Package ‘mcga’

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Description Machine coded genetic algorithm (MCGA) is a fast tool for real-valued optimization problems. It uses the byte representation of variables rather than real-values. It performs the classical crossover operations (uniform) on these byte representations. Mutation operator is also similar to classical mutation operator, which is to say, it changes a randomly selected byte value of a chromosome by +1 or -1 with probability 1/2. In MCGAs there is no need for encoding-decoding process and the classical operators are directly applicable on real-values. It is fast and can handle a wide range of a search space with high precision. Using a 256-unary alphabet is the main disadvantage of this algorithm but a moderate size population is convenient for many problems. Package also includes multi_mcga function for multi objective optimization problems. This function sorts the chromosomes using their ranks calculated from the non-dominated sorting algorithm.

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Description

Machine coded genetic algorithm (MCGA) is a fast tool for real-valued optimization problems. It uses the byte representation of variables rather than real-values. It performs the classical crossover operations (uniform) on these byte representations. Mutation operator is also similar to classical mutation operator, which is to say, it changes a randomly selected byte value of a chromosome by +1 or -1 with probability 1/2. In MCGAs there is no need for encoding-decoding process and the classical operators are directly applicable on real-values. It is fast and can handle a wide range of a search space with high precision. Using a 256-unary alphabet is the main disadvantage of this algorithm but a moderate size population is convenient for many problems.

Author(s)

Mehmet Hakan Satman

Maintainer: Mehmet Hakan Satman <mhsatman@istanbul.edu.tr>

Examples

```r
## Not run:
# A sample optimization problem
# Min f(xi) = (x1-7)^2 + (x2-77)^2 + (x3-777)^2 + (x4-7777)^2 + (x5-77777)^2
# The range of xi is unknown. The solution is
# x1 = 7
# x2 = 77
# x3 = 777
# x4 = 7777
# x5 = 77777
# Min f(xi) = 0
require("mcga")
f <- function(x){
  return ((x[1]-7)^2 + (x[2]-77)^2 +(x[3]-777)^2 + (x[4]-7777)^2 +(x[5]-77777)^2)
}
m <- mcga(popsize=200,
  chsize=5,
  minval=0.0,
  maxval=999999999.9,
  maxiter=2500,
  crossprob=1.0,
  mutateprob=0.01,
  evalFunc=f)

  cat("Best chromosome:
")
  print(m$population[1,])
  cat("Cost: ",m$costs[1],"\n")

## End(Not run)
```
arithmetic_crossover  Performs arithmetic crossover operation on a pair of two selected parent candidate solutions

Description

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter crossover= is set to arithmetic_crossover than the arithmetic crossover operator is applied in the genetic search. arithmetic_crossover generates offspring using the weighted mean of parents' genes. Weights are drawn randomly.

Usage

arithmetic_crossover(object, parents, ...)

Arguments

object  A GA::ga object
parents  Indices of the selected parents
...  Additional arguments to be passed to the function

Value

List of two generated offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

Examples

```r
f <- function(x){
  return(-sum( (x-5)^2 )
}
myga <- ga(type="real-valued", fitness = f, popSize = 100, maxiter = 100,
  min = rep(-50,5), max = rep(50,5), crossover = arithmetic_crossover)
print(myga@solution)
```
blx_crossover

**Performs blx (blend) crossover operation on a pair of two selected parent candidate solutions**

**Description**

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter crossover= is set to blx_crossover than the blx crossover operator is applied in the genetic search.

**Usage**

blx_crossover(object, parents, ...)

**Arguments**

- object: A GA::ga object
- parents: Indices of the selected parents
- ...: Additional arguments to be passed to the function

**Value**

List of two generated offspring

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**Examples**

```r
f <- function(x){
  return(-sum( (x-5)^2 ) )
}
myga <- ga(type="real-valued", fitness = f, popSize = 100, maxiter = 100,
            min = rep(-50,5), max = rep(50,5), crossover = blx_crossover)
print(myga@solution)
```
Description

This function is a C++ wrapper for mutating byte representation of a given candidate solution.

