best feasible point
- `values`: a vector of the objective and constraints at the point given in `par`,
- `history`: a list containing all the points visited by the algorithm ( $X$ ) and their corresponding objectives ( $y$ ).
- `model`: the last GP model, class `km`
- `control`: full list of control values, see "Details"
- `res`: the output of either [fastEGO.nsteps](#) or [TREGO.nsteps](#)

### Author(s)

Victor Picheny

### References

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

```
library(parallel)
library(DiceOptim)
set.seed(123)

#####
### 10 ITERATIONS OF TREGO ON THE BRANIN FUNCTION, ###
### STARTING FROM A 9-POINTS FACTORIAL DESIGN      ###
#####

# a 9-points factorial design, and the corresponding response
ylim=NULL
fun <- branin; d <- 2
budget <- 5*d
lower <- rep(0,d)
```



```

upper <- rep(1,d)
n.init <- 2*d

control <- list(n.init=2*d, TR=TRUE, nugget=1e-5, trcontrol=list(algo="TREGO"), multistart=1)

res1 <- easyEGO(fun=fun, budget=budget, lower=lower, upper=upper, control=control, n.cores=1)

par(mfrow=c(3,1))
y <- res1$history$y
steps <- res1$res$all.steps
success <- res1$res$all.success
sigma <- res1$res$all.sigma
ymin <- cummin(y)
pch <- rep(1, length(sigma))
col <- rep("red", length(sigma))
pch[which(!steps)] <- 2
col[which(success)] <- "darkgreen"

pch2 <- c(rep(3, n.init), pch)
col2 <- c(rep("black", n.init), col)
plot(y, col=col2, ylim=ylim, pch=pch2, lwd=2, xlim=c(0, budget))
lines(ymin, col="darkgreen")
abline(v=n.init+.5)

plot(n.init + (1:length(sigma)), sigma, xlim=c(0, budget), ylim=c(0, max(sigma)),
pch=pch, col=col, lwd=2, main="TR size")
lines(n.init + (1:length(sigma)), sigma, xlim=c(0, budget))
abline(v=n.init+.5)

plot(NA, xlim=c(0, budget), ylim=c(0, 1), main="x0 (coordinates)")
for (i in 1:d) {
  lines(n.init + (1:nrow(res1$res$all.x0)), res1$res$all.x0[,i])
  points(n.init + (1:nrow(res1$res$all.x0)), res1$res$all.x0[,i], pch=pch, col=col, lwd=2)
}
abline(v=n.init+.5)

par(mfrow=c(1,1))
pairs(res1$model@X, pch=pch2, col=col2)

```

**Description**

User-friendly wrapper of the function `EGO.cst` Generates initial DOEs and kriging models (objects of class `km`), and executes `nsteps` iterations of EGO methods integrating constraints.

**Usage**

```

easyEGO.cst(
  fun,
  constraint,
  n.cst = 1,
  budget,
  lower,
  upper,
  cheapfun = FALSE,
  equality = FALSE,
  X = NULL,
  y = NULL,
  C = NULL,
  control = list(method = "EFI", trace = 1, inneroptim = "genoud", maxit = 100, seed =
    42),
  ...
)

```

**Arguments**

fun	scalar function to be minimized,
constraint	vectorial function corresponding to the constraints, see details below,
n.cst	number of constraints,
budget	total number of calls to the objective and constraint functions,
lower	vector of lower bounds for the variables to be optimized over,
upper	vector of upper bounds for the variables to be optimized over,
cheapfun	optional boolean, TRUE if the objective is a fast-to-evaluate function that does not need a kriging model
equality	either FALSE if all constraints are inequalities, else a Boolean vector indicating which are equalities
X	initial design of experiments. If not provided, X is taken as a maximin LHD with budget/3 points
y	initial set of objective observations $f(X)$ . Computed if not provided.
C	initial set of constraint observations $g(X)$ . Computed if not provided.
control	an optional list of control parameters. See "Details".
...	additional parameters to be given to BOTH the objective fun and constraints.

**Details**

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

The problem considered is of the form:  $\min f(x)$  s.t.  $g(x) \leq 0$ ,  $g$  having a vectorial output. By default all its components are supposed to be inequalities, but one can use a Boolean vector in equality to specify which are equality constraints, hence of the type  $g(x) = 0$ . The control argument is a list that can supply any of the following components:

- method: choice of constrained improvement function: "AL", "EFI" or "SUR" (see [crit\\_EFI](#), [crit\\_AL](#), [crit\\_SUR\\_cst](#))
- trace: if positive, tracing information on the progress of the optimization is produced.
- inneroptim: choice of the inner optimization algorithm: "genoud" or "random" (see [genoud](#)).
- maxit: maximum number of iterations of the inner loop.
- seed: to fix the random variable generator

For additional details, see [EGO.cst](#).

### Value

A list with components:

- par: the best feasible point
- values: a vector of the objective and constraints at the point given in par,
- history: a list containing all the points visited by the algorithm (X) and their corresponding objectives (y) and constraints (C)

If no feasible point is found, par returns the most feasible point (in the least square sense).

### Author(s)

Victor Picheny

Mickael Binois

### References

- 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.
- M. Schonlau, W.J. Welch, and D.R. Jones (1998), Global versus local search in constrained optimization of computer models, *Lecture Notes-Monograph Series*, 11-25.
- M.J. Sasena, P. Papalambros, and P.Goovaerts (2002), Exploration of metamodeling sampling criteria for constrained global optimization, *Engineering optimization*, 34, 263-278.
- R.B. Gramacy, G.A. Gray, S. Le Digabel, H.K.H Lee, P. Ranjan, G. Wells, Garth, and S.M. Wild (2014+), Modeling an augmented Lagrangian for improved blackbox constrained optimization, *arXiv preprint arXiv:1403.4890*.
- J.M. Parr (2012), *Improvement criteria for constraint handling and multiobjective optimization*, University of Southampton.
- V. Picheny (2014), A stepwise uncertainty reduction approach to constrained global optimization, *Proceedings of the 17th International Conference on Artificial Intelligence and Statistics*, JMLR W&CP 33, 787-795.

## Examples

```

#-----
# 2D objective function, 3 cases
#-----

set.seed(25468)
library(DiceDesign)

n_var <- 2
fun <- goldsteinprice
fun1.cst <- function(x){return(-branin(x) + 25)}
fun2.cst <- function(x){return(3/2 - x[1] - 2*x[2] - .5*sin(2*pi*(x[1]^2 - 2*x[2])))}

# Constraint function with vectorial output
constraint <- function(x){return(c(fun1.cst(x), fun2.cst(x)))}

# For illustration purposes
n.grid <- 31
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun)
cst1.grid <- apply(test.grid, 1, fun1.cst)
cst2.grid <- apply(test.grid, 1, fun2.cst)

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

#-----
# 1- Expected Feasible Improvement criterion, expensive objective function,
# two inequality constraints, 15 observations budget, using genoud
#-----
res <- easyEGO.cst(fun=fun, constraint=constraint, n.cst=2, lower=lower, upper=upper, budget=15,
                  control=list(method="EFI", inneroptim="genoud", maxit=20))

cat("best design found:", res$par, "\n")
cat("corresponding objective and constraints:", res$value, "\n")

# Objective function in colour, constraint boundaries in red
# Initial DoE: white circles, added points: blue crosses, best solution: red cross

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(obj.grid, n.grid), main = "Two inequality constraints",
              xlab = expression(x[1]), ylab = expression(x[2]), color = terrain.colors,
              plot.axes = {axis(1); axis(2);
                contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                      matrix(cst1.grid, n.grid), level = 0, add=TRUE,
                      drawlabels=FALSE, lwd=1.5, col = "red")
                contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                      matrix(cst2.grid, n.grid), level = 0, add=TRUE, drawlabels=FALSE,
                      lwd=1.5, col = "red")
                points(res$history$X, col = "blue", pch = 4, lwd = 2)
                points(res$par[1], res$par[2], col = "red", pch = 4, lwd = 2, cex=2)
              }
}

```

```

)

#-----
# 2- Augmented Lagrangian Improvement criterion, expensive objective function,
# one inequality and one equality constraint, 25 observations budget, using random search
#-----
res2 <- easyEGO.cst(fun=fun, constraint=constraint, n.cst=2, lower=lower, upper=upper, budget=25,
  equality = c(TRUE, FALSE),
  control=list(method="AL", inneroptim="random", maxit=100))

cat("best design found:", res2$par, "\n")
cat("corresponding objective and constraints:", res2$value, "\n")

# Objective function in colour, inequality constraint boundary in red, equality
# constraint in orange
# Initial DoE: white circles, added points: blue crosses, best solution: red cross

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
  matrix(obj.grid, n.grid), xlab = expression(x[1]), ylab = expression(x[2]),
  main = "Inequality (red) and equality (orange) constraints", color = terrain.colors,
  plot.axes = {axis(1); axis(2);
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
    matrix(cst1.grid, n.grid), level = 0, add=TRUE,
    drawlabels=FALSE,lwd=1.5, col = "orange")
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
    matrix(cst2.grid, n.grid), level = 0, add=TRUE,
    drawlabels=FALSE,lwd=1.5, col = "red")
  points(res2$history$X, col = "blue", pch = 4, lwd = 2)
  points(res2$par[1], res2$par[2], col = "red", pch = 4, lwd = 2, cex=2)
  }
)

#-----
# 3- Stepwise Uncertainty Reduction criterion, fast objective function,
# single inequality constraint, with initial DOE given + 10 observations,
# using genoud
#-----
n.init <- 12
design.grid <- round(maximinESE_LHS(lhsDesign(n.init, n_var, seed = 42)$design)$design, 1)
cst2.init <- apply(design.grid, 1, fun2.cst)

res3 <- easyEGO.cst(fun=fun, constraint=fun2.cst, n.cst=1, lower=lower, upper=upper, budget=10,
  X=design.grid, C=cst2.init,
  cheapfun=TRUE, control=list(method="SUR", inneroptim="genoud", maxit=20))

cat("best design found:", res3$par, "\n")
cat("corresponding objective and constraint:", res3$value, "\n")

# Objective function in colour, inequality constraint boundary in red
# Initial DoE: white circles, added points: blue crosses, best solution: red cross

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
  matrix(obj.grid, n.grid), main = "Single constraint, fast objective",

```

```

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")
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
    matrix(obj.grid, n.grid), nlevels = 10, add = TRUE,
    drawlabels = TRUE)
  contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
    matrix(cst2.grid, n.grid), level = 0, add = TRUE,
    drawlabels = FALSE, lwd = 1.5, col = "red")
  points(res3$history$X, col = "blue", pch = 4, lwd = 2)
  points(res3$par[1], res3$par[2], col = "red", pch = 4, lwd = 2, cex=2)
}
)

```

---

EGO.cst

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

---

## Description

Executes `nsteps` iterations of EGO methods integrating constraints, based on 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 constrained Expected Improvement criteria available.

## Usage

```

EGO.cst(
  model.fun = NULL,
  fun,
  cheapfun = NULL,
  model.constraint,
  constraint,
  equality = FALSE,
  crit = "EFI",
  nsteps,
  lower,
  upper,
  type = "UK",
  cov.reestim = TRUE,
  critcontrol = NULL,
  optimcontrol = list(method = "genoud", threshold = 1e-05, distance = "euclidean",
    notrace = FALSE),
  ...
)

```

**Arguments**

<code>model.fun</code>	object of class <code>km</code> corresponding to the objective function,
<code>fun</code>	scalar function to be minimized, corresponding to <code>model.fun</code> found by a call to <code>match.fun</code> ,
<code>cheapfun</code>	optional scalar function to use if the objective is a fast-to-evaluate function (handled next with class <code>fastfun</code> , through the use of <code>match.fun</code> ), which does not need a kriging model, see details below,
<code>model.constraint</code>	either one or a list of models of class <code>km</code> , one per constraint,
<code>constraint</code>	vectorial function corresponding to the constraints, see details below,
<code>equality</code>	either FALSE if all constraints are for inequalities, else a vector of boolean indicating which are equalities
<code>crit</code>	choice of constrained improvement function: "AL", "EFI" or "SUR", see details below,
<code>nsteps</code>	an integer representing the desired number of iterations,
<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>type</code>	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account,
<code>cov.reestim</code>	optional boolean specifying if the kriging hyperparameters should be re-estimated at each iteration,
<code>critcontrol</code>	optional list of parameters for criterion <code>crit</code> , see details,
<code>optimcontrol</code>	an optional list of control parameters for optimization of the selected infill criterion: <ul style="list-style-type: none"> <li>• method can be set to "discrete" or "genoud". For "discrete", a matrix <code>candidate.points</code> must be given, For "genoud", specific parameters to the chosen method can also be specified (see <code>genoud</code>).</li> <li>• 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.</li> <li>• <code>notrace</code> can be set to TRUE to suppress printing of the optimization progresses.</li> </ul>
<code>...</code>	additional parameters to be given to the objective <code>fun</code> and <code>constraint</code> .

**Details**

Extension of the function `EGO.nsteps` to constrained optimization.

