

A genetic algorithm for the Knapsack problem

The aim of this document is to show how works a genetic algorithm (GA) for the knapsack problem. It is not a good alternative to solve these problems, but it can help to show how GAs work.

Defining test instances

To test the algorithm, we define first a function that generates instances of the knapsack problem. It is based in recommendations of

Pisinger, D. (2005). Where are the hard knapsack problems?. Computers & Operations Research, 32(9), 2271-2284.

The utility of each instance is quite close to its weight, so it can be somewhat hard to solve:

```
KPInstance01 <- function(R, size, h, seed=1212){  
  set.seed(1212)  
  w <- sample(1:R, size, replace=TRUE)  
  u <- sapply(1:size, function(x) sample((w[x] + R/10 - R/500):(w[x] + R/10 + R/500), 1))  
  W <- round(h/(101)*sum(w))  
  return(list(w=w,u=u,W=W))  
}
```

Every instance is encoded as a list containing the weights w , utilities u and knapsack capacity W .

Defining the GA

Crossover and mutation operators

Then we will define the building blocks of the genetic algorithm. First we define the **fitness function**. It is equal to total utility for feasible solutions, and equal to zero to non-feasible solutions:

```
FitnessKP <- function(dades, sol){  
  if(sum(dades$w*sol) <= dades$W) fit <- sum(dades$u*sol)  
  else fit <- 0  
  return(fit)  
}
```

The following function defines a 1-point **crossover operator** for the KP. It takes for granted that we encode the solution as a binary string of length equal to the number of items n .

```
CrossoverKP1point <- function(sol1, sol2){
  n <- length(sol1)
  point <- sample(1:(n-1), 1)
  son <- c(sol1[1:point], sol2[(point + 1):n])

  return(son)
}
```

To test the operator, here are several crossovers with a all-zeroes and a all-ones strings:

```
set.seed(1213)
a <- rep(1, 10)
b <- rep(0, 10)
for(i in 1:5) print(CrossoverKP1point(a,b))

## [1] 1 1 0 0 0 0 0 0 0 0
## [1] 1 1 0 0 0 0 0 0 0 0
## [1] 1 1 1 1 1 1 1 1 0 0
## [1] 1 1 1 1 0 0 0 0 0 0
## [1] 1 0 0 0 0 0 0 0 0 0
```

This function defines a three-point **mutation operator**. It changes the value of three consecutive positions of a string:

```
MutationKP3points <- function(sol){
  n <- length(sol)
  point <- sample(1:(n-2), 1)
  sol <- as.logical(sol)
  sol[point:(point+2)] <- !sol[point:(point+2)]
  sol <- as.numeric(sol)

  return(sol)
}
```

Let's run five times the mutation operator for a all-ones string:

```
ex <- rep(1, 10)
for(i in 1:5) print(MutationKP3points(ex))

## [1] 0 0 0 1 1 1 1 1 1 1
## [1] 0 0 0 1 1 1 1 1 1 1
## [1] 1 1 1 1 1 1 1 0 0 0
## [1] 1 1 1 1 1 1 0 0 0 1
## [1] 1 1 0 0 0 1 1 1 1 1
```

The genetic algorithm function

Here is the function that runs the GA for an instance of KP:

```
GeneticAlgorithmKP <- function(dades, npop, iterations, pmut, elitist = TRUE,
verbose = FALSE){
```

```

#problem size
n <- length(dades$w)

#initial population
rel.size <- min(dades$W/sum(dades$w),1)
population.parent <- replicate(npop, sample(0:1, n, replace = TRUE, prob=c(1-rel.size,
rel.size)), simplify=FALSE)

#initializing next generation
population.son <- replicate(npop, numeric(n), simplify=FALSE)

best.obj <- 0
best.sol <- numeric(n)

count <- 1

while(count <= iterations){

  #assess fitness of population parent

  fitness <- sapply(population.parent, function(x) FitnessKP(dades, x))

  #finding best solution
  iter.obj <- max(fitness)
  pos.max <- which(fitness == max(fitness))[1]
  iter.sol <- population.parent[[pos.max]]

  #update best solution
  if(iter.obj > best.obj){
    best.sol <- iter.sol
    best.obj <- iter.obj
  }

  #elitist strategy: replace worst solution by best solution so far
  if(elitist){
    pos.min <- which(fitness == min(fitness))[1]
    population.son[[pos.min]] <- best.sol
    fitness[pos.min] <- best.obj
  }

  #verbose printing
  if(verbose & (count%%10==0 | count==1)) {
    print(paste("population", count))
    sapply(1:npop, function(x) print(c(population.parent[[x]], fitness[x]), digits=0))
  }

  #computing probability of selection
  prob <- fitness^2
  prob <- prob/sum(prob)