Usage

`ByteCodeMutation(bytes1, pmutation)`

Arguments

- `bytes1` A vector of bytes of a candidate solution
- `pmutation` Probability of mutation

Value

Byte vector of mutated solution

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

See Also

`ByteCodeMutationUsingDoubles`

Examples

```r
set.seed(1246)
print(pi)
bytes <- DoubleToBytes(pi)
mutated.bytes <- ByteCodeMutation(bytes, 0.10)
new.var <- BytesToDouble(mutated.bytes)
print(new.var)
```
ByteCodeMutationUsingDoubles

*Mutation operator for byte representation of double values*

**Description**

This function is a C++ wrapper for mutating byte representation of a given candidate solution.

**Usage**

```
ByteCodeMutationUsingDoubles(d, pmutation)
```

**Arguments**

- `d`: A vector of doubles
- `pmutation`: Probability of mutation

**Value**

Double vector of mutated solution

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**See Also**

ByteCodeMutation

**Examples**

```
set.seed(1246)
print(pi)
print(exp(1))
new.var <- ByteCodeMutationUsingDoubles(c(pi, exp(1)), 0.10)
print(new.var)
```
ByteCodeMutationUsingDoublesRandom

*Mutation operator for byte representation of double values*

**Description**

This function is a C++ wrapper for mutating byte representation of a given candidate solution. This mutation operator randomly changes a byte in the range of [0,255].

**Usage**

```r
ByteCodeMutationUsingDoublesRandom(d, pmutation)
```

**Arguments**

- `d` A vector of doubles
- `pmutation` Probability of mutation

**Value**

Double vector of mutated solution

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**See Also**

ByteCodeMutation

**Examples**

```r
set.seed(1246)
print(pi)
print(exp(1))
new.var <- ByteCodeMutationUsingDoublesRandom(c(pi, exp(1)), 0.10)
print(new.var)
```
BytesToDouble

Converting sizeof(double) bytes to a double value

Description
This function converts sizeof(double) bytes to a double typed value

Usage
BytesToDouble(x)

Arguments
x A vector of bytes (unsigned chars in C++)

Value
Corresponding double typed value for a given vector of bytes

Author(s)
Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

See Also
DoubleVectorToBytes
DoubleToBytes
ByteVectorToDoubles

Examples
print(BytesToDouble(DoubleToBytes(56.43)))

ByteVectorToDoubles

Converting p * sizeof(double) bytes to a vector of p double values

Description
This function converts a byte vector to a vector of doubles

Usage
ByteVectorToDoubles(b)
Arguments

b A vector of bytes (unsigned chars in C++)

Value

Corresponding vector of double typed values for a given vector of bytes

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

See Also

DoubleVectorToBytes
BytesToDouble
ByteVectorToDoubles

Examples

a <- DoubleVectorToBytes(c(56.54, 89.7666, 98.565))
b <- ByteVectorToDoubles(a)
print(b)

Description

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter crossover= is set to byte_crossover than the byte-coded crossover operator is applied in the genetic search. In mcga2 function, the hard-coded crossover parameter is set to byte_crossover by definition. byte_crossover function simply takes two double vectors (parents) and combines the bytes of doubles using a Uniform distribution with parameters 0 and 1.

Usage

byte_crossover(object, parents, ...)

Arguments

object A GA::ga object
parents Indices of the selected parents
... Additional arguments to be passed to the function
Value

List of two generated offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

References


See Also

cmcga2

Examples

```r
f <- function(x)
  return(-sum((x-5)^2))

myga <- GA::ga(type="real-valued", fitness = f, popSize = 100, maxiter = 200,
  lower = rep(-50,5), upper = rep(50,5), crossover = byte_crossover,
  mutation = byte_mutation)
print(myga@solution)
```

byte_crossover_1p

Performs one-point crossover operation on a pair of two selected parent candidate solutions

Description

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter crossover= is set to byte_crossover_1p than the byte-coded one-point crossover operator is applied in the genetic search. In mcga2 function, the hard-coded crossover parameter is set to byte_crossover by definition. byte_crossover_1p function simply takes two double vectors (parents) and combines the bytes of doubles using given cut-point.

