

# Package ‘mcga’

May 13, 2018

**Type** Package

**Title** Machine Coded Genetic Algorithms for Real-Valued Optimization Problems

**Version** 3.0.3

**Date** 2018-05-12

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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.

**License** GPL (>= 2)

**Depends** GA

**Imports** Rcpp (>= 0.11.4)

**LinkingTo** Rcpp

**NeedsCompilation** yes

**LazyLoad** yes

**Repository** CRAN

**Date/Publication** 2018-05-13 20:30:12 UTC

**RoxygenNote** 5.0.1

**R topics documented:**

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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.

## Details

Package:	mcga
Type:	Package
Version:	2.0.3
Date:	2012-01-06
License:	GPL
LazyLoad:	yes

## Author(s)

Mehmet Hakan Satman

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

## Examples

```
## Not run:
# A sample optimization problem
# Min  $f(x_i) = (x_1-7)^2 + (x_2-77)^2 + (x_3-777)^2 + (x_4-7777)^2 + (x_5-77777)^2$ 
# The range of  $x_i$  is unknown. The solution is
#  $x_1 = 7$ 
#  $x_2 = 77$ 
#  $x_3 = 777$ 
#  $x_4 = 7777$ 
#  $x_5 = 77777$ 
# Min  $f(x_i) = 0$ 
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:\n")
  print(m$population[1,])
  cat("Cost: ",m$costs[1],"")

## 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

```

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	<i>Performs blx (blend) crossover operation on a pair of two selected parent candidate solutions</i>
---------------	--

---

### 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

```
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)
```

---

ByteCodeMutation	<i>Mutation operator for byte representation of double values</i>
------------------	---

---

## 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

```
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**

<code>d</code>	A vector of doubles
<code>pmutation</code>	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

```
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

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



---

BytesToDouble	<i>Converting sizeof(double) bytes to a double value</i>
---------------	--

---

**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	<i>Converting p * sizeof(double) bytes to a vector of p double values</i>
---------------------	---

---

**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)
```

---

byte_crossover	<i>Performs crossover operation on a pair of two selected parent candidate solutions</i>
----------------	--

---

**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**

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

**See Also**

mcga2

**Examples**

```
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	<i>Performs one-point crossover operation on a pair of two selected parent candidate solutions</i>
-------------------	--

---

**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**

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**

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

**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	<i>Performs two-point crossover operation on a pair of two selected parent candidate solutions</i>
-------------------	--

---

**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**

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

**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_2p,
              mutation = byte_mutation)
print(myga@solution)
```

---

byte_mutation	<i>Performs mutation operation on a given double vector</i>
---------------	---

---

**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 probability.

**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

**Value**

Mutated double vector

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**References**

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

**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,
              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 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 dynamically decreased and pre-determined mutation probability.

**Usage**

```
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

**Author(s)**

Mehmet Hakan Satman - mhsatman@istanbul.edu.tr

**References**

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

**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,
  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**

```
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

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

## 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,
  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 decreasing 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

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

## 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

*Byte representation of a double typed variable*

---

## 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**

```
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**

```
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	<i>Performs flat crossover operation on a pair of two selected parent candidate solutions</i>
----------------	---

---

**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**

```
flat_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 <- 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)
```

---

linear_crossover	<i>Performs linear crossover operation on a pair of two selected parent candidate solutions</i>
------------------	---

---

**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**

```
linear_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 <- 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())
```

---

mcga	<i>Performs machine coded genetic algorithms on a function subject to be minimized.</i>
------	---

---

### 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.

### Usage

```
mcga(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

## References

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

## 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,
  mutateprob=0.01,
  evalFunc=f)

cat("Best chromosome:\n")
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
```

```
* 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 <code>gaControl("real-valued")\$population</code> by default.
selection	Selection operator. It is <code>gaControl("real-valued")\$selection</code> by default.
crossover	Crossover operator. It is <code>byte_crossover</code> by default.
mutation	Mutation operator. It is <code>byte_mutation</code> by default. Other values can be given including <code>byte_mutation_random</code> , <code>byte_mutation_dynamic</code> and <code>byte_mutation_random_dynamic</code>
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 <code>base::max(1, round(popSize*0.05))</code> 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 <code>maxiter</code>
maxFitness	Upper bound of the fitness function. By default it is <code>Inf</code>
names	Vector of names of the variables. By default it is <code>NULL</code>
parallel	If <code>TRUE</code> , fitness calculations are performed parallel. It is <code>FALSE</code> by default
monitor	The monitoring function for printing some information about the current state of the genetic search. It is <code>gaMonitor</code> by default
seed	The seed for random number generating. It is <code>NULL</code> by default

### Value

Returns an object of class `ga-class`

### Author(s)

Mehmet Hakan Satman - [mhsatman@istanbul.edu.tr](mailto:mhsatman@istanbul.edu.tr)

### References

- M.H.Satman (2013), Machine Coded Genetic Algorithms for Real Parameter Optimization Problems, Gazi University Journal of Science, Vol 26, No 1, pp. 85-95
- Luca Scrucca (2013). GA: A Package for Genetic Algorithms in R. Journal of Statistical Software, 53(4), 1-37. URL <http://www.jstatsoft.org/v53/i04/>



**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**

Deb, K. (2000). An efficient constraint handling method for genetic algorithms. Computer methods in applied mechanics and engineering, 186(2), 311-338.

**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))
```

```
## End(Not run)
```

---

OnePointCrossOver      *One 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

```
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

```
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]])
```

---

sbx_crossover	<i>Performs sbx (simulated binary) crossover operation on a pair of two selected parent candidate solutions</i>
---------------	---

---

### 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 sbx\_crossover than the sbx crossover operator is applied in the genetic search. sbx\_crossover mimics the classical single-point crossover operator in binary genetic algorithms.

### Usage

```
sbx_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

Deb, Kalyanmoy, and Ram Bhushan Agrawal. "Simulated binary crossover for continuous search space." Complex systems 9.2 (1995): 115-148.

### 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 = sbx_crossover)
print(myga@solution)
```

---

SizeOfDouble	<i>Byte-length of a double typed variable</i>
--------------	---

---

**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	<i>Byte-length of a int typed variable</i>
-----------	--

---

**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	<i>Byte-length of a long typed variable</i>
------------	---

---

**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	<i>Two Point Crossover operation on the two vectors of bytes</i>
-------------------	--

---

**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

**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



**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 <- 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 = unfair_average_crossover)
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**

```
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

*Uniform 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

```
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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