MonteCarloSimulation-Genetic Algorithms

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MonteCarlo SImulations

Clear all the environment variables

```
rm(list=ls(all=TRUE))
```

Defining the function for the activity

```
timeRequired = function(number.simulations){
  time = 0
  for(i in 1: number.simulations){
    # Probability for the Full project is 90% and for Part project is 10%
   partOrFull = runif(1,0,1)
   if(partOrFull <= 0.1) {</pre>
      total.module = sample(10:12,1)
   }
   else{
      total.module = sample(20:24,1)
    # Task and the people per module are represented by two vectors randomly.
   tasks.per.module = sample(50:100, total.module, replace = T)
   total.tasks = sum(tasks.per.module)
    # Total time taken for all the tasks: Each task can take any time between 5 to 10 hours to get comp
   tasks.time = sum(sample(5:10,total.tasks,replace = T))
    # Randomly pick the number of resources
   resources = sample(3:7,1,replace = TRUE)
    # Time taken to complete the project
   time.taken.estimate = tasks.time/ resources
 }
 return (time.taken.estimate)
```

Running the simulations

```
simulations = c(10,100,10000, 100000, 200000, 300000, 100000, 200000, 300000)
for (i in simulations) {
   total.project.time = timeRequired(i)
   cat("Total Estimated Time to complete the project with",i," simulations is ",total.project.time,"\n")
}

## Total Estimated Time to complete the project with 10 simulations is 2882.2
## Total Estimated Time to complete the project with 100 simulations is 1787
## Total Estimated Time to complete the project with 10000 simulations is 2153.2
## Total Estimated Time to complete the project with 1e+05 simulations is 915.6667
## Total Estimated Time to complete the project with 2e+05 simulations is 2936.5
## Total Estimated Time to complete the project with 1e+05 simulations is 2814.2
## Total Estimated Time to complete the project with 2e+05 simulations is 2330.2
## Total Estimated Time to complete the project with 3e+05 simulations is 2581.6
```

Genetic Algorithm

Clear all the environment variables

```
rm(list = ls(all=TRUE))
```

Loading the required package

```
## Warning: package 'GA' was built under R version 3.4.1
## Loading required package: foreach
## Warning: package 'foreach' was built under R version 3.4.1
## Loading required package: iterators
## Package 'GA' version 3.0.2
## Type 'citation("GA")' for citing this R package in publications.
```

Setting the working directory

```
setwd("I:/DATA-SCIENCE/Insofe/Assignments/MonteCarlo-GA/assignment-mcs-ga-Thomas-K-John-master")
```

Reading the file

```
tsp.distance.data = read.csv("distanceinfo.csv", header =TRUE)
```

Implementation of the algorithm

```
rownames(tsp.distance.data) = colnames(tsp.distance.data)
tsp.distance.data = as.matrix(tsp.distance.data)
tour.length = function(tour, dist.matrix) {
 tour = c(tour, tour[1])
 route = embed(tour, 2)[,2:1]
 sum(dist.matrix[route])
tsp.fitness = function(tour, ...) 1/tour.length(tour, ...)
tsp.solution.model = ga(type="permutation", fitness=tsp.fitness, dist.matrix=tsp.distance.data, min=1,
summary(tsp.solution.model)
## +-----
          Genetic Algorithm
## +----+
##
## GA settings:
## Type
                     = permutation
## Population size = 50
## Number of generations = 100
## Elitism
## Crossover probability = 0.8
## Mutation probability = 0.2
##
## GA results:
## Iterations = 100
## Fitness function value = 0.000856898
## Solutions =
## x1 x2 x3 x4 x5 x6 x7 x8 x9 x10
## [1,] 9 2 4 1 3 6 8 7 10
## [2,] 10 5 9 2 4 1 3 6 8
```