Usage

`byte_crossover_1p(object, parents, ...)`

Arguments

<table>
<thead>
<tr>
<th>Object</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>object</td>
<td>A GA::ga object</td>
</tr>
<tr>
<td>parents</td>
<td>Indices of the selected parents</td>
</tr>
<tr>
<td>...</td>
<td>Additional arguments to be passed to the function</td>
</tr>
</tbody>
</table>
Value

List of two generated offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

References


See Also

mcga2

Examples

f <- function(x){
  return(-sum( (x-5)^2 ) )
}
myga <- GA::ga(type="real-valued", fitness = f, popSize = 100, maxiter = 200,
  min = rep(-50,5), max = rep(50,5), crossover = byte_crossover_1p,
  mutation = byte_mutation)
print(myga@solution)

---

### byte_crossover_2p

Performs two-point crossover operation on a pair of two selected parent candidate solutions

Description

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter crossover= is set to byte_crossover_2p than the byte-coded two-point crossover operator is applied in the genetic search. In mcga2 function, the hard-coded crossover parameter is set to byte_crossover by definition. byte_crossover_2p function simply takes two double vectors (parents) and combines the bytes of doubles using given cutpoint1 and cutpoint2.

Usage

byte_crossover_2p(object, parents, ...)

Arguments

- `object` A GA::ga object
- `parents` Indices of the selected parents
- `...` Additional arguments to be passed to the function
Value

List of two generated offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

References


See Also

mcga2

Examples

```r
f <- function(x){
  return(-sum( (x-5)^2 ) )
}
myga <- GA::ga(type="real-valued", fitness = f, popSize = 100, maxiter = 200, min = rep(-50,5), max = rep(50,5), crossover = byte_crossover_2p, mutation = byte_mutation)
print(myga@solution)
```

Description

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter mutation= is set to byte_mutation than the byte-coded mutation operator is applied in the genetic search. In mcga2 function, the hard-coded mutation parameter is set to byte_mutation by definition. Byte-mutation function simply takes an double vector and changes bytes of this values by +1 or -1 using the pre-determined mutation probailty.

Usage

```
byte_mutation(object, parent, ...)
```

Arguments

- `object` A GA::ga object
- `parent` Index of the candidate solution of the current population
- `...` Additional arguments to be passed to the function
byte_mutation_dynamic

Value
Mutated double vector

Author(s)
Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

References

Examples
```r
f <- function(x){
  return(-sum((x-5)^2))
}
myga <- GA::ga(type="real-valued", fitness = f, popSize = 100, maxiter = 200,
  min = rep(-50,5), max = rep(50,5), crossover = byte_crossover,
  mutation = byte_mutation)
print(myga@solution)
```

byte_mutation_dynamic  Performs mutation operation on a given double vector using dynamic mutation probabilities

Description
This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter mutation is set to byte_mutation_dynamic than the byte-coded mutation operator is applied in the genetic search. In mcega2 function, the hard-coded mutation parameter is set to byte_mutation by definition. Byte-mutation function simply takes an double vector and changes bytes of this values by +1 or -1 using the dynamically decreased and pre-determined mutation probability.

Usage
```r
byte_mutation_dynamic(object, parent, ...)
```

Arguments
- object A GA::ga object
- parent Index of the candidate solution of the current population
- ... Additional arguments to be passed to the function

Value
Mutated double vector
**byte_mutation_random**

**Author(s)**
Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**References**

**Examples**
```r
f <- function(x){
  return(-sum( (x-5)^2 ) )
}
myga <- GA::ga(type="real-valued", fitness = f, popSize = 100, maxiter = 200,
              min = rep(-50,5), max = rep(50,5), crossover = byte_crossover,
              mutation = byte_mutation_dynamic, pmutation = 0.10)
print(myga@solution)
```

---

**byte_mutation_random**  
Performs mutation operation on a given double vector

**Description**
This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter mutation= is set to byte_mutation_random than the byte-coded mutation operator is applied in the genetic search. In mcga2 function, the hard-coded mutation parameter is set to byte_mutation by definition. This function simply takes an double vector and changes bytes randomly in the range of [0,255] using the pre-determined mutation probability.

**Usage**
```r
byte_mutation_random(object, parent, ...)
```

**Arguments**
- **object**  
  A GA::ga object
- **parent**  
  Index of the candidate solution of the current population
- **...**  
  Additional arguments to be passed to the function

**Value**
Mutated double vector

**Author(s)**
Mehmet Hakan Satman - mhsatman@istanbul.edu.tr


References


Examples

```r
f <- function(x){
  return(-sum((x-5)^2))
}
myga <- GA::ga(type="real-valued", fitness = f, popSize = 100, maxiter = 200,
               min = rep(-50,5), max = rep(50,5), crossover = byte_crossover,
               mutation = byte_mutation_random, pmutation = 0.20)
print(myga@solution)
```

---

`byte_mutation_random_dynamic`

*Performs mutation operation on a given double vector with dynamic mutation probabilities*

---

Description

This function is not called directly but is given as a parameter in `GA::ga` function. In `GA::ga`, if the parameter `mutation=` is set to `byte_mutation_random_dynamic` than the byte-coded mutation operator with dynamic probabilities is applied in the genetic search. In `mcga2` function, the hard-coded mutation parameter is set to `byte_mutation` by definition. This function simply takes an double vector and changes bytes randomly in the range of \([0,255]\) using the descreasing values of pre-determined mutation probability by generations.