```

```

#creating son population
for(i in 1:npop){
  #crossover
  parents <- sample(1:npop, 2, prob = prob)
  v1 <- population.parent[[parents[1]]]
  v2 <- population.parent[[parents[2]]]
  population.son[[i]] <- CrossoverKP1point(v1, v2)
  #mutation
  if(pmut > runif(1)) population.son[[i]] <- MutationKP3points(population.son[[i]])
}

#population son becomes parent
population.parent <- population.son

count <- count + 1
}

return(list(sol=best.sol, obj=best.obj))
}

```

Some comments to this function:

- The function parameters are: instance data dades, population size npop, number of iterations to be performed and probability of mutation pmut. If the logical verbose is set to TRUE, the function prints the population and fitness of each element every ten runs.
- Selection probability is proportional to the square of the fitness function of each instance. Then, all unfeasible solutions will have selection probability equal to zero.
- The GA may be stuck if all solutions of a population are unfeasible. To avoid this, I have included in each element of the starting population a number of items proportional to the ratio weight of knapsack vs. total weight of items.
- If the flag elitist is true, in each generation one of the worst solutions is replaced by the best solution obtained so far.

Generating instances

To test the algorithm, I have generated three instances:

```

InstanceKP10 <- KPInstance01(1000, 10, 40)
InstanceKP50 <- KPInstance01(1000, 50, 40, 2424)
InstanceKP100 <- KPInstance01(1000, 100, 40, 3333)

```

In the case of the KP, we can obtain the optimal solution using linear programming. This function returns the results of linear programming in the same format of the GA function:

```
LPKP <- function(dades){  
  library(Rglpk)  
  n <- length(dades$u)  
  obj <- dades$u  
  mat <- matrix(dades$w, nrow=1)  
  types <- rep("B", n)  
  sol.lp <- Rglpk_solve_LP(obj=obj, mat=mat, dir="<=", rhs=dades$W, types=types,  
max = TRUE)  
  return(list(sol = sol.lp$solution, obj = sol.lp$optimum))  
}
```

Then we proceed to solve instances of size 10 and 50 with linear programming:

```
solPL.KP10 <- LPKP(InstanceKP10)  
## Warning: package 'slam' was built under R version 3.3.2  
solPL.KP50 <- LPKP(InstanceKP50)
```

GA in action

Let's run the genetic algorithm for InstanceKP10 and probability of mutation equal to zero:

```
set.seed(4444)  
solGA.KP10 <- GeneticAlgorithmKP(InstanceKP10, 10, 50, 0, elitist=FALSE,  
verbose=TRUE)  
  
## [1] "population 1"  
## [1] 1 0 0 0 0 0 1 1 1 0 1514  
## [1] 1 1 1 0 1 0 0 0 1 1 0  
## [1] 1 0 0 1 0 1 0 0 1 0 1758  
## [1] 0 0 0 1 1 1 0 0 0 0 1323  
## [1] 0 0 0 1 1 0 1 1 0 0 1532  
## [1] 1 1 0 0 1 0 0 0 0 0 1312  
## [1] 0 0 1 0 1 0 0 1 0 0 0  
## [1] 0 1 1 1 0 1 0 0 0 1 0  
## [1] 1 0 0 0 0 1 0 0 0 0 499  
## [1] 0 0 0 0 0 0 0 1 0 0 158  
## [1] "population 10"  
## [1] 1 0 0 1 0 1 0 0 0 0 952  
## [1] 1 0 0 1 0 1 0 0 1 0 1758  
## [1] 1 0 0 0 1 0 1 0 0 0 1286  
## [1] 1 0 0 1 0 1 0 0 1 0 1758  
## [1] 1 0 0 0 1 1 0 0 1 0 0  
## [1] 1 0 0 1 0 1 0 0 1 0 1758
```