The problem considered is of the form:  $\min f(x)$  s.t.  $g(x) \leq 0$ ,  $g$  having a vectorial output. By default all its components are supposed to be inequalities, but one can use a boolean vector in `equality` to specify which are equality constraints. In this case one can modify the tolerance on the constraints using the `tolConstraints` component of `critcontrol`: an optional vector giving a tolerance for each of the constraints (equality or inequality). It is highly recommended to use it

when there are equality constraints since the default tolerance of  $0.05$  in such case might not be suited.

Available infill criteria with `crit` are:

- Expected Probability of Feasibility (EFI) `crit_EFI`,
- Augmented Lagrangian (AL) `crit_AL`,
- Stepwise Uncertainty Reduction of the excursion volume (SUR) `crit_SUR_cst`.

Depending on the selected criterion, various parameters are available. More precisions are given in the corresponding help pages.

It is possible to consider a cheap to evaluate objective function submitted to expensive constraints. In this case, provide only a function in `cheapfun`, with both `model.fun` and `fun` to `NULL`, see examples below.

### Value

A list with components:

- `par`: a matrix representing the additional points visited during the algorithm,
- `values`: a vector representing the response (objective) values at the points given in `par`,
- `constraint`: a matrix representing the constraints values at the points given in `par`,
- `feasibility`: a boolean vector saying if points given in `par` respect the constraints,
- `nsteps`: an integer representing the desired number of iterations (given in argument),
- `lastmodel.fun`: an object of class `km` corresponding to the objective function,
- `lastmodel.constraint`: one or a list of objects of class `km` corresponding to the last kriging models fitted to the constraints.

If a problem occurs during either model updates or criterion maximization, the last working model and corresponding values are returned.

### Author(s)

Victor Picheny

Mickael Binois

### References

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.

M. Schonlau, W.J. Welch, and D.R. Jones (1998), Global versus local search in constrained optimization of computer models, *Lecture Notes-Monograph Series*, 11-25.

M.J. Sasena, P. Papalambros, and P.Goovaerts (2002), Exploration of metamodeling sampling criteria for constrained global optimization, *Engineering optimization*, 34, 263-278.



R.B. Gramacy, G.A. Gray, S. Le Digabel, H.K.H Lee, P. Ranjan, G. Wells, Garth, and S.M. Wild (2014+), Modeling an augmented Lagrangian for improved blackbox constrained optimization, *arXiv preprint arXiv:1403.4890*.

J.M. Parr (2012), *Improvement criteria for constraint handling and multiobjective optimization*, University of Southampton.

V. Picheny (2014), A stepwise uncertainty reduction approach to constrained global optimization, *Proceedings of the 17th International Conference on Artificial Intelligence and Statistics*, JMLR W&CP 33, 787-795.

### See Also

[critcst\\_optimizer](#), [crit\\_EFI](#), [crit\\_AL](#), [crit\\_SUR\\_cst](#), [easyEGO.cst](#)

### Examples

```
#-----
# 2D objective function, 3 cases
#-----

set.seed(25468)
library(DiceDesign)

n_var <- 2
fun <- goldsteinprice
fun1.cst <- function(x){return(-branin(x) + 25)}
fun2.cst <- function(x){return(3/2 - x[1] - 2*x[2] - .5*sin(2*pi*(x[1]^2 - 2*x[2])))}

# Constraint function with vectorial output
cstfun <- function(x){
  return(c(fun1.cst(x), fun2.cst(x)))
}

# For illustration purposes
n.grid <- 31
test.grid <- expand.grid(X1 = seq(0, 1, length.out = n.grid), X2 = seq(0, 1, length.out = n.grid))
obj.grid <- apply(test.grid, 1, fun)
cst1.grid <- apply(test.grid, 1, fun1.cst)
cst2.grid <- apply(test.grid, 1, fun2.cst)

# Initial set of observations and models
n.init <- 12
design.grid <- round(maximinESE_LHS(lhsDesign(n.init, n_var, seed = 42)$design)$design, 1)
obj.init <- apply(design.grid, 1, fun)
cst1.init <- apply(design.grid, 1, fun1.cst)
cst2.init <- apply(design.grid, 1, fun2.cst)
model.fun <- km(~., design = design.grid, response = obj.init)
model.constraint1 <- km(~., design = design.grid, response = cst1.init, lower=c(.2,.2))
model.constraint2 <- km(~., design = design.grid, response = cst2.init, lower=c(.2,.2))
model.constraint <- list(model.constraint1, model.constraint2)

lower <- rep(0, n_var)
```

```

upper <- rep(1, n_var)

#-----
# 1- Expected Feasible Improvement criterion, expensive objective function,
# two inequality constraints, 5 iterations, using genoud
#-----

cstEGO <- EGO.cst(model.fun = model.fun, fun = fun, model.constraint = model.constraint,
                 crit = "EFI", constraint = cstfun, equality = FALSE, lower = lower,
                 upper = upper, nsteps = 5, optimcontrol = list(method = "genoud", maxit = 20))

# Plots: objective function in colour, constraint boundaries in red
# Initial DoE: white circles, added points: blue crosses, best solution: red cross

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(obj.grid, n.grid), main = "Two inequality constraints",
              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")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                matrix(cst1.grid, n.grid), level = 0, add=TRUE,drawlabels=FALSE,
                                lwd=1.5, col = "red")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                matrix(cst2.grid, n.grid), level = 0, add=TRUE,drawlabels=FALSE,
                                lwd=1.5, col = "red")
                          points(cstEGO$par, col = "blue", pch = 4, lwd = 2)
                          }
              )

#-----
# 2- Augmented Lagrangian Improvement criterion, expensive objective function,
# one inequality and one equality constraint, using a discrete set of candidates (grid)
#-----

cstEG02 <- EGO.cst(model.fun = model.fun, fun = fun, model.constraint = model.constraint,
                  crit = "AL", constraint = cstfun, equality = c(TRUE, FALSE), lower = lower,
                  upper = upper, nsteps = 10,
                  critcontrol = list(tolConstraints = c(2, 0), always.update=TRUE),
                  optimcontrol=list(method="discrete", candidate.points=as.matrix(test.grid)))

# Plots: objective function in colour, inequality constraint boundary in red,
# equality constraint in orange
# Initial DoE: white circles, added points: blue crosses, best solution: red cross

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
              matrix(obj.grid, n.grid),
              main = "Inequality (red) and equality (orange) constraints",
              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")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                matrix(cst1.grid, n.grid), level = 0, add=TRUE,
                                drawlabels=FALSE,lwd=1.5, col = "orange")
                          contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),

```

```

                                matrix(cst2.grid, n.grid), level = 0, add=TRUE,
                                drawlabels=FALSE,lwd=1.5, col = "red")
                                points(cstEGO2$par, col = "blue", pch = 4, lwd = 2)
                                }
                                )

#-----
# 3- Stepwise Uncertainty Reduction criterion, fast objective function,
# single inequality constraint, 5 steps, importance sampling scheme
#-----

cstEGO3 <- EGO.cst(model.fun = NULL, fun = NULL, cheapfun = fun,
                  model.constraint = model.constraint2, constraint = fun2.cst,
                  crit = "SUR", lower = lower, upper = upper,
                  nsteps =5, critcontrol=list(distrib="SUR"))

# Plots: objective function in colour, inequality constraint boundary in red,
# Initial DoE: white circles, added points: blue crosses, best solution: red cross

filled.contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid), nlevels = 50,
               matrix(obj.grid, n.grid), main = "Single constraint, fast objective",
               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")
                           contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                   matrix(obj.grid, n.grid), nlevels = 10, add = TRUE,
                                   drawlabels = TRUE)
                           contour(seq(0, 1, length.out = n.grid), seq(0, 1, length.out = n.grid),
                                   matrix(cst2.grid, n.grid), level = 0, add=TRUE,
                                   drawlabels=FALSE,lwd=1.5, col = "black")
                           points(cstEGO3$par, col = "blue", pch = 4, lwd = 2)
                           }
               )

```

---

EGO.nsteps

*Sequential EI maximization and model re-estimation, with a number of iterations fixed in advance by the user*

---

### Description

Executes *nsteps* iterations of the EGO method to an object of class `km`. At each step, a kriging model is 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 the Expected Improvement criterion (EI).

### Usage

```
EGO.nsteps(
```

```

model,
fun,
nsteps,
lower,
upper,
parinit = NULL,
control = NULL,
kmcontrol = NULL
)

```

### Arguments

model	an object of class <a href="#">km</a> ,
fun	the objective function to be minimized,
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,
parinit	optional vector of initial values for the variables to be optimized over,
control	an optional list of control parameters for optimization. One can control "pop.size" (default : [4+3*log(nb of variables)]), "max.generations" (default :5), "wait.generations" (default :2), "BFGSburnin" (default :0), of the function <a href="#">genoud</a> .
kmcontrol	an optional list representing the control variables for the re-estimation of the kriging model. The items are the same as in <a href="#">km</a> : penalty, optim.method, parinit, control. The default values are those contained in model, typically corresponding to the variables used in <a href="#">km</a> to estimate a kriging model from the initial design points.

### Value

A list with components:

par	a data frame representing the additional points visited during the algorithm,
value	a data frame representing the response values at the points given in par,
npoints	an integer representing the number of parallel computations (=1 here),
nsteps	an integer representing the desired number of iterations (given in argument),
lastmodel	an object of class <a href="#">km</a> corresponding to the last kriging model fitted.

### Note

Most EGO-like methods (EI algorithms) usually work with Ordinary Kriging (constant trend), by maximization of the expected improvement. Here, the EI maximization is also possible with any linear trend. However, note that the optimization may perform much faster and better when the trend is a constant since it is the only case where the analytical gradient is available.

For more details on kmcontrol, see the documentation of [km](#).

**Author(s)**

David Ginsbourger  
Olivier Roustant

**References**

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.

J. Mockus (1988), *Bayesian Approach to Global Optimization*. Kluwer academic publishers.

T.J. Santner, B.J. Williams, and W.J. Notz (2003), *The design and analysis of computer experiments*, Springer.

M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

**See Also**

[EI](#), [max\\_EI](#), [EI.grad](#)

**Examples**

```
set.seed(123)
#####
### 10 ITERATIONS OF EGO ON THE BRANIN FUNCTION,   ###
### STARTING FROM A 9-POINTS FACTORIAL DESIGN     ###
#####

# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# EGO n steps
library(rgenoud)
nsteps <- 5 # Was 10, reduced to 5 for speeding up compilation
lower <- rep(0,d)
upper <- rep(1,d)
oEGO <- EGO.nsteps(model=fitted.model1, fun=branin, nsteps=nsteps,
lower=lower, upper=upper, control=list(pop.size=20, BFGSburnin=2))
```

```

print(oEGO$par)
print(oEGO$value)

# graphics
n.grid <- 15 # Was 20, reduced to 15 for speeding up compilation
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, branin)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid, y.grid, z.grid, 40)
title("Branin function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par, pch=19, col="red")
text(oEGO$par[,1], oEGO$par[,2], labels=1:nsteps, pos=3)

#####
### 20 ITERATIONS OF EGO ON THE GOLDSTEIN-PRICE,      ###
### STARTING FROM A 9-POINTS FACTORIAL DESIGN        ###
#####
## Not run:
# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.goldsteinPrice <- apply(design.fact, 1, goldsteinPrice)
response.goldsteinPrice <- data.frame(response.goldsteinPrice)
names(response.goldsteinPrice) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.goldsteinPrice,
covtype="gauss", control=list(pop.size=50, max.generations=50,
wait.generations=5, BFGSburnin=10,trace=FALSE), parinit=c(0.5, 0.5), optim.method="BFGS")

# EGO n steps
library(rgenoud)
nsteps <- 10 # Was 20, reduced to 10 for speeding up compilation
lower <- rep(0,d)
upper <- rep(1,d)
oEGO <- EGO.nsteps(model=fitted.model1, fun=goldsteinPrice, nsteps=nsteps,
lower, upper, control=list(pop.size=20, BFGSburnin=2))
print(oEGO$par)
print(oEGO$value)

# graphics
n.grid <- 15 # Was 20, reduced to 15 for speeding up compilation
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, goldsteinPrice)
z.grid <- matrix(response.grid, n.grid, n.grid)

```

```

contour(x.grid, y.grid, z.grid, 40)
title("Goldstein-Price Function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par, pch=19, col="red")
text(oEGO$par[,1], oEGO$par[,2], labels=1:nsteps, pos=3)

## End(Not run)

#####
###  nsteps ITERATIONS OF EGO ON THE HARTMAN6 FUNCTION,      ###
###  STARTING FROM A 10-POINTS UNIFORM DESIGN                ###
#####

## Not run:
fonction<-hartman6
data(mydata)
a <- mydata
nb<-10
nsteps <- 3 # Maybe be changed to a larger value
x1<-a[[1]][1:nb];x2<-a[[2]][1:nb];x3<-a[[3]][1:nb]
x4<-a[[4]][1:nb];x5<-a[[5]][1:nb];x6<-a[[6]][1:nb]
design <- data.frame(cbind(x1,x2,x3,x4,x5,x6))
names(design)<-c("x1", "x2", "x3", "x4", "x5", "x6")
n <- nrow(design)

response <- data.frame(q=apply(design,1,fonction))
names(response) <- "y"
fitted.model1 <- km(~1, design=design, response=response, covtype="gauss",
control=list(pop.size=50, max.generations=20, wait.generations=5, BFGSburnin=5,
trace=FALSE), optim.method="gen", parinit=rep(0.8,6))

res.nsteps <- EGO.nsteps(model=fitted.model1, fun=fonction, nsteps=nsteps,
lower=rep(0,6), upper=rep(1,6), parinit=rep(0.5,6), control=list(pop.size=50,
max.generations=20, wait.generations=5, BFGSburnin=5), kmcontrol=NULL)
print(res.nsteps)
plot(res.nsteps$value, type="l")

## End(Not run)

```

**Description**

Computes the Expected Improvement at current location. The current minimum of the observations can be replaced by an arbitrary value (plugin), which is useful in particular in noisy frameworks.