Usage

`byte_mutation_random_dynamic(object, parent, ...)`

Arguments

- `object`: A `GA::ga` object
- `parent`: Index of the candidate solution of the current population
- `...`: Additional arguments to be passed to the function

Value

Mutated double vector

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr
References

Examples
f <- function(x){
  return(-sum((x-5)^2 ))
}
# Increase popSize and maxiter for more precise solutions
myga <- GA::ga(type="real-valued", fitness = f, popSize = 100, maxiter = 200,
  min = rep(-50,5), max = rep(50,5), crossover = byte_crossover,
  mutation = byte_mutation_random_dynamic, pmutation = 0.20)
print(myga@solution)

DoubleToBytes

Description
This function returns a vector of byte values with the length of sizeof(double) for a given double typed value

Usage
DoubleToBytes(x)

Arguments
x A double typed value

Value
A vector of byte values with the length of sizeof(double) for a given double typed value

Author(s)
Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

See Also
DoubleVectorToBytes
BytesToDouble
ByteVectorToDoubles

Examples
print(DoubleToBytes(56.43))
DoubleVectorToBytes  
*Byte representation of a vector of double typed variables*

**Description**

This function returns a vector of byte values for a given vector of double typed values.

**Usage**

`DoubleVectorToBytes(d)`

**Arguments**

- `d`  
  A vector of double typed values

**Value**

returns a vector of byte values for a given vector of double typed values

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**See Also**

- `DoubleToBytes`
- `BytesToDouble`
- `ByteVectorToDoubles`

**Examples**

```r
print(DoubleVectorToBytes(c(56.54, 89.7666, 98.565)))
```

---

EnsureBounds  
*Altering vector of doubles to satisfy boundary constraints*

**Description**

Byte based crossover and mutation operators can generate variables out of bounds of the decision variables. This function controls if variables are between their lower and upper bounds and if not, draws random numbers between these ranges. This function directly modifies the argument doubles and does not return a value.

**Usage**

`EnsureBounds(doubles, mins, maxs)`
Arguments

- doubles: A vector of doubles
- mins: A vector of lower bounds of decision variables
- maxs: A vector of upper bounds of decision variables

Value

Function directly modifies the argument doubles and does not return a result.

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

See Also

- byte_crossover
- byte_mutation
- mcga2

Examples

```r
set.seed(1234)
x <- runif(10)
print(x)
# [1] 0.113703411 0.622299405 0.609274733 0.623379442 0.860915384 0.640310605
# [7] 0.009495756 0.232550506 0.666083758 0.514251141
EnsureBounds(x, mins=rep(0,10), maxs=rep(0.2,10))
print(x)
# [1] 0.113703411 0.138718258 0.108994967 0.056546717 0.184686697 0.058463168
# [7] 0.009495756 0.167459126 0.057244657 0.053364156
```

---

**flat_crossover**

*Performs flat crossover operation on a pair of two selected parent candidate solutions*

Description

This function is not called directly but is given as a parameter in `GA::ga` function. In `GA::ga`, if the parameter `crossover` is set to `flat_crossover` than the flat crossover operator is applied in the genetic search. `flat_crossover` draws a random number between parents’ genes and returns a pair of generated offspring

Usage

```r
flat_crossover(object, parents, ...)
```
linear_crossover

Arguments

<table>
<thead>
<tr>
<th>argument</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>object</td>
<td>A GA::ga object</td>
</tr>
<tr>
<td>parents</td>
<td>Indices of the selected parents</td>
</tr>
<tr>
<td>...</td>
<td>Additional arguments to be passed to the function</td>
</tr>
</tbody>
</table>

Value

List of two generated offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

Examples

```r
f <- function(x){
  return(-sum( (x-5)^2 )
}
myga <- ga(type="real-valued", fitness = f, popSize = 100, maxiter = 100,
  min = rep(-50,5), max = rep(50,5), crossover = flat_crossover)
print(myga@solution)
```

Description

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter crossover is set to linear_crossover than the linear crossover operator is applied in the genetic search. linear_crossover generates three offspring and performs a selection mechanism to determine best two of them.

