JawadAhmed_20P-0165

May 10, 2023

Lab 12: Genetic Algorithms

```
1.0.1 Name: Jawad Ahmed
1.0.2 Roll No: 20P-0165
1.0.3 Section: BCS-6A
In [1]: import numpy as np
In [2]: # Initialize a 2D numpy array with 3 rows and 4 columns filled with integer zeros
        arr = np.zeros((4, 4), dtype=int)
In [3]: arr
Out[3]: array([[0, 0, 0, 0],
                [0, 0, 0, 0],
                [0, 0, 0, 0],
                [0, 0, 0, 0]])
1.0.4 Step 1: Initial Population
Select any random states as initial population i.e.
   No. of population is a random or your choice.
```

Values in each population represents a random state.

```
In [4]: def fitness_function(arr):
            return np.count_nonzero(arr == 1)
In [5]: def generate_initial_chromosomes(n):
            arr = np.zeros((4,4), dtype=int)
            arr = arr.reshape(16)
            chromosomes = []
            for i in range(n):
                random_numbers = np.random.randint(0, 16, size=10)
                for random_number in random_numbers:
                    arr[random_number] = 1
                chromosomes.append(arr)
                arr = np.zeros((4, 4), dtype=int)
                arr = arr.reshape(16)
            return chromosomes
```

```
In [6]: chromosomes = generate_initial_chromosomes(30)
In [7]: chromosomes
Out[7]: [array([1, 1, 0, 1, 1, 0, 0, 0, 1, 1, 0, 1, 0, 1, 0, 1]),
        array([1, 1, 0, 0, 1, 1, 1, 0, 0, 1, 0, 1, 0, 0, 0, 1]),
         array([1, 1, 0, 0, 1, 0, 0, 1, 1, 0, 0, 1, 1, 1, 0, 1]),
        array([0, 1, 1, 1, 0, 0, 0, 1, 0, 0, 1, 0, 1, 0, 1, 0]),
         array([0, 1, 0, 0, 0, 0, 1, 1, 0, 1, 1, 1, 0, 0, 1]),
         array([1, 1, 0, 0, 1, 1, 1, 1, 0, 0, 0, 0, 1, 1, 0, 0]),
         array([0, 0, 1, 1, 1, 0, 0, 0, 1, 1, 0, 0, 0, 1, 1, 0]),
         array([0, 1, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0]),
         array([0, 1, 0, 0, 1, 1, 1, 0, 0, 0, 1, 1, 1, 0, 1, 0]),
         array([1, 0, 0, 0, 1, 0, 0, 1, 0, 1, 0, 1, 1, 0, 0]),
         array([1, 0, 1, 1, 1, 0, 1, 0, 0, 0, 1, 1, 1, 0, 0, 0]),
         array([1, 0, 0, 1, 0, 1, 1, 0, 1, 0, 0, 0, 1, 1, 1, 0]),
         array([0, 0, 1, 1, 1, 1, 1, 0, 0, 1, 0, 0, 0, 0, 1, 1]),
         array([0, 0, 1, 0, 0, 0, 1, 0, 1, 0, 1, 1, 1, 1, 1, 0]),
         array([0, 1, 0, 1, 0, 1, 0, 0, 0, 1, 0, 1, 1, 0, 1, 0]),
         array([0, 0, 0, 0, 0, 0, 1, 0, 1, 1, 0, 1, 1, 1]),
         array([0, 0, 0, 0, 0, 0, 1, 1, 1, 0, 1, 1, 1, 0, 0]),
        array([1, 0, 0, 0, 1, 0, 0, 1, 1, 0, 1, 1, 0, 0, 1]),
         array([0, 0, 1, 0, 0, 0, 0, 1, 0, 1, 0, 1, 0, 1, 0, 0]),
        array([0, 0, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 0, 0, 0, 0]),
         array([1, 0, 0, 0, 0, 0, 1, 1, 1, 0, 1, 1, 1, 0, 0]),
         array([1, 0, 0, 0, 1, 0, 1, 1, 1, 1, 0, 1, 0, 1, 1, 0]),
         array([1, 0, 0, 0, 1, 0, 1, 1, 1, 0, 1, 0, 1, 1, 0, 0]),
         array([0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 1, 1, 0, 1]),
        array([1, 0, 1, 1, 0, 1, 0, 0, 1, 1, 0, 1, 0, 0, 0, 1]),
         array([1, 0, 1, 0, 1, 0, 1, 0, 1, 1, 0, 0, 0, 0]),
         array([0, 0, 1, 0, 1, 0, 1, 1, 0, 1, 0, 0, 1, 0, 0, 1]),
         array([0, 1, 1, 0, 0, 0, 0, 1, 1, 1, 1, 1, 1, 0, 0, 1]),
         array([1, 1, 0, 0, 1, 0, 0, 1, 0, 0, 0, 1, 1, 1, 1]),
         array([0, 0, 1, 0, 1, 0, 1, 0, 1, 0, 0, 0, 1, 1, 1, 1])]
```