**Usage**

```

EI(
  x,
  model,
  plugin = NULL,
  type = "UK",
  minimization = TRUE,
  envir = NULL,
  proxy = FALSE
)

```

**Arguments**

x	a vector representing the input for which one wishes to calculate EI,
model	an object of class <code>km</code> ,
plugin	optional scalar: if provided, it replaces the minimum of the current observations,
type	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account,
minimization	logical specifying if EI is used in minimization or in maximization,
envir	an optional environment specifying where to assign intermediate values for future gradient calculations. Default is NULL.
proxy	an optional Boolean, if TRUE EI is replaced by the kriging mean (to minimize)

**Value**

The expected improvement, defined as

$$EI(x) := E[(\min(Y(X)) - Y(x))^+ | Y(X) = y(X)],$$

where X is the current design of experiments and Y is the random process assumed to have generated the objective function y. If a plugin is specified, it replaces

$$\min(Y(X))$$

in the previous formula.

**Author(s)**

David Ginsbourger  
 Olivier Roustant  
 Victor Picheny

**References**

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.  
 J. Mockus (1988), *Bayesian Approach to Global Optimization*. Kluwer academic publishers.



T.J. Santner, B.J. Williams, and W.J. Notz (2003), *The design and analysis of computer experiments*, Springer.

M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

### See Also

[max\\_EI](#), [EG0.nsteps](#), [qEI](#)

### Examples

```
set.seed(123)
#####
###   EI SURFACE ASSOCIATED WITH AN ORDINARY KRIGING MODEL   ###
###   OF THE BRANIN FUNCTION KNOWN AT A 9-POINTS FACTORIAL DESIGN   ###
#####

# a 9-points factorial design, and the corresponding response
d <- 2; n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# graphics
n.grid <- 12
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
#response.grid <- apply(design.grid, 1, branin)
EI.grid <- apply(design.grid, 1, EI,fitted.model1)
z.grid <- matrix(EI.grid, n.grid, n.grid)
contour(x.grid,y.grid,z.grid,25)
title("Expected Improvement for the Branin function known at 9 points")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
```

### Description

Computes the gradient of the Expected Improvement at the current location. The current minimum of the observations can be replaced by an arbitrary value (plugin), which is useful in particular in noisy frameworks.

### Usage

```
EI.grad(  
  x,  
  model,  
  plugin = NULL,  
  type = "UK",  
  minimization = TRUE,  
  envir = NULL,  
  proxy = FALSE  
)
```

### Arguments

x	a vector representing the input for which one wishes to calculate <a href="#">EI</a> .
model	an object of class <a href="#">km</a> .
plugin	optional scalar: if provided, it replaces the minimum of the current observations,
type	Kriging type: "SK" or "UK"
minimization	logical specifying if EI is used in minimization or in maximization,
envir	an optional environment specifying where to get intermediate values calculated in <a href="#">EI</a> .
proxy	an optional Boolean, if TRUE EI is replaced by the kriging mean (to minimize)

### Value

The gradient of the expected improvement criterion with respect to x. Returns 0 at design points (where the gradient does not exist).

### Author(s)

David Ginsbourger  
Olivier Roustant  
Victor Picheny

### References

D. Ginsbourger (2009), *Multiplés metamodels pour l'approximation et l'optimisation de fonctions numériques multivariées*, Ph.D. thesis, Ecole Nationale Supérieure des Mines de Saint-Etienne, 2009.

J. Mockus (1988), *Bayesian Approach to Global Optimization*. Kluwer academic publishers.

T.J. Santner, B.J. Williams, and W.J. Notz (2003), *The design and analysis of computer experiments*, Springer.

M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

## See Also

[EI](#)

## Examples

```

set.seed(123)
# a 9-points factorial design, and the corresponding response
d <- 2; n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# graphics
n.grid <- 9 # Increase to 50 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
#response.grid <- apply(design.grid, 1, branin)
EI.grid <- apply(design.grid, 1, EI,fitted.model1)
#EI.grid <- apply(design.grid, 1, EI.plot,fitted.model1, gr=TRUE)

z.grid <- matrix(EI.grid, n.grid, n.grid)

contour(x.grid,y.grid,z.grid,20)
title("Expected Improvement for the Branin function known at 9 points")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")

# graphics
n.gridx <- 5 # increase to 15 for nicer picture
n.gridy <- 5 # increase to 15 for nicer picture
x.grid2 <- seq(0,1,length=n.gridx)
y.grid2 <- seq(0,1,length=n.gridy)
design.grid2 <- expand.grid(x.grid2, y.grid2)

EI.envir <- new.env()
environment(EI) <- environment(EI.grad) <- EI.envir

for(i in seq(1, nrow(design.grid2)) )

```

```

{
x <- design.grid2[i,]
ei <- EI(x, model=fitted.model1, envir=EI.envir)
eigrad <- EI.grad(x , model=fitted.model1, envir=EI.envir)
if(!is.null(ei))
{
suppressWarnings(arrows(x$Var1,x$Var2,
x$Var1 + eigrad[1]*2.2*10e-5, x$Var2 + eigrad[2]*2.2*10e-5,
length = 0.04, code=2, col="orange", lwd=2))
}
}

```

EQI

*Expected Quantile Improvement***Description**

Evaluation of the Expected Quantile Improvement (EQI) criterion.

**Usage**

```

EQI(
  x,
  model,
  new.noise.var = 0,
  beta = 0.9,
  q.min = NULL,
  type = "UK",
  envir = NULL
)

```

**Arguments**

x	the input vector at which one wants to evaluate the criterion
model	a Kriging model of "km" class
new.noise.var	(scalar) noise variance of the future observation. Default value is 0 (noise-free observation).
beta	Quantile level (default value is 0.9)
q.min	Best kriging quantile. If not provided, this quantity is evaluated.
type	Kriging type: "SK" or "UK"
envir	environment for saving intermediate calculations and reusing them within EQI.grad

**Value**

Expected Quantile Improvement

**Author(s)**

Victor Picheny

David Ginsbourger

**References**

Picheny, V., Ginsbourger, D., Richet, Y., Caplin, G. (2013). Quantile-based optimization of noisy computer experiments with tunable precision. *Technometrics*, 55(1), 2-13.

**Examples**

```
#####
###   EQI SURFACE ASSOCIATED WITH AN ORDINARY KRIGING MODEL       ###
### OF THE BRANIN FUNCTION KNOWN AT A 12-POINT LATIN HYPERCUBE DESIGN ###
#####

set.seed(421)

# Set test problem parameters
doe.size <- 12
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {
  y.tilde[i] <- test.function(doe[i,]) + sqrt(noise.var)*rnorm(n=1)
}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Compute actual function and criterion on a grid
n.grid <- 12 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
nt <- nrow(design.grid)

crit.grid <- apply(design.grid, 1, EQI, model=model, new.noise.var=noise.var, beta=.9)
func.grid <- apply(design.grid, 1, test.function)

# Compute kriging mean and variance on a grid
```

```

names(design.grid) <- c("V1","V2")
pred <- predict(model, newdata=design.grid, type="UK", checkNames = FALSE)
mk.grid <- pred$m
sk.grid <- pred$sd

# Plot actual function
z.grid <- matrix(func.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Actual function");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot Kriging mean
z.grid <- matrix(mk.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Kriging mean");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot Kriging variance
z.grid <- matrix(sk.grid^2, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Kriging variance");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot EQI criterion
z.grid <- matrix(crit.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("EQI");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

```

---

EQI.grad

*EQI's Gradient*


---

### Description

Analytical gradient of the Expected Quantile Improvement (EQI) criterion.

### Usage

```

EQI.grad(
  x,
  model,
  new.noise.var = 0,
  beta = 0.9,
  q.min = NULL,
  type = "UK",

```

```

    envir = NULL
  )

```

### Arguments

x	the input vector at which one wants to evaluate the criterion
model	a Kriging model of "km" class
new.noise.var	(scalar) noise variance of the future observation. Default value is 0 (noise-free observation).
beta	Quantile level (default value is 0.9)
q.min	Best kriging quantile. If not provided, this quantity is evaluated.
type	Kriging type: "SK" or "UK"
envir	environment for inheriting intermediate calculations from EQI

### Value

Gradient of the Expected Quantile Improvement

### Author(s)

Victor Picheny  
David Ginsbourger

### Examples

```

set.seed(421)

# Set test problem parameters
doe.size <- 12
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {
  y.tilde[i] <- test.function(doe[i,]) + sqrt(noise.var)*rnorm(n=1)
}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

```

```

# Compute actual function and criterion on a grid
n.grid <- 9 # change to 21 for nicer visuals
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
nt <- nrow(design.grid)

crit.grid <- apply(design.grid, 1, EQI, model=model, new.noise.var=noise.var, beta=.9)
crit.grad <- t(apply(design.grid, 1, EQI.grad, model=model, new.noise.var=noise.var, beta=.9))

z.grid <- matrix(crit.grid, n.grid, n.grid)
contour(x.grid,y.grid, z.grid, 30)
title("EQI and its gradient")
points(model@X[,1],model@X[,2],pch=17,col="blue")

for (i in 1:nt)
{
  x <- design.grid[i,]
  suppressWarnings(arrows(x$Var1,x$Var2, x$Var1+crit.grad[i,1]*.2,x$Var2+crit.grad[i,2]*.2,
length=0.04,code=2,col="orange",lwd=2))
}

```

---

fastEGO.nsteps	<i>Sequential EI maximization and model re-estimation, with a number of iterations fixed in advance by the user</i>
----------------	---

---

## Description

Executes *nsteps* iterations of the EGO method to an object of class `km`. At each step, a kriging model is 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 the Expected Improvement criterion (EI).

## Usage

```

fastEGO.nsteps(
  model,
  fun,
  nsteps,
  lower,
  upper,
  control = NULL,
  trace = 0,
  n.cores = 1,
  ...
)

```



**Arguments**

model	an object of class <code>km</code> ,
fun	the objective function to be minimized,
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,
control	an optional list of control parameters for EGO. One can control "warping" whether or not a warping is applied to the outputs (default FALSE) "cov.reestim" whether or not the covariance parameters are estimated at each step (default TRUE) "gpmean.trick" whether or not EI should be replaced periodically by the GP mean (default FALSE) "gpmean.freq" frequency at which EI is replaced by the GP mean (default 1e4) "always.sample" if TRUE, forces observation even if it creates poor conditioning
trace	between -1 (no trace) and 3 (full messages)
n.cores	number of cores used for EI maximisation
...	additional parameters to be given to fun

**Value**

A list with components:

par	a data frame representing the additional points visited during the algorithm,
value	a data frame representing the response values at the points given in par,
npoints	an integer representing the number of parallel computations (=1 here),
nsteps	an integer representing the desired number of iterations (given in argument),
lastmodel	an object of class <code>km</code> corresponding to the last kriging model fitted. If warping is true, y values are normalized (warped) and will not match value.

**Author(s)**

Victor Picheny

**References**

- 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.
- J. Mockus (1988), *Bayesian Approach to Global Optimization*. Kluwer academic publishers.
- T.J. Santner, B.J. Williams, and W.J. Notz (2003), *The design and analysis of computer experiments*, Springer.
- M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

**See Also**

[EI](#), [max\\_crit](#), [EI.grad](#)

**Examples**

```

set.seed(123)
#####
### 10 ITERATIONS OF EGO ON THE BRANIN FUNCTION, ###
### STARTING FROM A 9-POINTS FACTORIAL DESIGN ###
#####

# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# EGO n steps
nsteps <- 5
lower <- rep(0,d)
upper <- rep(1,d)
oEGO <- fastEGO.nsteps(model=fitted.model1, fun=branin, nsteps=nsteps, lower=lower, upper=upper)
print(oEGO$par)
print(oEGO$value)

# graphics
n.grid <- 15 # Was 20, reduced to 15 for speeding up compilation
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, branin)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid, y.grid, z.grid, 40)
title("Branin function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par, pch=19, col="red")
text(oEGO$par[,1], oEGO$par[,2], labels=1:nsteps, pos=3)

```

**Description**

Modification of an R function to be used as 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`.