Usage

```r
linear_crossover(object, parents, ...)
```

Arguments

<table>
<thead>
<tr>
<th>argument</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>object</td>
<td>A GA::ga object</td>
</tr>
<tr>
<td>parents</td>
<td>Indices of the selected parents</td>
</tr>
<tr>
<td>...</td>
<td>Additional arguments to be passed to the function</td>
</tr>
</tbody>
</table>

Value

List of two generated offspring
MaxDouble

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

Examples

f <- function(x){
return(-sum((x-5)^2))
}
myga <- ga(type="real-valued", fitness = f, popSize = 100, maxiter = 100,
           min = rep(-50,5), max = rep(50,5), crossover = linear_crossover)
print(myga@solution)

MaxDouble

Maximum value of a double typed variable

Description

Maximum value of a double typed variable

Usage

MaxDouble()

Value

Returns maximum value of a double typed variable in C++ compiler

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

Examples

print(MaxDouble())
mca

Performs machine coded genetic algorithms on a function subject to
be minimized.

Description

Machine coded genetic algorithm (MCGA) is a fast tool for real-valued optimization problems. It uses the byte representation of variables rather than real-values. It performs the classical crossover operations (uniform) on these byte representations. Mutation operator is also similar to classical mutation operator, which is to say, it changes a randomly selected byte value of a chromosome by +1 or -1 with probability 1/2. In MCGA there is no need for encoding-decoding process and the classical operators are directly applicable on real-values. It is fast and can handle a wide range of a search space with high precision. Using a 256-unary alphabet is the main disadvantage of this algorithm but a moderate size population is convenient for many problems.

Usage

mca(popsize, chsize, crossprob = 1.0, mutateprob = 0.01,
    elitism = 1, minval, maxval, maxiter = 10, evalFunc)

Arguments

popsize Number of chromosomes.
chsize Number of parameters.
crossprob Crossover probability. By default it is 1.0
mutateprob Mutation probability. By default it is 0.01
elitism Number of best chromosomes to be copied directly into next generation. By default it is 1
minval The lower bound of the randomized initial population. This is not a constraint for parameters.
maxval The upper bound of the randomized initial population. This is not a constraint for parameters.
maxiter The maximum number of generations. By default it is 10
evalFunc An R function. By default, each problem is a minimization.

Value

population Sorted population resulted after generations
costs Cost values for each chromosomes in the resulted population

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr
mcga2

References


Examples

# A sample optimization problem
# Min f(xi) = (x1-7)^2 + (x2-77)^2 + (x3-777)^2 + (x4-7777)^2 + (x5-77777)^2
# The range of xi is unknown. The solution is
# x1 = 7
# x2 = 77
# x3 = 777
# x4 = 7777
# x5 = 77777
# Min f(xi) = 0
require("mcga")
f<-function(x){
  return ((x[1]-7)^2 + (x[2]-77)^2 +(x[3]-777)^2 +(x[4]-7777)^2 +(x[5]-77777)^2)
}
m <- mcga( popsize=200,
chsize=5,
minval=0.0,
maxval=999999999.9,
maxiter=2500,
crossprob=1.0,
mutableprob=0.01,
evalFunc=f)

  cat("Best chromosome:
")
  print(m$population[1,])
  cat("Cost: ",m$costs[1],"\n")

mcga2

Performs a machine-coded genetic algorithm search for a given optimization problem

Description

mcga2 is the improvement version of the standard mcga function as it is based on the GA::ga function. The byte_crossover and the byte_mutation operators are the main reproduction operators and these operators uses the byte representations of parents in the computer memory.