[illegible]

The algorithm gets stuck in the best feasible solution of the initial population.
Let's run the algorithm with nonzero probability of mutation and elitist strategy:

```
set.seed(4444)
solGA.KP10 <- GeneticAlgorithmKP(InstanceKP10, 10, 100, 0.2, elitist=TRUE,
verbose=TRUE)
```

```
## [1] "population 1"
## [1] 1 0 0 0 0 0 1 1 1 0 1514
## [1] 1 1 1 0 1 0 0 0 1 1 1758
## [1] 1 0 0 1 0 1 0 0 1 0 1758
## [1] 0 0 0 1 1 1 0 0 0 0 1323
## [1] 0 0 0 1 1 0 1 1 0 0 1532
## [1] 1 1 0 0 1 0 0 0 0 0 1312
## [1] 0 0 1 0 1 0 0 1 0 0 0
## [1] 0 1 1 1 0 1 0 0 0 1 0
## [1] 1 0 0 0 0 1 0 0 0 0 499
## [1] 0 0 0 0 0 0 0 1 0 0 158
## [1] "population 10"
## [1] 0 0 0 1 0 0 0 0 1 0 1259
## [1] 1 0 0 1 0 0 0 0 1 0 1624
## [1] 1 1 1 1 0 1 0 1 0 0 1846
## [1] 0 0 0 1 1 0 0 0 1 0 0
## [1] 1 0 0 1 1 0 0 0 0 0 1554
## [1] 0 0 0 1 0 1 0 0 1 0 1393
## [1] 1 0 0 1 1 0 0 0 1 0 0
## [1] 0 0 0 1 0 0 0 0 0 0 453
## [1] 0 0 0 1 0 0 1 1 1 0 1602
## [1] 0 0 0 1 0 1 0 0 1 1 1734
## [1] "population 20"
## [1] 0 0 0 1 0 0 0 0 1 1 1600
## [1] 0 0 0 1 0 0 0 0 1 1 1600
## [1] 0 0 0 1 0 0 0 0 1 1 1600
## [1] 1 0 0 1 0 1 0 0 1 1 1846
## [1] 0 0 0 1 0 1 0 0 1 0 1393
## [1] 0 0 0 1 0 1 0 0 1 1 1734
## [1] 0 0 0 1 0 1 0 0 1 1 1734
## [1] 0 1 1 0 0 0 0 0 1 1 0
## [1] 0 0 0 1 0 0 0 0 1 1 1600
## [1] 0 0 0 1 1 0 1 0 1 0 0
## [1] "population 30"
## [1] 0 0 0 1 0 1 0 0 1 1 1734
## [1] 0 0 0 1 0 1 0 0 1 1 1734
## [1] 0 0 0 1 0 1 0 1 1 1 1846
## [1] 0 0 0 1 0 1 0 0 1 1 1734
## [1] 0 0 0 1 0 1 0 0 1 1 1734
## [1] 0 0 1 0 1 1 1 0 1 0
## [1] 0 0 0 1 0 1 0 0 1 1 1734
## [1] 0 0 0 1 0 0 0 0 0 1 794
## [1] 0 0 0 1 0 0 0 0 1 1 1600
```

```

## [1] 0 0 0 1 0 1 0 1 1 1 0
## [1] "population 40"
## [1] 0 0 0 0 1 1 0 1 1 1 1846
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 1 0 1 0 0 1 1 1 0
## [1] 1 1 1 1 0 0 0 1 1 1 0
## [1] 0 0 0 1 0 0 0 0 1 1 1758
## [1] 1 1 1 1 0 0 0 0 1 1 0
## [1] 1 1 1 1 0 0 0 1 1 1 0
## [1] 0 0 0 1 0 0 0 0 1 1 1758
## [1] 0 0 0 1 0 0 1 0 0 1 979
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] "population 50"
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 1 1 1 1 0 0 0 1 1 1 1846
## [1] 0 0 0 1 1 1 1 1 1 1 0
## [1] 0 0 1 1 0 0 0 0 0 0 1518
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] "population 60"
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 1 1 1 1 0 0 0 1 1 1 1846
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 1 1 0 1 1 0
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] "population 70"
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 0 0 0 453
## [1] 0 0 0 0 1 1 0 1 1 1 1846
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] "population 80"
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 1 1 0 1 1 1846
## [1] 0 0 0 1 0 0 0 1 1 1 1758