---

```
integration_design_cst
```

*Generic 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 with constraints.

**Usage**

```
integration_design_cst(
  integcontrol = NULL,
  lower,
  upper,
  model.fun = NULL,
  model.constraint = NULL,
  equality = FALSE,
  critcontrol = NULL,
  min.prob = 0.001
)
```

**Arguments**

<code>integcontrol</code>	<p>Optional list specifying the procedure to build the integration points and weights. Many options are possible.</p> <p>A) If nothing is specified, <math>100*d</math> points are chosen using the Sobol sequence.</p> <p>B) One can directly set the field <code>integration.points</code> (<math>p * d</math> matrix) for pre-specified integration points. In this case these integration points and the corresponding vector <code>integration.weights</code> will be used for all the iterations of the algorithm.</p> <p>C) If the field <code>integration.points</code> is not set then the integration points are renewed at each iteration. In that case one can control the number of integration points <code>n.points</code> (default: <math>100*d</math>) and a specific distribution <code>distrib</code>. Possible values for <code>distrib</code> are: "sobol", "MC" and "SUR" (default: "sobol").</p> <p>C.1) The choice "sobol" corresponds to integration points chosen with the Sobol sequence in dimension <math>d</math> (uniform weight).</p> <p>C.2) The choice "MC" corresponds to points chosen randomly, uniformly on the domain.</p> <p>C.3) The choice "SUR" corresponds to importance sampling distributions (unequal weights).</p> <p>When important sampling procedures are chosen, <code>n.points</code> points are chosen using importance sampling among a discrete set of <code>n.candidates</code> points (default: <math>n.points*10</math>) which are distributed according to a distribution <code>init.distrib</code> (default: "sobol"). Possible values for <code>init.distrib</code> 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 <code>init.distrib.spec</code> to specify himself a <math>n.candidates * d</math> matrix of points in dimension <math>d</math>.</p>
<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.fun</code>	object of class <code>km</code> corresponding to the objective functions, or, if the objective function is fast-to-evaluate, a <code>fastfun</code> object,
<code>model.constraint</code>	either one or a list of objects of class <code>km</code> , one for each constraint function,
<code>equality</code>	either FALSE if all constraints are for inequalities, else a vector of boolean indicating which are equalities
<code>critcontrol</code>	optional list of parameters (see <code>crit_SUR_cst</code> ); here only the component <code>tolConstraints</code> is used.
<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 $M$ points, the maximum weight becomes $1/\text{min.prob} * 1/M$ .

**Value**

A list with components:

- `integration.points`  $p \times d$  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`

### Author(s)

Victor Picheny  
Mickael Binois

### References

Chevalier C., Picheny V., Ginsbourger D. (2012), The KrigInv package: An efficient and user-friendly R implementation of Kriging-based inversion algorithms, *Computational Statistics and Data Analysis*, 71, 1021-1034.

Chevalier C., Bect J., Ginsbourger D., Vazquez E., Picheny V., Richet Y. (2011), Fast parallel kriging-based stepwise uncertainty reduction with application to the identification of an excursion set, *Technometrics*, 56(4), 455-465.

V. Picheny (2014), A stepwise uncertainty reduction approach to constrained global optimization, *Proceedings of the 17th International Conference on Artificial Intelligence and Statistics*, JMLR W&CP 33, 787-795.

### See Also

[crit\\_SUR\\_cst](#) KrigInv [integration\\_design](#)

---

<code>kriging.quantile</code>	<i>Kriging quantile</i>
-------------------------------	-------------------------

---

### Description

Evaluation of a kriging quantile at a new point. To be used in an optimization loop.

### Usage

```
kriging.quantile(x, model, beta = 0.1, type = "UK", envir = NULL)
```

### Arguments

<code>x</code>	the input vector at which one wants to evaluate the criterion
<code>model</code>	a Kriging model of "km" class
<code>beta</code>	Quantile level (default value is 0.1)
<code>type</code>	Kriging type: "SK" or "UK"
<code>envir</code>	an optional environment specifying where to assign intermediate values for future gradient calculations. Default is <code>NULL</code> .

**Value**

Kriging quantile

**Author(s)**

Victor Picheny

David Ginsbourger

**Examples**

```
#####
###   KRIGING QUANTILE SURFACE                               #####
### OF THE BRANIN FUNCTION KNOWN AT A 12-POINT LATIN HYPERCUBE DESIGN #####
#####

set.seed(421)

# Set test problem parameters
doe.size <- 12
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {
  y.tilde[i] <- test.function(doe[i,]) + sqrt(noise.var)*rnorm(n=1)
}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Compute actual function and criterion on a grid
n.grid <- 12 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
nt <- nrow(design.grid)

crit.grid <- apply(design.grid, 1, kriging.quantile, model=model, beta=.1)
func.grid <- apply(design.grid, 1, test.function)

# Compute kriging mean and variance on a grid
names(design.grid) <- c("V1","V2")
pred <- predict(model, newdata=design.grid, type="UK", checkNames = FALSE)
```

```

mk.grid <- pred$m
sk.grid <- pred$sd

# Plot actual function
z.grid <- matrix(func.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Actual function");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot Kriging mean
z.grid <- matrix(mk.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Kriging mean");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot Kriging variance
z.grid <- matrix(sk.grid^2, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("Kriging variance");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

# Plot kriging.quantile criterion
z.grid <- matrix(crit.grid, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title("kriging.quantile");
points(model@X[,1],model@X[,2],pch=17,col="blue");
axis(1); axis(2)})

```

---

kriging.quantile.grad *Analytical gradient of the Kriging quantile of level beta*

---

### Description

Computes the gradient of the Kriging quantile of level beta at the current location. Only available for Universal Kriging with constant trend (Ordinary Kriging).

### Usage

```
kriging.quantile.grad(x, model, beta = 0.1, type = "UK", envir = NULL)
```

### Arguments

x	a vector representing the input for which one wishes to calculate kriging.quantile.grad.
model	an object of class <a href="#">km</a> .
beta	A quantile level (between 0 and 1)

type           Kriging type: "SK" or "UK"  
 envir         environment for inheriting intermediate calculations from "kriging.quantile"

### Value

The gradient of the Kriging mean predictor with respect to x. Returns 0 at design points (where the gradient does not exist).

### Author(s)

Victor Picheny  
 David Ginsbourger

### References

O. Roustant, D. Ginsbourger, Y. Deville, *DiceKriging, DiceOptim: Two R packages for the analysis of computer experiments by kriging-based metamodeling and optimization*, J. Stat. Soft., 2010. <https://www.jstatsoft.org/article/view/v051i01>

D. Ginsbourger (2009), *Multiplés metamodelés pour l'approximation et l'optimisation de fonctions numériques multivariées*, Ph.D. thesis, Ecole Nationale Supérieure des Mines de Saint-Etienne, 2009.

### See Also

[EI.grad](#)

### Examples

```
#####
###   KRIGING QUANTILE SURFACE AND ITS GRADIENT FOR   ###
###   THE BRANIN FUNCTION KNOWN AT A 12-POINT LATIN HYPERCUBE DESIGN   ###
#####
set.seed(421)

# Set test problem parameters
doe.size <- 12
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {
  y.tilde[i] <- test.function(doe[i,]) + sqrt(noise.var)*rnorm(n=1)
}
y.tilde <- as.numeric(y.tilde)
```



```

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Compute actual function and criterion on a grid
n.grid <- 9 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
nt <- nrow(design.grid)

crit.grid <- apply(design.grid, 1, kriging.quantile, model=model, beta=.1)
crit.grad <- t(apply(design.grid, 1, kriging.quantile.grad, model=model, beta=.1))

z.grid <- matrix(crit.grid, n.grid, n.grid)
contour(x.grid,y.grid, z.grid, 30)
title("kriging.quantile and its gradient")
points(model@X[,1],model@X[,2],pch=17,col="blue")

for (i in 1:nt)
{
  x <- design.grid[i,]
  arrows(x$Var1,x$Var2, x$Var1+crit.grad[i,1]*.01,x$Var2+crit.grad[i,2]*.01,
        length=0.04,code=2,col="orange",lwd=2)
}

```

---

max\_AEI

---

*Maximizer of the Augmented Expected Improvement criterion function*


---

## Description

Maximization, based on the package rgenoud of the Augmented Expected Improvement (AEI) criterion.

## Usage

```

max_AEI(
  model,
  new.noise.var = 0,
  y.min = NULL,
  type = "UK",
  lower,
  upper,
  parinit = NULL,
  control = NULL
)

```

**Arguments**

model	a Kriging model of "km" class
new.noise.var	the (scalar) noise variance of the new observation.
y.min	The kriging mean prediction at the current best point (point with smallest kriging quantile). If not provided, this quantity is evaluated inside the AEI function (may increase computational time).
type	Kriging type: "SK" or "UK"
lower	vector containing the lower bounds of the variables to be optimized over
upper	optional vector containing the upper bounds of the variables to be optimized over
parinit	optional vector containing the initial values for the variables to be optimized over
control	optional list of control parameters for optimization. One can control "pop.size" (default : [N=3*2^dim for dim<6 and N=32*dim otherwise]), "max.generations" (12), "wait.generations" (2) and "BFGSburnin" (2) of function "genoud" (see <a href="#">genoud</a> ). Numbers into brackets are the default values

**Value**

A list with components:

par	the best set of parameters found.
value	the value AEI at par.

**Author(s)**

Victor Picheny

David Ginsbourger

**Examples**

```
library(DiceDesign)
set.seed(100)

# Set test problem parameters
doe.size <- 10
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(lhsDesign(doe.size, dim)$design)
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {y.tilde[i] <- test.function(doe[i,])}
```

```

+ sqrt(noise.var)*rnorm(n=1)}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
  covtype="gauss", noise.var=rep(noise.var,1,doe.size),
  lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Optimisation using max_AEI
res <- max_AEI(model, new.noise.var=noise.var, type = "UK",
  lower=c(0,0), upper=c(1,1))
X.genoud <- res$par

# Compute actual function and criterion on a grid
n.grid <- 12 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
names(design.grid) <- c("V1","V2")
nt <- nrow(design.grid)
crit.grid <- apply(design.grid, 1, AEI, model=model, new.noise.var=noise.var)

## Not run:
# # 2D plots
z.grid <- matrix(crit.grid, n.grid, n.grid)
tit <- "Green: best point found by optimizer"
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
  plot.axes = {title(tit);points(model@X[,1],model@X[,2],pch=17,col="blue");
  points(X.genoud[1],X.genoud[2],pch=17,col="green");
  axis(1); axis(2)})

## End(Not run)

```

---

max\_AKG

*Maximizer of the Expected Quantile Improvement criterion function*


---

## Description

Maximization, based on the package rgenoud of the Expected Quantile Improvement (AKG) criterion.

## Usage

```

max_AKG(
  model,
  new.noise.var = 0,
  type = "UK",
  lower,
  upper,
  parinit = NULL,

```

```

    control = NULL
  )

```

### Arguments

model	a Kriging model of "km" class
new.noise.var	the (scalar) noise variance of an observation. Default value is 0 (noise-free observation).
type	Kriging type: "SK" or "UK"
lower	vector containing the lower bounds of the variables to be optimized over
upper	vector containing the upper bounds of the variables to be optimized over
parinit	optional vector containing the initial values for the variables to be optimized over
control	optional list of control parameters for optimization. One can control "pop.size" (default : $[N=3*2^{\dim}$ for $\dim < 6$ and $N=32*\dim$ otherwise]), "max.generations" (12), "wait.generations" (2) and "BFGSburnin" (2) of function "genoud" (see <a href="#">genoud</a> ). Numbers into brackets are the default values

### Value

A list with components:

par	the best set of parameters found.
value	the value AKG at par.

### Author(s)

Victor Picheny  
David Ginsbourger

### Examples

```

#####
###   AKG SURFACE AND OPTIMIZATION PERFORMED BY GENOUD           #####
###   FOR AN ORDINARY KRIGING MODEL                             #####
###   OF THE BRANIN FUNCTION KNOWN AT A 12-POINT LATIN HYPERCUBE DESIGN #####
#####
set.seed(10)

# Set test problem parameters
doe.size <- 10
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response

```

```

library(DiceDesign)
doe <- as.data.frame(lhsDesign(doe.size, dim)$design)
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {y.tilde[i] <- test.function(doe[i,])
+ sqrt(noise.var)*rnorm(n=1)}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
  covtype="gauss", noise.var=rep(noise.var,1,doe.size),
  lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Optimisation using max_AKG
res <- max_AKG(model, new.noise.var=noise.var, type = "UK",
  lower=c(0,0), upper=c(1,1))
X.genoud <- res$par

## Not run:
# Compute actual function and criterion on a grid
n.grid <- 12 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
names(design.grid) <- c("V1","V2")
nt <- nrow(design.grid)
crit.grid <- apply(design.grid, 1, AKG, model=model, new.noise.var=noise.var)

# # 2D plots
z.grid <- matrix(crit.grid, n.grid, n.grid)
tit <- "Green: best point found by optimizer"
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
  plot.axes = {title(tit);points(model@X[,1],model@X[,2],pch=17,col="blue");
  points(X.genoud[1],X.genoud[2],pch=17,col="green");
  axis(1); axis(2)})

## End(Not run)

```

---

max\_crit

*Maximization of the Expected Improvement criterion*


---

## Description

For a number of control\$restarts, generates a large number of random samples, then picks the one with best EI value to start L-BFGS.