Usage

mcga2(fitness, ..., min, max,
  population = gaControl("real-valued")$population,
  selection = gaControl("real-valued")$selection,
  crossover = byte_crossover, mutation = byte_mutation, popSize = 50,
  pcrossover = 0.8, pmutation = 0.1, elitism = base::max(1, round(popSize
mcga2

* 0.05)), maxiter = 100, run = maxiter, maxFitness = Inf,
  names = NULL, parallel = FALSE, monitor = gaMonitor, seed = NULL)

Arguments

  fitness  The goal function to be maximized
  ...      Additional arguments to be passed to the fitness function
  min      Vector of lower bounds of variables
  max      Vector of upper bounds of variables
  population  Initial population. It is gaControl("real-valued")$population by default.
  selection  Selection operator. It is gaControl("real-valued")$selection by default.
  crossover  Crossover operator. It is byte_crossover by default.
  mutation  Mutation operator. It is byte_mutation by default. Other values can be given
            including byte_mutation_random, byte_mutation_dynamic and byte_mutation_random_dynamic
  popSize   Population size. It is 50 by default
  pcrossover Probability of crossover. It is 0.8 by default
  pmutation Probability of mutation. It is 0.1 by default
  elitism   Number of elitist solutions. It is base::max(1, round(popSize*0.05)) by default
  maxiter   Maximum number of generations. It is 100 by default
  run       The genetic search is stopped if the best solution has not any improvements in
            last run generations. By default it is maxiter
  maxFitness Upper bound of the fitness function. By default it is Inf
  names     Vector of names of the variables. By default it is NULL
  parallel  If TRUE, fitness calculations are performed parallel. It is FALSE by default
  monitor   The monitoring function for printing some information about the current state
            of the genetic search. It is gaMonitor by default
  seed      The seed for random number generating. It is NULL by default

Value

  Returns an object of class ga-class

Author(s)

  Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

References

  M.H.Satman (2013), Machine Coded Genetic Algorithms for Real Parameter Optimization Prob-
  lems, Gazi University Journal of Science, Vol 26, No 1, pp. 85-95
  53(4), 1-37.
multi_mcga

See Also

GA::ga

Examples

f <- function(x){
  return(-sum( (x-5)^2 ) )
}
myga <- mcga2(fitness = f, popSize = 100, maxiter = 300,
               min = rep(-50,5), max = rep(50,5))
print(myga@solution)

---

multi_mcga  Performs multi objective machine coded genetic algorithms.

Description

Machine coded genetic algorithm (MCGA) is a fast tool for real-valued optimization problems. It uses the byte representation of variables rather than real-values. It performs the classical crossover operations (uniform) on these byte representations. Mutation operator is also similar to classical mutation operator, which is to say, it changes a randomly selected byte value of a chromosome by +1 or -1 with probability 1/2. In MCGAs there is no need for encoding-decoding process and the classical operators are directly applicable on real-values. It is fast and can handle a wide range of a search space with high precision. Using a 256-unary alphabet is the main disadvantage of this algorithm but a moderate size population is convenient for many problems.

This function performs multi objective optimization using the same logic underlying the mcga. Chromosomes are sorted by their objective values using a non-dominated sorting algorithm.

Usage

multi_mcga(popsize, chsize, crossprob = 1.0, mutateprob = 0.01,
           elitism = 1, minval, maxval, maxiter = 10, numfunc, evalFunc)

Arguments

  popsize  Number of chromosomes.
  chsize   Number of parameters.
  crossprob Crossover probability. By default it is 1.0
  mutateprob Mutation probability. By default it is 0.01
  elitism   Number of best chromosomes to be copied directly into next generation. By default it is 1
  minval    The lower bound of the randomized initial population. This is not a constraint for parameters.
  maxval    The upper bound of the randomized initial population. This is not a constraint for parameters.
maxiter The maximum number of generations. By default it is 10.
numfunc Number of objective functions.
evalFunc An R function. By default, each problem is a minimization. This function must return a cost vector with dimension of numfunc. Each element of this vector points to the corresponding function to optimize.

Value
population Sorted population resulted after generations
costs Cost values for each chromosomes in the resulted population
ranks Calculated ranks using a non-dominated sorting for each chromosome

Author(s)
Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

References

Examples
## Not run:
# We have two objective functions.
f1<-function(x){
  return(sin(x))
}

f2<-function(x){
  return(sin(2*x))
}

# This function returns a vector of cost functions for a given x sent from mcga
f<-function(x){
  return ( c( f1(x), f2(x)) )
}

# main loop
m<-multi_mcga(popsize=200, chsize=1, minval= 0, elitism=2,
  maxval= 2.0 * pi, maxiter=1000, crossprob=1.0,
  mutateprob=0.01, evalFunc=f, numfunc=2)

# Points show best five solutions.
curve(f1, 0, 2*pi)
curve(f2, 0, 2*pi, add=TRUE)

p <- m$population[1:5,]
points(p, f1(p))
points(p, f2(p))
OnePointCrossOver

## Description

This function is a C++ wrapper for crossing-over of two byte vectors of candidate solutions.

