

EPC3

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A)

```
def initPopulation(self, num_individuals):  
  
    num_individuals  
  
    population = []  
    #Cria todos os individuos e insere na populacao inicial  
    for i in range(num_individuals):  
        individual = np.random.binomial(1,0.5,self.individual_size)  
        #print(individual)  
        population.append(individual.tolist())  
    return population
```

B)

```
def fitness(self, population):  
  
    print(population)  
    fitnessPop=[]  
    #calculate the fitness for each individual of population  
    for individual in population:  
        fitnessPop.append(self.getFitness(individual))  
  
    #print(fitnessPop)  
    return fitnessPop
```

C)

```
def __selection(self):

    sorted_population = sorted(self.population, key=self.problem.getFitness,
reverse=True)
    pop_fit = self.problem.fitness(sorted_population)

    prob_fit = []
    for individual in pop_fit:
        prob_fit.append(1.0*individual/np.sum(pop_fit))

    #Get the cumulative sum of the probabilities.
    cumSumP = np.cumsum(prob_fit)
    #Get our random numbers - one for each column.
    randomNumber = np.random.rand()
    #Get the values from A.
    #If the random number is less than the cumulative probability then
    #that's the number to use from A.
    for i, total in enumerate(cumSumP):
        if randomNumber < total:
            break
    self.population = sorted_population
    selected = self.population[i]
    #Display it. Uncomment for log.
    print("Selected individual [%d] = %s" %(i,selected))
    return selected
```

D)

```
def __crossover(self, individual_x, individual_y):

    n=self.problem.getIndividualSize()

    c = int(np.random.uniform(0,n-1))
    print("crossing point: %d" %c)

    new_individual_x=[]
    new_individual_y=[]

    # concatenate the two fathers in the C element chosen randomly
    for gene in range(c):
        new_individual_x.append(individual_x[gene])
        new_individual_y.append(individual_y[gene])
    for gene in range(c,n):
        new_individual_x.append(individual_y[gene])
        new_individual_y.append(individual_x[gene])

    return new_individual_x, new_individual_y
```

E)

```
def __mutation(self, individual):  
  
    if self.__mutationTest():  
        randomPosition = int(np.random.uniform(0, self.problem.getIndividualSize()-1))  
        print('Mutation at position: %d' % randomPosition)  
        #get a random value for changing in the individual position selected before  
        randomValue =  
        np.random.uniform(self.problem.getMinGeneSymbol(), self.problem.getMaxGeneSymbol())  
  
        if (randomValue <= 0.5):  
            randomValue = int(randomValue)  
        else:  
            randomValue = int(randomValue+1)  
  
        print('New gene value: %d' % randomValue)  
        individual[randomPosition] = randomValue  
    return individual
```

3-

MaxGeneration = 100

4-

Ótimo da função: $x_1 = 1$ e $x_2 = 1$, que gera $f(x_1, x_2) = 0$

Ótimo obtido: $x_1 = 0.996078$ e $x_2 = 0.992157$, que gerou $f(x_1, x_2) = 1.54024e-05$

População: 100 indivíduos

Gerações: 100 gerações

5-

Ótimo obtido: $x_1 = 0.996078$ e $x_2 = 0.992157$, que gerou $f(x_1, x_2) = 1.54024e-05$

População: 350 indivíduos

Gerações: 150 gerações