```



```

## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 0 0 0 453
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 1 1 0 0 0 1 1 1 0
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] "population 90"
## [1] 1 1 1 1 0 0 0 1 1 1 1846
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 1 0 0 1 979
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 1 1 1 1 0 0 0 1 1 1 0
## [1] 1 1 1 1 0 0 0 1 1 1 0
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 1 1 0 0 0 1 0 0 1 0
## [1] 1 1 1 1 0 0 0 1 1 1 0
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] "population 100"
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 1 1 1 1 1 1846
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758
## [1] 0 0 0 1 0 0 0 1 1 1 1758

```

Finally, we will run the algorithm with 1000 iterations and :

```

set.seed(4444)
solGA.KP10 <- GeneticAlgorithmKP(InstanceKP10, 10, 1000, 0.2, elitist=TRUE,
verbose=FALSE)
solGA.KP10

## $sol
## [1] 1 1 0 0 1 1 0 1 0 1
##
## $obj
## [1] 1945

solPL.KP10

## $sol
## [1] 0 1 0 1 0 1 1 1 1 0
##
## $obj
## [1] 1947

```

For the instances of size 50 and 100 we have:

```

set.seed(4444)
solGA.KP50 <- GeneticAlgorithmKP(InstanceKP50, 10, 1000, 0.2, elitist=TRUE,
verbose=FALSE)
solGA.KP50

## $sol
## [1] 1 0 0 1 0 1 1 1 1 1 1 1 1 1 0 0 1 1 0 1 0 1 0 1 1 0 0 0 1 0 1 0 1 0 1
## [36] 1 0 0 0 1 1 1 1 0 1 0 0 0 0 0
##
## $obj
## [1] 12071

solPL.KP50

## $sol
## [1] 1 1 0 1 0 1 1 1 1 0 1 0 0 1 1 0 1 1 1 1 1 0 1 0 0 1 0 1 1 1 0 1
## [36] 1 0 0 0 1 1 1 0 1 1 1 1 0 0 0
##
## $obj
## [1] 12477

set.seed(4444)
solGA.KP100 <- GeneticAlgorithmKP(InstanceKP100, 10, 1000, 0.2, elitist=TRUE,
verbose=FALSE)
solGA.KP100

## $sol
## [1] 0 1 0 0 1 1 0 1 1 0 1 0 0 0 0 0 1 1 1 0 0 0 1 1 0 0 1 0 1 0 1 0 1 0 1
## [36] 1 0 1 1 1 0 0 1 1 0 0 1 0 1 0 1 1 0 0 1 1 0 0 1 1 1 1 0 0 0 0 1 0 0 0
## [71] 0 0 1 1 0 0 1 1 0 1 1 1 1 0 0 0 0 1 0 1 1 0 0 1 0 0 0 0 1 1
##
## $obj
## [1] 23002

```

The optimal solution of the instance with 100 items has objective value equal to 24643. So the genetic algorithm work well solving small instances of the knapsack problem.