## Usage

```r
OnePointCrossOver(bytes1, bytes2, cutpoint)
```

## Arguments

- `bytes1`: A vector of bytes of the first parent.
- `bytes2`: A vector of bytes of the second parent.
- `cutpoint`: Cut-point for the single point crossing-over.

## Value

List of two byte vectors of offspring.

## Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

## See Also

- `UniformCrossOver`
- `UniformCrossOverOnDoublesUsingBytes`

## Examples

```r
b1 <- DoubleVectorToBytes(c(56.54, 89.7666, 98.565))
b2 <- DoubleVectorToBytes(c(79.76, 56.4443, 34.22121))
result <- OnePointCrossOver(b1, b2, round(runif(1, 1, SizeOfDouble() * 3)))
print(ByteVectorToDoubles(result[[1]]))
print(ByteVectorToDoubles(result[[2]]))
```
OnePointCrossOverOnDoublesUsingBytes

One-point Crossover operation on the two vectors of doubles using their byte representations

Description

This function is a C++ wrapper for crossing-over of two double vectors of candidate solutions using their byte representations

Usage

OnePointCrossOverOnDoublesUsingBytes(d1, d2, cutpoint)

Arguments

d1  A vector of doubles of the first parent
d2  A vector of doubles of the second parent
cutpoint  An integer between 1 and chromosome length for crossover cutting

Value

List of two double vectors of offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

See Also

OnePointCrossOver
UniformCrossOverOnDoublesUsingBytes

Examples

d1 <- runif(3)
d2 <- runif(3)
cutp <- sample(1:(length(d1)*SizeOfDouble()), 1)[1]
offspring <- OnePointCrossOverOnDoublesUsingBytes(d1, d2, cutp)
print("Parents:"
print(d1)
print(d2)
print("Offspring:"
print(offspring[[1]])
print(offspring[[2]])
sbox_crossover

Performs sbox (simulated binary) crossover operation on a pair of two selected parent candidate solutions

Description

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter crossover= is set to sbox_crossover than the sbox crossover operator is applied in the genetic search. sbox_crossover mimics the classical single-point crossover operator in binary genetic algorithms.

Usage

sbox_crossover(object, parents, ...)

Arguments

object A GA::ga object
parents Indices of the selected parents
... Additional arguments to be passed to the function

Value

List of two generated offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

References


Examples

f <- function(x){
  return(-sum( (x-5)^2 ) )
}
myga <- ga(type="real-valued", fitness = f, popSize = 100, maxiter = 100,
  min = rep(-50,5), max = rep(50,5), crossover = sbox_crossover)
print(myga@solution)
**SizeOfDouble**

*Byte-length of a double typed variable*

**Description**

Byte-length of a double typed variable in computer memory

**Usage**

SizeOfDouble()

**Value**

Returns the byte-length of a double typed variable in computer memory

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**Examples**

print(SizeOfDouble())

---

**SizeOfInt**

*Byte-length of a int typed variable*

**Description**

Byte-length of a int typed variable in computer memory

**Usage**

SizeOfInt()

**Value**

Returns the byte-length of a int typed variable in computer memory

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**Examples**

print(SizeOfInt())
**SizeOfLong**

**Byte-length of a long typed variable**

**Description**

Byte-length of a long typed variable in computer memory

**Usage**

SizeOfLong()

**Value**

Returns the byte-length of a long typed variable in computer memory

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**Examples**

print(SizeOfLong())

---

**TwoPointCrossOver**

*Two Point Crossover operation on the two vectors of bytes*

**Description**

This function is a C++ wrapper for crossing-over of two byte vectors of candidate solutions

**Usage**

TwoPointCrossOver(bytes1, bytes2, cutpoint1, cutpoint2)

**Arguments**

- `bytes1` : A vector of bytes of the first parent
- `bytes2` : A vector of bytes of the second parent
- `cutpoint1` : First cut-point for the single point crossing-over
- `cutpoint2` : Second cut-point for the single point crossing-over

**Value**

List of two byte vectors of offspring
TwoPointCrossOverOnDoublesUsingBytes

Author(s)
Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

See Also
OnePointCrossOver
OnePointCrossOverOnDoublesUsingBytes
UniformCrossOverOnDoublesUsingBytes