## Usage

```

max_crit(
  model,
  type = "UK",

```

```

    lower,
    upper,
    minimization = TRUE,
    control = NULL,
    proxy = FALSE,
    trcontrol = NULL,
    n.cores = 1
  )

```

### Arguments

model	an object of class <code>km</code> ,
type	Kriging type: "SK" or "UK"
lower, upper	vectors of lower and upper bounds for the variables to be optimized over,
minimization	logical specifying if EI is used in minimization or in maximization,
control	optional list of control parameters for optimization. For now only the number of restarts can be set.
proxy	Boolean, if TRUE, then EI maximization is replaced by the minimization of the kriging mean.
trcontrol	an optional list to activate the Trust-region management (see <code>TREGO.nsteps</code> )
n.cores	Number of cores if parallel computation is used

### Value

A list with components:

par	The best set of parameters found.
value	The value of expected improvement at par.

### Author(s)

Victor Picheny

### Examples

```

set.seed(123)
library(parallel)
#####
### "ONE-SHOT" EI-MAXIMIZATION OF THE BRANIN FUNCTION ###
### KNOWN AT A 9-POINTS FACTORIAL DESIGN          ###
#####

# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact) <- c("x1", "x2")

```

```

design.fact <- data.frame(design.fact)
names(design.fact) <- c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# EGO one step
lower <- rep(0,d)
upper <- rep(1,d)      # domain for Branin function
oEGO <- max_crit(fitted.model1, lower=lower, upper=upper)
print(oEGO)

# graphics
n.grid <- 20
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, branin)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid,y.grid,z.grid,40)
title("Branin Function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par[1], oEGO$par[2], pch=19, col="red")

```

---

max\_EI

*Maximization of the Expected Improvement criterion*


---

## Description

Given an object of class `km` and a set of tuning parameters (`lower`, `upper`, `parinit`, and `control`), `max_EI` performs the maximization of the Expected Improvement criterion and delivers the next point to be visited in an EGO-like procedure.

## Usage

```

max_EI(
  model,
  plugin = NULL,
  type = "UK",
  lower,
  upper,
  parinit = NULL,
  minimization = TRUE,
  control = NULL
)

```

**Arguments**

model	an object of class <code>km</code> ,
plugin	optional scalar: if provided, it replaces the minimum of the current observations,
type	Kriging type: "SK" or "UK"
lower	vector of lower bounds for the variables to be optimized over,
upper	vector of upper bounds for the variables to be optimized over,
parinit	optional vector of initial values for the variables to be optimized over,
minimization	logical specifying if EI is used in minimization or in maximization,
control	optional list of control parameters for optimization. One can control "pop.size" (default : $[N=3*2^{\dim}$ for $\dim < 6$ and $N=32*\dim$ otherwise]), "max.generations" (12), "wait.generations" (2) and "BFGSburnin" (2) of function "genoud" (see <code>genoud</code> ). Numbers into brackets are the default values

**Details**

The latter maximization relies on a genetic algorithm using derivatives, `genoud`. This function plays a central role in the package since it is in constant use in the proposed algorithms. It is important to remark that the information needed about the objective function reduces here to the vector of response values embedded in `model` (no call to the objective function or simulator).

The current minimum of the observations can be replaced by an arbitrary value (`plugin`), which is useful in particular in noisy frameworks.

**Value**

A list with components:

par	The best set of parameters found.
value	The value of expected improvement at par.

**Author(s)**

David Ginsbourger

Olivier Roustant

Victor Picheny

**References**

D. Ginsbourger (2009), *Multiplés metamodeles pour l'approximation et l'optimisation de fonctions numeriques multivariables*, Ph.D. thesis, Ecole Nationale Superieure des Mines de Saint-Etienne, 2009.

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.

W.R. Jr. Mebane and J.S. Sekhon (2009), in press, Genetic optimization using derivatives: The `genoud` package for R, *Journal of Statistical Software*.



## Examples

```

set.seed(123)
#####
### "ONE-SHOT" EI-MAXIMIZATION OF THE BRANIN FUNCTION ###
### KNOWN AT A 9-POINTS FACTORIAL DESIGN          ###
#####

# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact) <- c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact) <- c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# EGO one step
library(rgenoud)
lower <- rep(0,d)
upper <- rep(1,d)      # domain for Branin function
oEGO <- max_EI(fitted.model1, lower=lower, upper=upper,
control=list(pop.size=20, BFGSburnin=2))
print(oEGO)

# graphics
n.grid <- 20
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, branin)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid,y.grid,z.grid,40)
title("Branin Function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par[1], oEGO$par[2], pch=19, col="red")

#####
### "ONE-SHOT" EI-MAXIMIZATION OF THE CAMELBACK FUNCTION ###
### KNOWN AT A 16-POINTS FACTORIAL DESIGN          ###
#####
## Not run:
# a 16-points factorial design, and the corresponding response
d <- 2
n <- 16
design.fact <- expand.grid(seq(0,1,length=4), seq(0,1,length=4))

```

```

names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact) <- c("x1", "x2")
response.camelback <- apply(design.fact, 1, camelback)
response.camelback <- data.frame(response.camelback)
names(response.camelback) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.camelback,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# EI maximization
library(rgenoud)
lower <- rep(0,d)
upper <- rep(1,d)
oEGO <- max_EI(fitted.model1, lower=lower, upper=upper,
control=list(pop.size=20, BFGSburnin=2))
print(oEGO)

# graphics
n.grid <- 20
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, camelback)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid,y.grid,z.grid,40)
title("Camelback Function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par[1], oEGO$par[2], pch=19, col="red")

## End(Not run)

#####
### "ONE-SHOT" EI-MAXIMIZATION OF THE GOLDSTEIN-PRICE FUNCTION #####
###          KNOWN AT A 9-POINTS FACTORIAL DESIGN          #####
#####

## Not run:
# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.goldsteinPrice <- apply(design.fact, 1, goldsteinPrice)
response.goldsteinPrice <- data.frame(response.goldsteinPrice)
names(response.goldsteinPrice) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.goldsteinPrice,
covtype="gauss", control=list(pop.size=50, max.generations=50,
wait.generations=5, BFGSburnin=10, trace=FALSE), parinit=c(0.5, 0.5), optim.method="gen")

```

```

# EI maximization
library(rgenoud)
lower <- rep(0,d); upper <- rep(1,d); # domain for Branin function
oEGO <- max_EQI(fitted.model1, lower=lower, upper=upper, control
=list(pop.size=50, max.generations=50, wait.generations=5, BFGSburnin=10))
print(oEGO)

# graphics
n.grid <- 20
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, goldsteinPrice)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid,y.grid,z.grid,40)
title("Goldstein-Price Function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par[1], oEGO$par[2], pch=19, col="red")

## End(Not run)

```

---

max\_EQI

*Maximizer of the Expected Quantile Improvement criterion function*


---

### Description

Maximization, based on the package `rgenoud` of the Expected Quantile Improvement (EQI) criterion.

### Usage

```

max_EQI(
  model,
  new.noise.var = 0,
  beta = 0.9,
  q.min = NULL,
  type = "UK",
  lower,
  upper,
  parinit = NULL,
  control = NULL
)

```

### Arguments

<code>model</code>	a Kriging model of "km" class
<code>new.noise.var</code>	the (scalar) noise variance of an observation. Default value is 0 (noise-free observation).

beta	Quantile level (default value is 0.9)
q.min	The current best kriging quantile. If not provided, this quantity is evaluated inside the EQI function (may increase computational time).
type	Kriging type: "SK" or "UK"
lower	vector containing the lower bounds of the variables to be optimized over
upper	optional vector containing the upper bounds of the variables to be optimized over
parinit	optional vector containing the initial values for the variables to be optimized over
control	optional list of control parameters for optimization. One can control "pop.size" (default: $[N=3*2^{\dim}$ for $\dim < 6$ and $N=32*\dim$ otherwise]), "max.generations" (12), "wait.generations" (2) and "BFGSburnin" (2) of function "genoud" (see <a href="#">genoud</a> ). Numbers into brackets are the default values

### Value

A list with components:

par	the best set of parameters found.
value	the value EQI at par.

### Author(s)

Victor Picheny  
David Ginsbourger

### Examples

```
set.seed(10)

# Set test problem parameters
doe.size <- 10
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {y.tilde[i] <- test.function(doe[i,])
+ sqrt(noise.var)*rnorm(n=1)}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
covtype="gauss", noise.var=rep(noise.var,1,doe.size),
```

```

        lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Optimisation using max_EQI
res <- max_EQI(model, new.noise.var=noise.var, type = "UK",
lower=c(0,0), upper=c(1,1))
X.genoud <- res$par

## Not run:
# Compute actual function and criterion on a grid
n.grid <- 12 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
names(design.grid) <- c("V1","V2")
nt <- nrow(design.grid)
crit.grid <- apply(design.grid, 1, EQI, model=model, new.noise.var=noise.var, beta=.9)

# # 2D plots
z.grid <- matrix(crit.grid, n.grid, n.grid)
tit <- "Green: best point found by optimizer"
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = rainbow,
plot.axes = {title(tit);points(model@X[,1],model@X[,2],pch=17,col="blue");
points(X.genoud[1],X.genoud[2],pch=17,col="green");
axis(1); axis(2)})

## End(Not run)

```

---

max\_qEI

*Maximization of multipoint expected improvement criterion (qEI)*


---

## Description

Maximization of the qEI criterion. Two options are available : Constant Liar (CL), and brute force qEI maximization with Broyden-Fletcher-Goldfarb-Shanno (BFGS) algorithm, or GENetic Optimization Using Derivative (genoud) algorithm.

## Usage

```

max_qEI(
  model,
  npoints,
  lower,
  upper,
  crit = "exact",
  minimization = TRUE,
  optimcontrol = NULL
)

```

**Arguments**

model	an object of class <code>km</code> ,
npoints	an integer representing the desired number of iterations,
lower	vector of lower bounds,
upper	vector of upper bounds,
crit	"exact", "CL" : a string specifying the criterion used. "exact" triggers the maximization of the multipoint expected improvement at each iteration (see <code>max_qEI</code> ), "CL" applies the Constant Liar heuristic,
minimization	logical specifying if the qEI to be maximized is used in minimization or in maximization,
optimcontrol	an optional list of control parameters for optimization. See details.

**Details**

- CL is a heuristic method. First, the regular Expected Improvement EI is maximized (`max_EI`). Then, for the next points, the Expected Improvement is maximized again, but with an artificially updated Kriging model. Since the response values corresponding to the last best point obtained are not available, the idea of CL is to replace them by an arbitrary constant value L (a "lie") set by the user (default is the minimum of all currently available observations).

- The BFGS algorithm is implemented in the standard function `optim`. Analytical formulae of `qEI` and its gradient `qEI.grad` are used. The `nStarts` starting points are by default sampled with respect to the regular EI (`sampleFromEI`) criterion.

- The "genoud" method calls the function `genoud` using analytical formulae of `qEI` and its gradient `qEI.grad`.

The parameters of list `optimcontrol` are :

- `optimcontrol$method` : "BFGS" (default), "genoud" ; a string specifying the method used to maximize the criterion (irrelevant when `crit` is "CL" because this method always uses `genoud`),

- when `crit="CL"` :

+ `optimcontrol$parinit` : optional matrix of initial values (must have `model@d` columns, the number of rows is not constrained),

+ `optimcontrol$L` : "max", "min", "mean" or a scalar value specifying the liar ; "min" takes `model@min`, "max" takes `model@max`, "mean" takes the prediction of the model ; When L is NULL, "min" is taken if `minimization==TRUE`, else it is "max".

+ The parameters of function `genoud`. Main parameters are : "pop.size" (default :  $[N=3*2^{\text{model@d}}$  for  $\text{dim}<6$  and  $N=32*\text{model@d}$  otherwise]), "max.generations" (default : 12), "wait.generations" (default : 2) and "BFGSburnin" (default : 2).

- when `optimcontrol$method = "BFGS"` :

+ `optimcontrol$nStarts` (default : 4),

+ `optimcontrol$fastCompute` : if TRUE (default), a fast approximation method based on a semi-analytic formula is used, see [Marmin 2014] for details,

+ `optimcontrol$samplingFun` : a function which sample a batch of starting point (default : `sampleFromEI`),

+ `optimcontrol$parinit` : optional 3d-array of initial (or candidate) batches (for all `k`, `parinit[,k]` is a matrix of size `npoints*model@d` representing one batch). The number of initial batches (`length(parinit[1,1])`) is not constrained and does not have to be equal to `nStarts`. If there is too few initial batches for `nStarts`, missing batches are drawn with `samplingFun` (default : NULL),

- when `optimcontrol$method = "genoud"` :

+ `optimcontrol$fastCompute` : if TRUE (default), a fast approximation method based on a semi-analytic formula is used, see [Marmin 2014] for details,

+ `optimcontrol$parinit` : optional matrix of candidate starting points (one row corresponds to one point),

+ The parameters of the [genoud](#) function. Main parameters are "pop.size" (default : `[50*(model@d)*(npoints)]`), "max.generations" (default : 5), "wait.generations" (default : 2), "BFGSburnin" (default : 2).

