

BIRLA INSTITUTE OF TECHNOLOGY & SCIENCE, PILANI
WORK INTEGRATED LEARNING PROGRAMMES
DSECL ZG557, Artificial and Computational Intelligence

Familiarize with the working of Local Search algorithms:
Genetic Algorithm

Tool: Python

Libraries Used: numpy, sys

Sample Problem: The N Queen is the problem of placing N chess queens on an N×N chessboard so that no two queens attack each other.

Queens can attack either on the same row, on the same column or across the diagonal.

If none of the queens are located on the same row, same column or across the diagonals for each other then we call the positioning/configuration of Queens to be a solution.

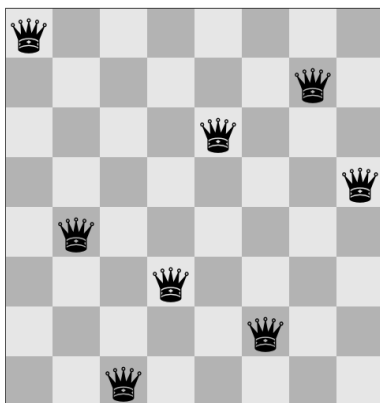
Input: Population with multiple Board Configurations of N-Queens

Example: [1,2,3,7,5,0,4,6]

This represents 0th queen lies in 1st row, 1st queen lies in 2nd row, 2nd queen lies in 3rd row, 3rd queen lies in 7th row, 4th queen lies in 5th row, 5th queen lies in 0th row, 6th queen lies in 4th row and 7th queen lies in 6th row.

Output: A possible configuration of queens such that none of the attack each other.

Example (8-Queens):



[0,4,7,5,2,6,1,3]

Explanation: If 0th queen lies in 0th row, 1st queen lies in 4th row, 2nd queen lies in 7th row, 3rd queen lies in 5th row, 4th queen lies in 2nd row, 5th queen lies in 6th row, 6th queen lies in 1st row and 7th queen lies in 3rd row, then none can attack each other.

Implementation:

```
import numpy as np
import sys
```

```
nQueens = 8
STOP_CTR = 28
MUTATE = 0.01
MUTATE_FLAG = True
# MAX_ITER = 100000
MAX_ITER = 1000000
POPULATION = None
```

```
class BoardPosition:
    def __init__(self):
        self.sequence = None
        self.fitness = None
        self.survival = None

    def setSequence(self, val):
        self.sequence = val

    def setFitness(self, fitness):
        self.fitness = fitness

    def setSurvival(self, val):
        self.survival = val

    def getAttr(self):
        return {'sequence': self.sequence, 'fitness': self.fitness, 'survival':
self.survival}
```

```
def fitness(chromosome=None):
    """
    returns 28 - <number of conflicts>
    to test for conflicts, we check for
    -> row conflicts
    -> columnar conflicts
    -> diagonal conflicts

    The ideal case can yield upto 28 arrangements of non attacking pairs.
    for iteration 0 -> there are 7 non attacking queens
    for iteration 1 -> there are 6 no attacking queens ..... and so on

    Therefore max fitness = 7 + 6+ 5+4 +3 +2 +1 = 28

    hence fitness val returned will be 28 - <number of clashes>

    """
    # calculate row and column clashes
    # just subtract the unique length of array from total length of array
    # [1,1,1,2,2,2] - [1,2] => 4 clashes
    clashes = 0
    row_col_clashes = abs(len(chromosome) - len(np.unique(chromosome)))
    clashes += row_col_clashes
```

```

# calculate diagonal clashes
for i in range(len(chromosome)):
    for j in range(len(chromosome)):
        if (i != j):
            dx = abs(i - j)
            dy = abs(chromosome[i] - chromosome[j])
            if (dx == dy):
                clashes += 1

return 28 - clashes

def generateChromosome():
    # randomly generates a sequence of board states.
    global nQueens
    init_distribution = np.arange(nQueens)
    np.random.shuffle(init_distribution)
    return init_distribution

def generatePopulation(population_size=100):
    global POPULATION

    POPULATION = population_size

    population = [BoardPosition() for i in range(population_size)]
    for i in range(population_size):
        population[i].setSequence(generateChromosome())
        population[i].setFitness(fitness(population[i].sequence))

    summation_fitness = np.sum([x.fitness for x in population])
    for each in population:
        each.survival = each.fitness / (summation_fitness * 1.0)

    return population

def getParent():
    globals()
    parent1, parent2 = None, None
    # parent is decided by random probability of survival.
    # since the fitness of each board position is an integer >0,
    # we need to normaliza the fitness in order to find the solution

    while True:
        parent1_random = np.random.rand()
        parent1_rn = [x for x in population if x.survival <= parent1_random]
        try:
            parent1 = parent1_rn[0]
            break
        except:
            pass

    while True:
        parent2_random = np.random.rand()
        parent2_rn = [x for x in population if x.survival <= parent2_random]

