

Package ‘ParBayesianOptimization’

December 3, 2018

Title Parallel Bayesian Optimization of Hyperparameters

Version 0.0.1

Description Fast, flexible framework for implementing Bayesian optimization of model hyperparameters according to the methods described in Snoek et al. <arXiv:1206.2944>. The package allows the user to run scoring function in parallel, save intermediary results, and tweak other aspects of the process to fully utilize the computing resources available to the user.

URL <https://github.com/AnotherSamWilson/ParBayesianOptimization>

BugReports <https://github.com/AnotherSamWilson/ParBayesianOptimization/issues>

Depends R (>= 3.5.0), data.table (>= 1.11.8), GauPro, stats, foreach, dbscan

License GPL-2

Encoding UTF-8

LazyData true

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Suggests knitr,rmarkdown,xgboost,doParallel,ggplot2

VignetteBuilder knitr

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NeedsCompilation no

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 applyCluster

Find Clusters

Description

Applies DBSCAN algorithm to find local optimum candidate parameter sets

Usage

```
applyCluster(e = parent.frame())
```

Arguments

e the entire parent environment is made available to this function

Value

the number of values that are outside the bounds/

BayesianOptimization

Bayesian Optimization

Description

Flexible Bayesian optimization of model hyperparameters.

Usage

```
BayesianOptimization(FUN, bounds, saveIntermediate = NULL,
  leftOff = NULL, parallel = FALSE, packages = NULL, export = NULL,
  initialize = TRUE, initGrid = NULL, initPoints = 0, bulkNew = 1,
  nIters = 0, kern = "Matern52", beta = 0, acq = "ucb",
  stopImpatient = list(newAcq = "ucb", rounds = Inf), kappa = 2.576,
  eps = 0, gsPoints = 100, convThresh = 1e+07,
  minClusterUtility = NULL, noiseAdd = 0.25, verbose = 1)
```

Arguments

FUN the function to be maximized. This function should return a named list with at least 1 component. The first component must be named Score and should contain the metric to be maximized. You may return other named scalar elements that you wish to include in the final summary table.

bounds named list of lower and upper bounds for each hyperparameter. The names of the list should be arguments passed to FUN. Use "L" suffix to indicate integer hyperparameters.

<code>saveIntermediate</code>	character filepath (including file name) that specifies the location to save intermediary results. This will save a <code>data.table</code> as an RDS that can be specified as the <code>leftOff</code> parameter.
<code>leftOff</code>	<code>data.table</code> containing parameter-Score pairs. If supplied, the process will rbind this table to the parameter-Score pairs obtained through initialization. This table should be obtained from the file saved by <code>saveIntermediate</code> .
<code>parallel</code>	should the process run in parallel? If TRUE, several criteria must be met: <ul style="list-style-type: none"> • A parallel backend must be registered • FUN must be executable using only packages specified in <code>packages</code> (and base packages) • FUN must be executable using only the the objects specified in <code>export</code> • The function must be thread safe.
<code>packages</code>	character vector of the packages needed to run FUN.
<code>export</code>	character vector of object names needed to evaluate FUN.
<code>initialize</code>	should the process initialize a parameter-Score pair set? If FALSE, <code>leftOff</code> must be provided.
<code>initGrid</code>	user specified points to sample the target function, should be a <code>data.frame</code> or <code>data.table</code> with identical column names as <code>bounds</code> .
<code>initPoints</code>	number of randomly chosen points to sample the scoring function before Bayesian Optimization fitting the Gaussian Process.
<code>bulkNew</code>	integer that specifies the number of parameter combinations to try between each Gaussian process fit.
<code>nIters</code>	total number of parameter sets to be sampled, including initial set.
<code>kern</code>	a character that gets mapped to one of GauPro's <code>GauPro_kernel_beta</code> S6 classes. Determines the covariance function used in the gaussian process. Can be one of: <ul style="list-style-type: none"> • "Gaussian" • "Exponential" • "Matern52" • "Matern32"
<code>beta</code>	the kernel lengthscale parameter $\log_{10}(\theta)$. Passed to <code>GauPro_kernel_beta</code> specified in <code>kern</code> .
<code>acq</code>	acquisition function type to be used. Can be "ucb", "ei", "eips" or "poi". <ul style="list-style-type: none"> • ucb Upper Confidence Bound • ei Expected Improvement • eips Expected Improvement Per Second • poi Probability of Improvement
<code>stopImpatient</code>	a list containing <code>rounds</code> and <code>newAcq</code> , if <code>acq = "eips"</code> you can switch the acquisition function to <code>newAcq</code> after <code>rounds</code> parameter-score pairs are found.
<code>kappa</code>	tunable parameter kappa of GP Upper Confidence Bound, to balance exploitation against exploration, increasing kappa will incentivise exploration.