### Value

A list with components:

<code>par</code>	A matrix containing the <code>npoints</code> input vectors found.
<code>value</code>	A value giving the qEI computed in <code>par</code> .

### Author(s)

Sebastien Marmin

Clement Chevalier

David Ginsbourger

### References

C. Chevalier and D. Ginsbourger (2014) Learning and Intelligent Optimization - 7th International Conference, Lion 7, Catania, Italy, January 7-11, 2013, Revised Selected Papers, chapter Fast computation of the multipoint Expected Improvement with applications in batch selection, pages 59-69, Springer.

D. Ginsbourger, R. Le Riche, L. Carraro (2007), A Multipoint Criterion for Deterministic Parallel Global Optimization based on Kriging. The International Conference on Non Convex Programming, 2007.

D. Ginsbourger, R. Le Riche, and L. Carraro. Kriging is well-suited to parallelize optimization (2010), In Lim Meng Hiot, Yew Soon Ong, Yoel Tenne, and Chi-Keong Goh, editors, *Computational Intelligence in Expensive Optimization Problems*, Adaptation Learning and Optimization, pages 131-162. Springer Berlin Heidelberg.

J. Mockus (1988), *Bayesian Approach to Global Optimization*. Kluwer academic publishers.

M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

### See Also

[qEI](#), [qEI.grad](#)

**Examples**

```

set.seed(000)
# 3-points EI maximization.
# 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"
lower <- c(0,0)
upper <- c(1,1)

# number of point in the batch
batchSize <- 3

# model identification
fitted.model <- km(~1, design=design.fact, response=response.branin,
                  covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# maximization of qEI

# With a multistarted BFGS algorithm
maxBFGS <- max_qEI(model = fitted.model, npoints = batchSize, lower = lower, upper = upper,
                  crit = "exact",optimcontrol=list(nStarts=3,method = "BFGS"))

# comparison
print(maxBFGS$value)
## Not run:
# With a genetic algorithm using derivatives
maxGen <- max_qEI(model = fitted.model, npoints = batchSize, lower = lower, upper = upper,
                  crit = "exact", optimcontrol=list(nStarts=3,method = "genoud",pop.size=100,max.generations = 15))
# With the constant liar heuristic
maxCL <- max_qEI(model = fitted.model, npoints = batchSize, lower = lower, upper = upper,
                  crit = "CL",optimcontrol=list(pop.size=20))
print(maxGen$value)
print(maxCL$value)

## End(Not run)

```



**Description**

Minimization, based on the package rgenoud of the kriging quantile.

**Usage**

```
min_quantile(
  model,
  beta = 0.1,
  type = "UK",
  lower,
  upper,
  parinit = NULL,
  control = NULL
)
```

**Arguments**

model	a Kriging model of "km" class
beta	Quantile level (default value is 0.1)
type	Kriging type: "SK" or "UK"
lower	vector containing the lower bounds of the variables to be optimized over
upper	vector containing the upper bounds of the variables to be optimized over
parinit	optional vector containing the initial values for the variables to be optimized over
control	optional list of control parameters for optimization. One can control "pop.size" (default : [N=3*2^dim for dim<6 and N=32*dim otherwise]), "max.generations" (12), "wait.generations" (2) and "BFGSburnin" (2) of function "genoud" (see <a href="#">genoud</a> ). Numbers into brackets are the default values

**Value**

A list with components:

par	the best set of parameters found.
value	the value of the kriging quantile at par.

**Author(s)**

Victor Picheny  
David Ginsbourger

**Examples**

```
#####
### KRIGING QUANTILE SURFACE AND OPTIMIZATION PERFORMED BY GENOUD ###
```

```

###   FOR AN ORDINARY KRIGING MODEL                                     #####
### OF THE BRANIN FUNCTION KNOWN AT A 12-POINT LATIN HYPERCUBE DESIGN #####
#####
set.seed(10)

# Set test problem parameters
doe.size <- 10
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.2

# Generate DOE and response
doe <- as.data.frame(matrix(runif(doe.size*dim),doe.size))
y.tilde <- rep(0, 1, doe.size)
for (i in 1:doe.size) {y.tilde[i] <- test.function(doe[i,])
+ sqrt(noise.var)*rnorm(n=1)}
y.tilde <- as.numeric(y.tilde)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
           covtype="gauss", noise.var=rep(noise.var,1,doe.size),
           lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Optimisation using max_kriging.quantile
res <- min_quantile(model, beta=0.1, type = "UK", lower=c(0,0), upper=c(1,1))
X.genoud <- res$par

# Compute actual function and criterion on a grid
n.grid <- 12 # Change to 21 for a nicer picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
names(design.grid) <- c("V1","V2")
nt <- nrow(design.grid)
crit.grid <- apply(design.grid, 1, kriging.quantile, model=model, beta=.1)

# # 2D plots
z.grid <- matrix(crit.grid, n.grid, n.grid)
tit <- "Green: best point found by optimizer"
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
plot.axes = {title(tit);points(model@X[,1],model@X[,2],pch=17,col="blue");
points(X.genoud[1],X.genoud[2],pch=17,col="green");
axis(1); axis(2)})

```

## Description

Sequential optimization of kriging-based criterion conditional on noisy observations, with model update after each evaluation. Eight criteria are proposed to choose the next observation: random search, sequential parameter optimization (SPO), reinterpolation, Expected Improvement (EI) with plugin, Expected Quantile Improvement (EQI), quantile minimization, Augmented Expected Improvement (AEI) and Approximate Knowledge Gradient (AKG). The criterion optimization is based on the package rgenoud.

## Usage

```
noisy.optimizer(
  optim.crit,
  optim.param = NULL,
  model,
  n.ite,
  noise.var = NULL,
  funnoise,
  lower,
  upper,
  parinit = NULL,
  control = NULL,
  CovReEstimate = TRUE,
  NoiseReEstimate = FALSE,
  nugget.LB = 1e-05,
  estim.model = NULL,
  type = "UK"
)
```

## Arguments

<code>optim.crit</code>	String defining the criterion to be optimized at each iteration. Possible values are: "random.search", "SPO", "reinterpolation", "EI.plugin", "EQI", "min.quantile", "AEI", "AKG".
<code>optim.param</code>	List of parameters for the chosen criterion. For "EI.plugin": <code>optim.param\$plugin.type</code> is a string defining which plugin is to be used. Possible values are "ytilde", "quantile" and "other". If "quantile" is chosen, <code>optim.param\$quantile</code> defines the quantile level. If "other" is chosen, <code>optim.param\$plugin</code> directly sets the plugin value. For "EQI": <code>optim.param\$quantile</code> defines the quantile level. If not provided, default value is 0.9. For "min.quantile": <code>optim.param\$quantile</code> defines the quantile level. If not provided, default value is 0.1. For "AEI": <code>optim.param\$quantile</code> defines the quantile level to choose the current best point. If not provided, default value is 0.75.
<code>model</code>	a Kriging model of "km" class
<code>n.ite</code>	Number of iterations

noise.var	Noise variance (scalar). If noiseReEstimate=TRUE, it is an initial guess for the unknown variance (used in optimization).
funnoise	objective (noisy) function
lower	vector containing the lower bounds of the variables to be optimized over
upper	vector containing the upper bounds of the variables to be optimized over
parinit	optional vector of initial values for the variables to be optimized over
control	optional list of control parameters for optimization. One can control "pop.size" (default : $[N=3*2^{\dim}$ for $\dim < 6$ and $N=32*\dim$ otherwise]), "max.generations" (N), "wait.generations" (2) and "BFGSburnin" (0) of function "genoud" (see <a href="#">genoud</a> ). Numbers into brackets are the default values
CovReEstimate	optional boolean specifying if the covariance parameters should be re-estimated at every iteration (default value = TRUE)
NoiseReEstimate	optional boolean specifying if the noise variance should be re-estimated at every iteration (default value = FALSE)
nugget.LB	optional scalar of minimal value for the estimated noise variance. Default value is $1e-5$ .
estim.model	optional kriging model of "km" class with homogeneous nugget effect (no noise.var). Required if noise variance is reestimated and the initial "model" has heterogeneous noise variances.
type	"SK" or "UK" for Kriging with known or estimated trend

### Value

A list with components:

model	the current (last) kriging model of "km" class
best.x	The best design found
best.y	The objective function value at best.x
best.index	The index of best.x in the design of experiments
history.x	The added observations
history.y	The added observation values
history.hyperparam	The history of the kriging parameters
estim.model	If noiseReEstimate=TRUE, the current (last) kriging model of "km" class for estimating the noise variance.
history.noise.var	If noiseReEstimate=TRUE, the history of the noise variance estimate.

### Author(s)

Victor Picheny

## References

V. Picheny and D. Ginsbourger (2013), Noisy kriging-based optimization methods: A unified implementation within the DiceOptim package, *Computational Statistics & Data Analysis*

## Examples

```
#####
### EXAMPLE 1: 3 OPTIMIZATION STEPS USING EQI WITH KNOWN NOISE      ###
### AND KNOWN COVARIANCE PARAMETERS FOR THE BRANIN FUNCTION      ###
#####

set.seed(10)
library(DiceDesign)
# Set test problem parameters
doe.size <- 9
dim <- 2
test.function <- get("branin2")
lower <- rep(0,1,dim)
upper <- rep(1,1,dim)
noise.var <- 0.1

# Build noisy simulator
funnoise <- function(x)
{   f.new <- test.function(x) + sqrt(noise.var)*rnorm(n=1)
    return(f.new)}

# Generate DOE and response
doe <- as.data.frame(lhsDesign(doe.size, dim)$design)
y.tilde <- funnoise(doe)

# Create kriging model
model <- km(y~1, design=doe, response=data.frame(y=y.tilde),
            covtype="gauss", noise.var=rep(noise.var,1,doe.size),
            lower=rep(.1,dim), upper=rep(1,dim), control=list(trace=FALSE))

# Optimisation with noisy.optimizer (n.ite can be increased)
n.ite <- 2
optim.param <- list()
optim.param$quantile <- .9
optim.result <- noisy.optimizer(optim.crit="EQI", optim.param=optim.param, model=model,
n.ite=n.ite, noise.var=noise.var, funnoise=funnoise, lower=lower, upper=upper,
NoiseReEstimate=FALSE, CovReEstimate=FALSE)

new.model <- optim.result$model
best.x <- optim.result$best.x
new.doe <- optim.result$history.x

## Not run:
##### DRAW RESULTS #####
# Compute actual function on a grid
n.grid <- 12
```

```

x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
names(design.grid) <- c("V1","V2")
nt <- nrow(design.grid)
func.grid <- rep(0,1,nt)

for (i in 1:nt)
{ func.grid[i] <- test.function(x=design.grid[i,])}

# Compute initial and final kriging on a grid
pred <- predict(object=model, newdata=design.grid, type="UK", checkNames = FALSE)
mk.grid1 <- pred$m
sk.grid1 <- pred$sd

pred <- predict(object=new.model, newdata=design.grid, type="UK", checkNames = FALSE)
mk.grid2 <- pred$m
sk.grid2 <- pred$sd

# Plot initial kriging mean
z.grid <- matrix(mk.grid1, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
plot.axes = {title("Initial kriging mean");
points(model@X[,1],model@X[,2],pch=17,col="black");
axis(1); axis(2)})

# Plot initial kriging variance
z.grid <- matrix(sk.grid1^2, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
plot.axes = {title("Initial kriging variance");
points(model@X[,1],model@X[,2],pch=17,col="black");
axis(1); axis(2)})

# Plot final kriging mean
z.grid <- matrix(mk.grid2, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
plot.axes = {title("Final kriging mean");
points(new.model@X[,1],new.model@X[,2],pch=17,col="black");
axis(1); axis(2)})

# Plot final kriging variance
z.grid <- matrix(sk.grid2^2, n.grid, n.grid)
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
plot.axes = {title("Final kriging variance");
points(new.model@X[,1],new.model@X[,2],pch=17,col="black");
axis(1); axis(2)})

# Plot actual function and observations
z.grid <- matrix(func.grid, n.grid, n.grid)
tit <- "Actual function - Black: initial points; red: added points"
filled.contour(x.grid,y.grid, z.grid, nlevels=50, color = topo.colors,
plot.axes = {title(tit);points(model@X[,1],model@X[,2],pch=17,col="black");
points(new.doe[1,],new.doe[2,],pch=15,col="red");
axis(1); axis(2)})

```

```
## End(Not run)

#####
### EXAMPLE 2: 3 OPTIMIZATION STEPS USING EQI WITH UNKNOWN NOISE      ###
### AND UNKNOWN COVARIANCE PARAMETERS FOR THE BRANIN FUNCTION        ###
#####
# Same initial model and parameters as for example 1
n.ite <- 2 # May be changed to a larger value
res <- noisy.optimizer(optim.crit="min.quantile",
  optim.param=list(type="quantile",quantile=0.01),
  model=model, n.ite=n.ite, noise.var=noise.var, funnoise=funnoise,
  lower=lower, upper=upper,
  control=list(print.level=0),CovReEstimate=TRUE, NoiseReEstimate=TRUE)

# Plot actual function and observations
plot(model@X[,1], model@X[,2], pch=17,xlim=c(0,1),ylim=c(0,1))
points(res$history.x[1,], res$history.x[2,], col="blue")

# Restart: requires the output estim.model of the previous run
# to deal with potential repetitions
res2 <- noisy.optimizer(optim.crit="min.quantile",
  optim.param=list(type="quantile",quantile=0.01),
  model=res$model, n.ite=n.ite, noise.var=noise.var, funnoise=funnoise,
  lower=lower, upper=upper, estim.model=res$estim.model,
  control=list(print.level=0),CovReEstimate=TRUE, NoiseReEstimate=TRUE)

# Plot new observations
points(res2$history.x[1,], res2$history.x[2,], col="red")
```

---

ParrConstraint

*2D constraint function*


---

### Description

Strongly multimodal constraint function from Parr et al. (standardized version)

### Usage

```
ParrConstraint(x)
```

### Arguments

*x* a 2-dimensional vector or a two-column matrix specifying the location(s) where the function is to be evaluated.