```

```

        try:
            t = np.random.randint(len(parent2_rn))
            parent2 = parent2_rn[t]
            if parent2 != parent1:
                break
            else:
                continue
        except:
            continue

    if parent1 is not None and parent2 is not None:
        return parent1, parent2
    else:
        sys.exit(-1)

def reproduce_crossover(parent1, parent2):
    globals()
    n = len(parent1.sequence)
    c = np.random.randint(0, n)
    child = BoardPosition()
    child.sequence = []
    child.sequence.extend(parent1.sequence[0:c])
    child.sequence.extend(parent2.sequence[c:])
    child.setFitness(fitness(child.sequence))
    return child

def mutate(child):
    """
    - according to genetic theory, a mutation will take place
    when there is an anomaly during cross over state

    - since a computer cannot determine such anomaly, we can define
    the probability of developing such a mutation

    """
    if child.survival < MUTATE:
        c = np.random.randint(8)
        child.sequence[c] = np.random.randint(8)
    return child.sequence

def GA(iteration):
    print(" #" * 10, "Executing Genetic generation : ", iteration, " #" * 10)
    globals()
    newpopulation = []
    for i in range(len(population)):
        parent1, parent2 = getParent()
        # print "Parents generated : ", parent1, parent2

        child = reproduce_crossover(parent1, parent2)
        newpopulation.append(child)

    summation_fitness = np.sum([x.fitness for x in newpopulation])
    for each in newpopulation:
        each.survival = each.fitness / (summation_fitness * 1.0)

```

```

if (MUTATE_FLAG):
    for each in newpopulation:
        presentVal = each.sequence
        mightBeChangedVal = mutate(each)
        if presentVal!=mightBeChangedVal:
            each.sequence = presentVal
            each.fitness = each.setFitness(fitness(each.sequence))

summation_fitness = np.sum([x.fitness for x in newpopulation])
for each in newpopulation:
    each.survival = each.fitness / (summation_fitness * 1.0)

return newpopulation

def stop():
    globals()

    fitnessvals = [pos.fitness for pos in population]
    if STOP_CTR in fitnessvals:
        return True
    if MAX_ITER == iteration:
        return True
    return False

population = generatePopulation(100)

iteration = 0
while not stop():
    # keep iteratin till you find the best position
    population = GA(iteration)
    iteration += 1

print("Iteration Number: ", iteration)
for each in population:
    if each.fitness == 28:
        print(each.sequence)

```

Output :

[5, 2, 0, 7, 4, 1, 3, 6]

Output Explanation: The output shows which queen would lie in which row so that there are no clashes. 0th queen in 5th row, 1st queen in 2nd row and so on. Refer table below

Column	Row
0	5
1	2
2	0
3	7
4	4
5	1
6	3
7	6

Screenshot:

[illegible]

Lab Exercises:

1. Experiment with number of queens as 16
2. Experiment with population size of 500, 1000
3. Experiment with different MUTATE values.
4. Implement cryptarithmic using Genetic Algorithm