eps	tunable parameter epsilon of ei, eips and poi. Balances exploitation against exploration. Increasing eps will make the "improvement" threshold higher.
gsPoints	integer that specifies how many initial points to try when searching for the optimal parameter set. Increase this for a higher chance to find global optimum, at the expense of more time.
convThresh	convergence threshold passed to factr when the optim function (L-BFGS-B) is called. Lower values will take longer to converge, but may be more accurate.
minClusterUtility	number 0-1. Represents the minimum percentage of the optimal utility required for a less optimal local maximum to be included as a candidate parameter set in the next scoring function. If NULL, only the global optimum will be used as a candidate parameter set.
noiseAdd	if bulkNew > 1, specifies how much noise to add to the optimal candidate parameter set to obtain the other bulkNew-1 candidate parameter sets. New random draws are pulled from a shape(4,4) beta distribution centered at the optimal candidate parameter set with a range equal to noiseAdd*(Upper Bound - Lower Bound)
verbose	Whether or not to print progress. If 0, nothing will be printed. If 1, progress will be printed. If 2, progress and information about new parameter-score pairs will be printed.

Value

A list containing details about the process:

GPlist	The list of the gaussian process objects that were fit.
acqMaximums	The optimal parameters according to each gaussian process
ScoreDT	A list of all parameter-score pairs, as well as extra columns from FUN
BestPars	The best parameter set at each iteration

References

Jasper Snoek, Hugo Larochelle, Ryan P. Adams (2012) *Practical Bayesian Optimization of Machine Learning Algorithms*

Examples

```
# Example 1 - Optimization of a Linear Function
scoringFunction <- function(x) {
  a <- exp(-(2-x)^2)*1.5
  b <- exp(-(4-x)^2)*2
  c <- exp(-(6-x)^2)*1
  return(list(Score = a+b+c))
}

bounds <- list(x = c(0,8))

Results <- BayesianOptimization(
  FUN = scoringFunction
```

```

    , bounds = bounds
    , initPoints = 5
    , nIters = 8
    , gsPoints = 10
  )

## Not run:
# Example 2 - Hyperparameter Tuning in xgboost
library("xgboost")

data(agaricus.train, package = "xgboost")

Folds <- list( Fold1 = as.integer(seq(1,nrow(agaricus.train$data),by = 3))
               , Fold2 = as.integer(seq(2,nrow(agaricus.train$data),by = 3))
               , Fold3 = as.integer(seq(3,nrow(agaricus.train$data),by = 3)))

scoringFunction <- function(max_depth, min_child_weight, subsample) {

  dtrain <- xgb.DMatrix(agaricus.train$data,label = agaricus.train$label)

  Pars <- list( booster = "gbtree"
               , eta = 0.01
               , max_depth = max_depth
               , min_child_weight = min_child_weight
               , subsample = subsample
               , objective = "binary:logistic"
               , eval_metric = "auc")

  xgbcv <- xgb.cv( params = Pars
                  , data = dtrain
                  , nround = 100
                  , folds = Folds
                  , prediction = TRUE
                  , showsd = TRUE
                  , early_stopping_rounds = 5
                  , maximize = TRUE
                  , verbose = 0)

  return(list( Score = max(xgbcv$evaluation_log$test_auc_mean)
              , nrounds = xgbcv$best_iteration
              )
         )
}

bounds <- list(max_depth = c(2L, 10L)
              , min_child_weight = c(1, 100)
              , subsample = c(0.25, 1))

kern <- "Matern52"

acq <- "ei"

ScoreResult <- BayesianOptimization(

```

```
    FUN = scoringFunction
  , bounds = bounds
  , initPoints = 10
  , bulkNew = 1
  , nIters = 12
  , kern = kern
  , acq = acq
  , kappa = 2.576
  , verbose = 1
  , parallel = FALSE
  , gsPoints = 50)

## End(Not run)
```

CheckBounds

Check Bounds

Description

Checks if a list of parameters is within the supplied bounds

Usage

```
CheckBounds(x, Table, bounds)
```

Arguments

x	Parameter Name
Table	A data.table of parameter values to check
bounds	the bounds list

Value

the number of values that are outside the bounds/

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