Examples
b1 <- DoubleVectorToBytes(c(56.54, 89.7666, 98.565))
b2 <- DoubleVectorToBytes(c(79.76, 56.4443, 34.22121))
cutpoints <- sort(sample(1:(length(b1)*SizeOfDouble()), 2, replace = FALSE))
result <- TwoPointCrossOver(b1,b2, cutpoints[1], cutpoints[2])
print(ByteVectorToDoubles(result[[1]]))
print(ByteVectorToDoubles(result[[2]]))

TwoPointCrossOverOnDoublesUsingBytes
Two-point Crossover operation on the two vectors of doubles using their byte representations

Description
This function is a C++ wrapper for crossing-over of two double vectors of candidate solutions using their byte representations

Usage
TwoPointCrossOverOnDoublesUsingBytes(d1, d2, cutpoint1, cutpoint2)

Arguments
d1 A vector of doubles of the first parent
d2 A vector of doubles of the second parent
cutpoint1 An integer between 1 and chromosome length for crossover cutting
cutpoint2 An integer between cutpoint1 and chromosome length for crossover cutting

Value
List of two double vectors of offspring

Author(s)
Mehmet Hakan Satman - mhsatman@istanbul.edu.tr
unfair_average_crossover

See Also

TwoPointCrossOver
OnePointCrossOver
UniformCrossOver
OnePointCrossOverOnDoublesUsingBytes

Examples

d1 <- runif(3)
d2 <- runif(3)
cutpoints <- sort(sample(1:(length(d1)*SizeOfDouble())), 2, replace = FALSE))
offspring <- TwoPointCrossOverOnDoublesUsingBytes(d1,d2,cutpoints[1], cutpoints[2])
print("Parents:")
print(d1)
print(d2)
print("Offspring:")
print(offspring[[1]])
print(offspring[[2]])

unfair_average_crossover

Performs unfair average crossover operation on a pair of two selected parent candidate solutions

Description

This function is not called directly but is given as a parameter in GA::ga function. In GA::ga, if the parameter crossover= is set to unfair_average_crossover than the unfair average crossover operator is applied in the genetic search.

Usage

unfair_average_crossover(object, parents, ...)

Arguments

object A GA::ga object
parents Indices of the selected parents
... Additional arguments to be passed to the function

Value

List of two generated offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr
Examples

\[ f \left( x \right) = \text{return} \left( -\sum \left( x-5 \right)^2 \right) \]

\[
\text{myga <- ga(type="real-valued", fitness = f, popSize = 100, maxiter = 100,}
\text{ min = rep(-50,5), max = rep(50,5), crossover = unfair_average_crossover)}
\]

\text{print(myga@solution)}

---

**UniformCrossOver**

**Uniform Crossover operation on the two vectors of bytes**

**Description**

This function is a C++ wrapper for crossing-over of two byte vectors of candidate solutions

**Usage**

```
UniformCrossOver(bytes1, bytes2)
```

**Arguments**

- `bytes1` A vector of bytes of the first parent
- `bytes2` A vector of bytes of the second parent

**Value**

List of two byte vectors of offspring

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**See Also**

OnePointCrossOver

UniformCrossOverOnDoublesUsingBytes

**Examples**

```r
b1 <- DoubleVectorToBytes(c(56.54, 89.7666, 98.565))
b2 <- DoubleVectorToBytes(c(79.76, 56.4443, 34.22121))
result <- UniformCrossOver(b1,b2)
print(ByteVectorToDoubles(result[[1]]))
print(ByteVectorToDoubles(result[[2]]))
```
UniformCrossOverOnDoublesUsingBytes

Description

This function is a C++ wrapper for crossing-over of two double vectors of candidate solutions using their byte representations.

Usage

UniformCrossOverOnDoublesUsingBytes(d1, d2)

Arguments

d1: A vector of doubles of the first parent
d2: A vector of doubles of the second parent

Value

List of two double vectors of offspring

Author(s)

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

See Also

OnePointCrossOver
OnePointCrossOverOnDoublesUsingBytes

Examples

d1 <- runif(3)
d2 <- runif(3)
offspring <- UniformCrossOverOnDoublesUsingBytes(d1, d2)
print("Parents:")
print(d1)
print(d2)
print("Offspring:")
print(offspring[[1]])
print(offspring[[2]])
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