### Value

A scalar

**Examples**

```

n.grid <- 20
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, ParrConstraint)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid,y.grid,z.grid,40)
title("Parr constraint function")

```

---

qEGO.nsteps	<i>Sequential multipoint Expected improvement (qEI) maximizations and model re-estimation</i>
-------------	---

---

**Description**

Executes `nsteps` iterations of the multipoint EGO method to an object of class `km`. At each step, a kriging model (including covariance parameters) is re-estimated based on the initial design points plus the points visited during all previous iterations; then a new batch of points is obtained by maximizing the multipoint Expected Improvement criterion (qEI).

**Usage**

```

qEGO.nsteps(
  fun,
  model,
  npoints,
  nsteps,
  lower = rep(0, model@d),
  upper = rep(1, model@d),
  crit = "exact",
  minimization = TRUE,
  optimcontrol = NULL,
  cov.reestim = TRUE,
  ...
)

```

**Arguments**

<code>fun</code>	the objective function to be optimized,
<code>model</code>	an object of class <code>km</code> ,
<code>npoints</code>	an integer representing the desired batchsize,
<code>nsteps</code>	an integer representing the desired number of iterations,
<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,



crit	"exact", "CL" : a string specifying the criterion used. "exact" triggers the maximization of the multipoint expected improvement at each iteration (see <a href="#">max_qEI</a> ), "CL" applies the Constant Liar heuristic,
minimization	logical specifying if we want to minimize or maximize fun,
optimcontrol	an optional list of control parameters for the qEI optimization (see details or <a href="#">max_qEI</a> ),
cov.reestim	optional boolean specifying if the kriging hyperparameters should be re-estimated at each iteration,
...	optional arguments for fun.

## Details

The parameters of list `optimcontrol` are :

- `optimcontrol$method` : "BFGS" (default), "genoud" ; a string specifying the method used to maximize the criterion (irrelevant when `crit` is "CL" because this method always uses `genoud`),

- when `crit="CL"` :

- + `optimcontrol$parinit` : optional matrix of initial values (must have `model@d` columns, the number of rows is not constrained),

- + `optimcontrol$L` : "max", "min", "mean" or a scalar value specifying the liar ; "min" takes `model@min`, "max" takes `model@max`, "mean" takes the prediction of the model ; When `L` is NULL, "min" is taken if `minimization==TRUE`, else it is "max".

- + The parameters of function `genoud`. Main parameters are : "pop.size" (default :  $[N=3*2^{\text{model@d}}$  for  $\text{dim}<6$  and  $N=32*\text{model@d}$  otherwise]), "max.generations" (default : 12), "wait.generations" (default : 2) and "BFGSburnin" (default : 2).

- when `optimcontrol$method = "BFGS"` :

- + `optimcontrol$nStarts` (default : 4),

- + `optimcontrol$fastCompute` : if TRUE (default), a fast approximation method based on a semi-analytic formula is used, see [Marmin 2014] for details,

- + `optimcontrol$samplingFun` : a function which sample a batch of starting point (default : `sampleFromEI`),

- + `optimcontrol$parinit` : optional 3d-array of initial (or candidate) batches (for all `k`, `parinit[,k]` is a matrix of size `npoints*model@d` representing one batch). The number of initial batches (`length(parinit[1,1])`) is not constrained and does not have to be equal to `nStarts`. If there is too few initial batches for `nStarts`, missing batches are drawn with `samplingFun` (default : NULL),

- when `optimcontrol$method = "genoud"` :

- + `optimcontrol$fastCompute` : if TRUE (default), a fast approximation method based on a semi-analytic formula is used, see [Marmin 2014] for details,

- + `optimcontrol$parinit` : optional matrix of candidate starting points (one row corresponds to one point),

- + The parameters of the `genoud` function. Main parameters are "pop.size" (default :  $[50*(\text{model@d})*(\text{npoints})]$ ), "max.generations" (default : 5), "wait.generations" (default : 2), "BFGSburnin" (default : 2).

**Value**

A list with components:

par	a data frame representing the additional points visited during the algorithm,
value	a data frame representing the response values at the points given in par,
npoints	an integer representing the number of parallel computations,
nsteps	an integer representing the desired number of iterations (given in argument),
lastmodel	an object of class <code>km</code> corresponding to the last kriging model fitted,
history	a vector of size nsteps representing the current known optimum at each step.

**Author(s)**

Sebastien Marmin

Clement Chevalier

David Ginsbourger

**References**

C. Chevalier and D. Ginsbourger (2014) Learning and Intelligent Optimization - 7th International Conference, Lion 7, Catania, Italy, January 7-11, 2013, Revised Selected Papers, chapter Fast computation of the multipoint Expected Improvement with applications in batch selection, pages 59-69, Springer.

D. Ginsbourger, R. Le Riche, L. Carraro (2007), A Multipoint Criterion for Deterministic Parallel Global Optimization based on Kriging. The International Conference on Non Convex Programming, 2007.

D. Ginsbourger, R. Le Riche, and L. Carraro. Kriging is well-suited to parallelize optimization (2010), In Lim Meng Hiot, Yew Soon Ong, Yoel Tenne, and Chi-Keong Goh, editors, *Computational Intelligence in Expensive Optimization Problems*, Adaptation Learning and Optimization, pages 131-162. Springer Berlin Heidelberg.

S. Marmin. Developpements pour l'evaluation et la maximisation du critere d'amelioration esperee multipoint en optimisation globale (2014). Master's thesis, Mines Saint-Etienne (France) and University of Bern (Switzerland).

J. Mockus (1988), *Bayesian Approach to Global Optimization*. Kluwer academic publishers.

M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

**See Also**

[qEI](#), [max\\_qEI](#), [qEI.grad](#)

## Examples

```

set.seed(123)
#####
### 2 ITERATIONS OF EGO ON THE BRANIN FUNCTION, ###
### STARTING FROM A 9-POINTS FACTORIAL DESIGN ###
#####

# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# EGO n steps
library(rgenoud)
nsteps <- 2 # increase to 10 for a more meaningful example
lower <- rep(0,d)
upper <- rep(1,d)
npoints <- 3 # The batchsize
oEGO <- qEGO.nsteps(model = fitted.model1, branin, npoints = npoints, nsteps = nsteps,
crit="exact", lower, upper, optimcontrol = NULL)
print(oEGO$par)
print(oEGO$value)
plot(c(1:nsteps),oEGO$history,xlab='step',ylab='Current known minimum')

## Not run:
# graphics
n.grid <- 15 # increase to 21 for better picture
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, branin)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid, y.grid, z.grid, 40)
title("Branin function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par, pch=19, col="red")
text(oEGO$par[,1], oEGO$par[,2], labels=c(tcrossprod(rep(1,npoints),1:nsteps)), pos=3)

## End(Not run)

```

---

qEI	<i>Analytical expression of the multipoint expected improvement (qEI) criterion</i>
-----	---

---

### Description

Computes the multipoint expected improvement criterion.

### Usage

```
qEI(
  x,
  model,
  plugin = NULL,
  type = "UK",
  minimization = TRUE,
  fastCompute = TRUE,
  eps = 10^(-5),
  envir = NULL
)
```

### Arguments

<code>x</code>	a matrix representing the set of input points (one row corresponds to one point) where to evaluate the qEI criterion,
<code>model</code>	an object of class <code>km</code> ,
<code>plugin</code>	optional scalar: if provided, it replaces the minimum of the current observations,
<code>type</code>	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account,
<code>minimization</code>	logical specifying if EI is used in minimization or in maximization,
<code>fastCompute</code>	if TRUE, a fast approximation method based on a semi-analytic formula is used (see [Marmin 2014] for details),
<code>eps</code>	the value of <i>epsilon</i> of the fast computation trick. Relevant only if <code>fastComputation</code> is TRUE,
<code>envir</code>	an optional environment specifying where to get intermediate values calculated in <code>qEI</code> .

### Value

The multipoint Expected Improvement, defined as

$$qEI(X_{new}) := E[(\min(Y(X)) - \min(Y(X_{new})))_+ | Y(X) = y(X)],$$

where  $X$  is the current design of experiments,  $X_{new}$  is a new candidate design, and  $Y$  is a random process assumed to have generated the objective function  $y$ .

**Author(s)**

Sebastien Marmin  
 Clement Chevalier  
 David Ginsbourger

**References**

- C. Chevalier and D. Ginsbourger (2014) Learning and Intelligent Optimization - 7th International Conference, Lion 7, Catania, Italy, January 7-11, 2013, Revised Selected Papers, chapter Fast computation of the multipoint Expected Improvement with applications in batch selection, pages 59-69, Springer.
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- D. Ginsbourger, R. Le Riche, and L. Carraro. Kriging is well-suited to parallelize optimization (2010), In Lim Meng Hiot, Yew Soon Ong, Yoel Tenne, and Chi-Keong Goh, editors, *Computational Intelligence in Expensive Optimization Problems*, Adaptation Learning and Optimization, pages 131-162. Springer Berlin Heidelberg.
- J. Mockus (1988), *Bayesian Approach to Global Optimization*. Kluwer academic publishers.
- M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

**See Also**

[EI](#)

**Examples**

```
set.seed(007)

# Monte-Carlo validation

# a 4-d, 81-points grid design, and the corresponding response
d <- 4; n <- 3^d
design <- do.call(expand.grid,rep(list(seq(0,1,length=3)),d))
names(design) <- paste("x",1:d,sep="")
y <- data.frame(apply(design, 1, hartman4))
names(y) <- "y"

# learning
model <- km(~1, design=design, response=y, control=list(trace=FALSE))

# pick up 10 points sampled from the 1-point expected improvement
```

```

q <- 10
X <- sampleFromEI(model,n=q)
# simulation of the minimum of the kriging random vector at X
t1 <- proc.time()
newdata <- as.data.frame(X)
colnames(newdata) <- colnames(model@X)

krig <- predict(object=model, newdata=newdata,type="UK",se.compute=TRUE, cov.compute=TRUE)
mk <- krig$mean
Sigma.q <- krig$cov
mychol <- chol(Sigma.q)
nsim <- 300000
white.noise <- rnorm(n=nsim*q)
minYsim <- apply(crossprod(mychol,matrix(white.noise,nrow=q)) + mk,2,min)
# simulation of the improvement (minimization)
qImprovement <- (min(model@y)-minYsim)*((min(model@y)-minYsim) > 0)

# empirical expectation of the improvement and confident interval (95%)
eiMC <- mean(qImprovement)
confInterv <- c(eiMC - 1.96*sd(qImprovement)*1/sqrt(nsim),eiMC + 1.96*sd(qImprovement)*1/sqrt(nsim))

# MC estimation of the qEI
print(eiMC)
t2 <- proc.time()
# qEI with analytical formula
qEI(X,model,fastCompute= FALSE)
t3 <- proc.time()
# qEI with fast computation trick
qEI(X,model)
t4 <- proc.time()
t2-t1 # Time of MC computation
t3-t2 # Time of normal computation
t4-t3 # Time of fast computation

```

---

qEI.grad

*Gradient of the multipoint expected improvement (qEI) criterion*


---

### Description

Computes an exact or approximate gradient of the multipoint expected improvement criterion

### Usage

```

qEI.grad(
  x,
  model,
  plugin = NULL,
  type = "UK",

```

```

    minimization = TRUE,
    fastCompute = TRUE,
    eps = 10^(-6),
    envir = NULL
)

```

### Arguments

x	a matrix representing the set of input points (one row corresponds to one point) where to evaluate the gradient,
model	an object of class <a href="#">km</a> ,
plugin	optional scalar: if provided, it replaces the minimum of the current observations,
type	"SK" or "UK" (by default), depending whether uncertainty related to trend estimation has to be taken into account,
minimization	logical specifying if EI is used in minimization or in maximization,
fastCompute	if TRUE, a fast approximation method based on a semi-analytic formula is used (see [Marmin 2014] for details),
eps	the value of <i>epsilon</i> of the fast computation trick. Relevant only if fastComputation is TRUE,
envir	an optional environment specifying where to get intermediate values calculated in <a href="#">qEI</a> .