1.0.5 Step 2: Selection of Parents

- 1. Parents will be selected random from the population in the first step.
- 2. High fitness value population will have high chances of selection as a parent i.e. P4 will have high weightage to be selected as randomly Selection_Probability P(i) = Fitness of P(i)/Total Fitness of all Populations

```
In [8]: def calculate_probabilities(chromosomes):
    #    Calculate Total Fitness of all the population
    total_fitness_all_population = 0
    for chromosome in chromosomes:
        total_fitness_all_population += fitness_function(chromosome)
    #    Calculate selection probabitlies
    selection_probabilities = []
```

```
for chromosome in chromosomes:
                selection_probabilities.append((fitness_function(chromosome)) / total_fitness_
            return selection_probabilities
In [9]: def convert_chromosomes_to_indices(chromosomes):
            indices_chromosomes = []
            for i in range(len(chromosomes)):
                indices_chromosomes.append(i)
            return indices_chromosomes
In [10]: def roulette_wheel_selection(chromosomes):
             selection_probabilities = calculate_probabilities(chromosomes)
             indices_chromosomes = convert_chromosomes_to_indices(chromosomes)
             selected_parents_indices = np.random.choice(indices_chromosomes, 2, p=selection_p:
             selected_parents = []
             selected_parents.append(chromosomes[selected_parents_indices[0]])
             selected_parents.append(chromosomes[selected_parents_indices[1]])
             return selected_parents
In [11]: selected_parents = roulette_wheel_selection(chromosomes)
In [12]: selected_parents
Out[12]: [array([0, 1, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0]),
          array([0, 1, 0, 0, 0, 0, 1, 1, 0, 1, 1, 1, 0, 0, 1])]
1.0.6 Step 3: Modification
```

Crossover: 1. This is called convergence step because it will generate children (new states) 2. Create children from meeting of the parents 3. Generate two children/successor from two parents (new version of genetic algorithm produce one child instead)

```
In [13]: def generate_childs_using_crossover(selected_parents):
             crossover_point = np.random.randint(0, len(selected_parents[0]))
               Create child
             child_one = np.concatenate((selected_parents[0][:crossover_point], selected_parent
             child_two = np.concatenate((selected_parents[1][:crossover_point], selected_paren
             childs = []
             childs.append(child_one)
             childs.append(child_two)
             return childs
In [14]: childs = generate_childs_using_crossover(selected_parents)
In [15]: childs
Out[15]: [array([0, 1, 0, 0, 0, 0, 1, 1, 0, 1, 1, 1, 0, 0, 1]),
          array([0, 1, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0])]
```

1.0.7 Mutation:

- 1. Apply mutation on children in order to get the new/updated state (assuming it will quickly lead us to goal)
- 2. Mutation probability or mutation rate is fixed and chosen a very small value i.e. 0.01 means generate a random no. between (1-100), mutate the child if random no. is 1 else skip. i.e. 0.2 means generate a random no. between (1-10), mutate the child if random no. is 1 or 2. Skip otherwise
- 3. Mutation rate will be applied and checked for each digit/char in a population i.e. You will keep repeating this process for each array value in a single population means 16 times random no. will be generated and checked respectively.

```
In [16]: def mutation(childs, mutation_rate = 0.2):
             random_number = 0
             if (mutation_rate == 0.01):
                 random_number = np.random.randint(1, 100)
                 for child in childs:
                     for i in range(len(child)):
                           If random number is 1 mutate the child
         #
                         if random number is 1:
                             if (childs[i] is 1):
                                 childs[i] = 0
                             else:
                                 childs[i] = 1
             else:
                 random_number = np.random.randint(1, 10)
                 for child in childs:
                     for i in range(len(child)):
                           If random number is 1 or 2 mutate it
         #
                         if random_number is 1 or random_number is 2:
                             if childs[i] is 1:
                                 childs[i] = 0
                             else:
                                  childs[i] = 1
In [17]: childs
Out[17]: [array([0, 1, 0, 0, 0, 0, 0, 1, 1, 0, 1, 1, 1, 0, 0, 1]),
          array([0, 1, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0])]
In [18]: mutation(childs)
In [19]: childs
Out[19]: [array([0, 1, 0, 0, 0, 0, 0, 1, 1, 0, 1, 1, 1, 0, 0, 1]),
          array([0, 1, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0])]
```

1.0.8 Step 4: Evaluation

- 1. Compute the fitness values of newly generated children
- 2. Apply the goal test
- 3. Replace the old population with newly created population having new children
- 4. Repeat the steps 1-4 if goal is not found.

```
In [20]: def evaluation(goal_fitness_value = 16, no_of_iterations = 10000):
             new_fitness_value = 0
             goal_state = np.array([])
             while(new_fitness_value <= goal_fitness_value and a < no_of_iterations):</pre>
                 chromosomes = generate_initial_chromosomes(30)
                 selected_parents = roulette_wheel_selection(chromosomes)
                 childs = generate_childs_using_crossover(selected_parents)
                 if (fitness_function(childs[0]) > fitness_function(childs[1])) and (fitness_function(childs[1]))
                     new_fitness_value = fitness_function(childs[0])
                     goal_state = childs[0]
                 elif (fitness_function(childs[1]) > new_fitness_value):
                     new_fitness_value = fitness_function(childs[1])
                      goal_state = childs[1]
                 a += 1
             return goal_state, new_fitness_value
In [21]: evaluation()
Out[21]: (array([0, 1, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 0, 1]), 13)
In []:
In []:
```