### Value

The gradient of the multipoint expected improvement criterion with respect to  $x$ . A 0-matrix is returned if the batch of input points contains twice the same point or a point from the design experiment of the `km` object (the gradient does not exist in these cases).

### Author(s)

Sebastien Marmin  
 Clement Chevalier  
 David Ginsbourger

### References

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J. Mockus (1988), *Bayesian Approach to Global Optimization*. Kluwer academic publishers.

M. Schonlau (1997), *Computer experiments and global optimization*, Ph.D. thesis, University of Waterloo.

## See Also

[qEI](#)

## Examples

```
set.seed(15)
# Example 1 - validation by comparison to finite difference approximations

# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design)<-c("x1", "x2")
design <- data.frame(design)
names(design)<-c("x1", "x2")
y <- apply(design, 1, branin)
y <- data.frame(y)
names(y) <- "y"

# learning
model <- km(~1, design=design, response=y)

# pick up 2 points sampled from the simple expected improvement
q <- 2 # increase to 4 for a more meaningful test
X <- sampleFromEI(model,n=q)

# compute the gradient at the 4-point batch
grad.analytic <- qEI.grad(X,model)
# numerically compute the gradient
grad.numeric <- matrix(NaN,q,d)
eps <- 10^(-6)
EPS <- matrix(0,q,d)
for (i in 1:q) {
  for (j in 1:d) {
    EPS[i,j] <- eps
    grad.numeric[i,j] <- 1/eps*(qEI(X+EPS,model,fastCompute=FALSE)-qEI(X,model,fastCompute=FALSE))
    EPS[i,j] <- 0
  }
}
print(grad.numeric)
print(grad.analytic)

## Not run:
```



```

# graphics: displays the EI criterion, the design points in black,
# the batch points in red and the gradient in blue.
nGrid <- 15
gridAxe1 <- seq(lower[1],upper[1],length=nGrid)
gridAxe2 <- seq(lower[2],upper[2],length=nGrid)
grid <- expand.grid(gridAxe1,gridAxe2)
aa <- apply(grid,1,EI,model=model)
myMat <- matrix(aa,nrow=nGrid)
image(x = gridAxe1, y = gridAxe2, z = myMat,
      col = colorRampPalette(c("darkgray","white"))(5*10),
      ylab = names(design)[1], xlab=names(design)[2],
      main = "qEI-gradient of a batch of 4 points", axes = TRUE,
      zlim = c(min(myMat), max(myMat)))
contour(x = gridAxe1, y = gridAxe2, z = myMat,
        add = TRUE, nlevels = 10)
points(X[,1],X[,2],pch=19,col='red')
points(model@X[,1],model@X[,2],pch=19)
arrows(X[,1],X[,2],X[,1]+0.012*grad.analytic[,1],X[,2]+0.012*grad.analytic[,2],col='blue')

## End(Not run)

```

---

sampleFromEI

*Sampling points according to the expected improvement criterion*


---

## Description

Samples  $n$  points from a distribution proportional to the expected improvement (EI) computed from a `km` object.

## Usage

```

sampleFromEI(
  model,
  minimization = TRUE,
  n = 1,
  initdistrib = NULL,
  lower = rep(0, model@d),
  upper = rep(1, model@d),
  T = NULL
)

```

## Arguments

<code>model</code>	an object of class <code>km</code> ,
<code>minimization</code>	logical specifying if EI is used in minimization or in maximization,
<code>n</code>	number of points to be sampled,
<code>initdistrib</code>	matrix of candidate points.

lower	vector of lower bounds,
upper	vector of upper bounds,
T	optional scalar : if provided, it replaces the current minimum (or maximum) of observations.

**Value**

A  $n \times d$  matrix containing the sampled points. If NULL,  $1000 \times d$  points are obtained by latin hypercube sampling,

**Author(s)**

Sebastien Marmin  
 Clement Chevalier  
 David Ginsbourger

**References**

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.

**See Also**

[EI](#), [km](#), [qEI](#)

**Examples**

```
set.seed(004)

# a 9-points factorial design, and the corresponding responses
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
lower <- c(0,0)
upper <- c(1,1)
names(response.branin) <- "y"

# model identification
fitted.model <- km(~1, design=design.fact, response=response.branin,
                  covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))
```

```

# sample a 30 point batch
batchSize <- 30
x <- sampleFromEI(model = fitted.model, n = batchSize, lower = lower, upper = upper)

# graphics
# displays the EI criterion, the design points in black and the EI-sampled points in red.
nGrid <- 15
gridAxe1 <- seq(lower[1],upper[1],length=nGrid)
gridAxe2 <- seq(lower[2],upper[2],length=nGrid)
grid <- expand.grid(gridAxe1,gridAxe2)
aa <- apply(grid,1,EI,model=fitted.model)
myMat <- matrix(aa,nrow=nGrid)
image(x = gridAxe1, y = gridAxe2, z = myMat,
      col = colorRampPalette(c("darkgray","white"))(5*10),
      ylab = names(design.fact)[1], xlab=names(design.fact)[2],
      main = "Sampling from the expected improvement criterion",
      axes = TRUE, zlim = c(min(myMat), max(myMat)))
contour(x = gridAxe1, y = gridAxe2, z = myMat,
        add = TRUE, nlevels = 10)
points(x[,1],x[,2],pch=19,col='red')
points(fitted.model@X[,1],fitted.model@X[,2],pch=19)

```

---

test\_feas\_vec

*Test constraints violation (vectorized)*


---

## Description

Test whether a set of constraints are violated or not, depending on their nature (equality or inequality) and tolerance parameters

## Usage

```
test_feas_vec(cst, equality = FALSE, tolConstraints = NULL)
```

## Arguments

cst	matrix of constraints (one column for each constraint function)
equality	either FALSE or a Boolean vector defining which constraints are treated as equalities
tolConstraints	tolerance (vector) for all constraints. If not provided, set to zero for inequalities and 0.05 for equalities

## Value

A Boolean vector, TRUE if the point is feasible, FALSE if at least one constraint is violated

**Author(s)**

Mickael Binois

Victor Picheny

TREGO.nsteps

*Trust-region based EGO algorithm.***Description**

Executes *nsteps* iterations of the TREGO method to an object of class `km`. At each step, a kriging model is 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 the Expected Improvement criterion (EI) over either the entire search space or restricted to a trust region. The trust region is updated at each iteration based on a sufficient decrease condition.

**Usage**

```
TREGO.nsteps(
  model,
  fun,
  nsteps,
  lower,
  upper,
  control = NULL,
  kmcontrol = NULL,
  trcontrol = NULL,
  trace = 0,
  n.cores = 1,
  ...
)
```

**Arguments**

<code>model</code>	an object of class <code>km</code> ,
<code>fun</code>	the objective function to be minimized,
<code>nsteps</code>	an integer representing the desired number of iterations,
<code>lower, upper</code>	vector of lower and upper bounds for the variables to be optimized over,
<code>control</code>	an optional list of control parameters for optimization. For now only the number of restarts can be set.
<code>kmcontrol</code>	an optional list representing the control variables for the re-estimation of the kriging model.

trcontrol	an optional list of control parameters for the trust-region scheme: <code>sigma</code> the initial size of the trust region, <code>x0</code> its initial center, <code>beta</code> the contraction factor, <code>alpha</code> its dilatation factor, <code>kappa</code> the forcing factor, <code>crit</code> the criterion used inside the TR (either "EI" or "gpmean"), <code>GLratio</code> number of consecutive global and local steps, <code>algo</code> either "TREGO" or "TRIKE", <code>minsigma</code> minimal sigma value, <code>maxsigma</code> maximal sigma value, <code>minEI</code> stopping criterion for TRIKE, <code>local.model</code> Boolean; if TRUE, a local model is used within the trust region, <code>local.trend</code> , <code>local.covtype</code> trend and covariance for the local model, <code>n.local.min</code> minimal number of points used to build the local model,
trace	between -1 (no trace) and 3 (full messages)
n.cores	number of cores used for EI maximisation
...	additional parameters to be given to fun

**Value**

A list with components:

par	a data frame representing the additional points visited during the algorithm,
value	a data frame representing the response values at the points given in par,
npoints	an integer representing the number of parallel computations (=1 here),
nsteps	an integer representing the desired number of iterations (given in argument),
lastmodel	an object of class <code>km</code> corresponding to the last kriging model fitted. If warping is true, y values are normalized (warped) and will not match value.
all.success	a vector of Boolean indicating the successful steps according to the sufficient decrease condition
all.steps	a vector of Boolean indicating which steps were global
all.sigma	history of trust region size
all.x0	history of trust region centers
local.model	if <code>trcontrol\$local.model=TRUE</code> , the latest local model

**Author(s)**

Victor Picheny

**References**

Diouane, Picheny, Le Riche, Scotto Di Perrotolo (2021), *TREGO: a Trust-Region Framework for Efficient Global Optimization*, ArXiv

**See Also**

[EI](#), [max\\_crit](#), [EI.grad](#)

**Examples**

```

set.seed(123)
#####
### 10 ITERATIONS OF TREGO ON THE BRANIN FUNCTION, ###
### STARTING FROM A 9-POINTS FACTORIAL DESIGN      ###
#####

# a 9-points factorial design, and the corresponding response
d <- 2
n <- 9
design.fact <- expand.grid(seq(0,1,length=3), seq(0,1,length=3))
names(design.fact)<-c("x1", "x2")
design.fact <- data.frame(design.fact)
names(design.fact)<-c("x1", "x2")
response.branin <- apply(design.fact, 1, branin)
response.branin <- data.frame(response.branin)
names(response.branin) <- "y"

# model identification
fitted.model1 <- km(~1, design=design.fact, response=response.branin,
covtype="gauss", control=list(pop.size=50,trace=FALSE), parinit=c(0.5, 0.5))

# TREGO n steps
nsteps <- 5
lower <- rep(0, d)
upper <- rep(1, d)
oEGO <- TREGO.nsteps(model=fitted.model1, fun=branin, nsteps=nsteps,
lower=lower, upper=upper)
print(oEGO$par)
print(oEGO$value)

# graphics
n.grid <- 15 # Was 20, reduced to 15 for speeding up compilation
x.grid <- y.grid <- seq(0,1,length=n.grid)
design.grid <- expand.grid(x.grid, y.grid)
response.grid <- apply(design.grid, 1, branin)
z.grid <- matrix(response.grid, n.grid, n.grid)
contour(x.grid, y.grid, z.grid, 40)
title("Branin function")
points(design.fact[,1], design.fact[,2], pch=17, col="blue")
points(oEGO$par, pch=19, col="red")
text(oEGO$par[,1], oEGO$par[,2], labels=1:nsteps, pos=3)

```

**Description**

Update of a noisy Kriging model when adding new observation, with or without covariance parameter re-estimation. When the noise level is unknown, a twin model "estim.model" is also updated.

**Usage**

```
update_km_noisyEGO(
  model,
  x.new,
  y.new,
  noise.var = 0,
  type = "UK",
  add.obs = TRUE,
  index.in.DOE = NULL,
  CovReEstimate = TRUE,
  NoiseReEstimate = FALSE,
  estim.model = NULL,
  nugget.LB = 1e-05
)
```

**Arguments**

model	a Kriging model of "km" class
x.new	a matrix containing the new points of experiments
y.new	a matrix containing the function values on the points NewX
noise.var	scalar: noise variance
type	kriging type: "SK" or "UK"
add.obs	boolean: if TRUE, the new point does not exist already in the design of experiment model@X
index.in.DOE	optional integer: if add.obs=TRUE, it specifies the index of the observation in model@X corresponding to x.new
CovReEstimate	optional boolean specifying if the covariance parameters should be re-estimated (default value = TRUE)
NoiseReEstimate	optional boolean specifying if the noise variance should be re-estimated (default value = TRUE)
estim.model	optional input of "km" class. Required if NoiseReEstimate=TRUE, in order to deal with repetitions.
nugget.LB	optional scalar: is used to define a lower bound on the noise variance.

**Value**

A list containing:

model	The updated Kriging model
estim.model	If NoiseReEstimate=TRUE, the updated estim.model
noise.var	If NoiseReEstimate=TRUE, the re-estimated noise variance

**Author(s)**

Victor Picheny

**References**

V. Picheny and D. Ginsbourger (2013), Noisy kriging-based optimization methods: A unified implementation within the DiceOptim package, *Computational Statistics & Data Analysis*



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