## Analysing the Convergence Rate of Genetic Algorithms

CS-3111

Dr Sudip Sanyal

#### **Group Members**

Arjun Bakshi - BT18GEC134

Somanshu Singh - BT18GCS111

Yukta Sharma - BT18GCS181

#### The Basic Problem

#### **Most Dominating Set of Queens Problem**

Given a squared chess board we have to find the position of queens such that they dominate/threaten the maximum squares of the board.

 $A = \begin{bmatrix} 00000110 & 00010000 \end{bmatrix}$ 

1	2	3	4
5	6	7	8
9	10	11	12
13	14	15	16

#### Genetic Algorithm

- An approach towards solving a problem by considering a large population of possible solutions and then applying the theory of evolution on the population.
- This yields fitter and fitter generations.

#### MDSQP

- Analogous to a popular classical puzzle game called the n-queens problem.
- With the use of this problem, we aim to test out various genetic operators and analyze their results.

#### Expressed as an equation:

## Methodology

$$E(m(H,t+1)) \geq \frac{m(H,t)f(H)}{a_t}[1-p]$$

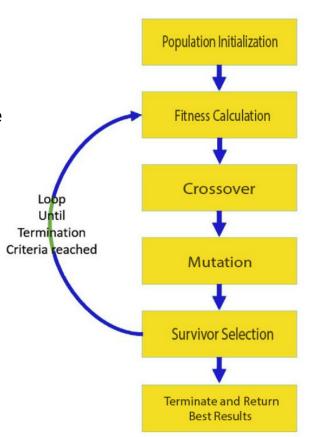
**Fitness Calculation** - Calculating the fitness of each chromosome in the population.

**Selection -** Selecting the best/strongest chromosomes in the population for crossover.

**CrossOver**: Selecting the best features of the parent to be added in the offspring.

**Mutation** -Ensuring the same population is not passed on. Deliberately Inserting some variations in some individuals of the population

**Survivor Selection**: Removing the weakest of the population as each generation comes along.

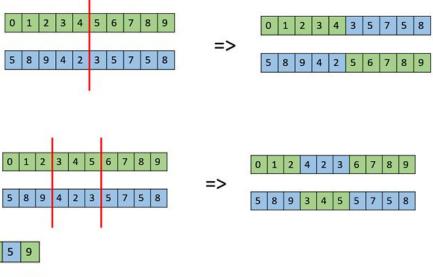


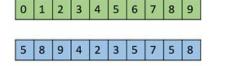
#### **Types of Crossovers:**

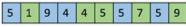
1. Single Point Crossover

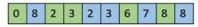
=>

- 2. Two point Crossover
- 3. Uniform Crossover



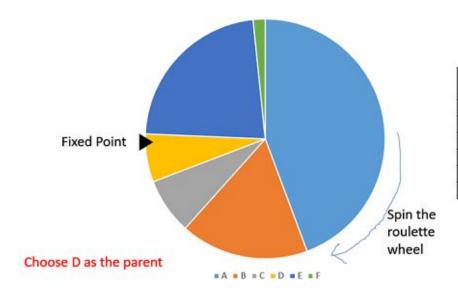




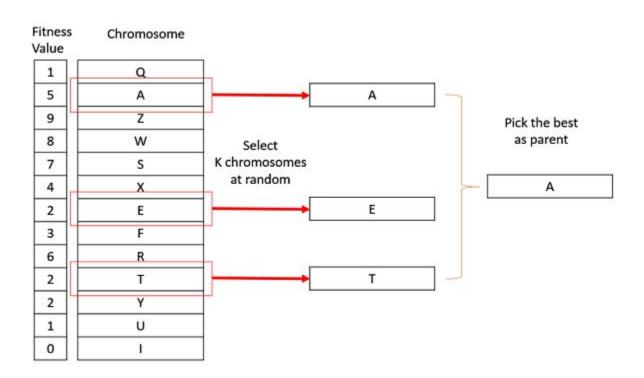


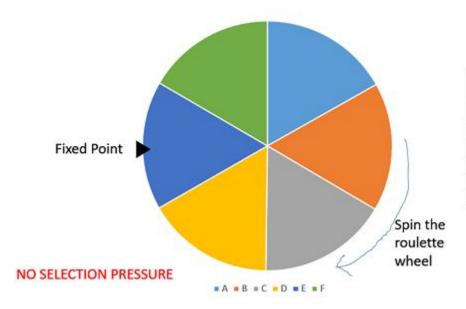
#### **Types of Selections:**

- Elitist selection
- Roulette-wheel selection
- 3. Tournament selection
- 4. Rank Based Selection



Chromosome	Fitness Value	
Α	8.2	
В	3.2	
С	1.4	
D	1.2	
E	4.2	
F	0.3	





Chromosome	Fitness Value
Α	8.1
В	8.0
С	8.05
D	7.95
E	8.02
F	7.99

#### **Types of Mutation:**

- 1. Bit Flip Mutation
- 2. Swap Mutation
- 3. Scramble Mutation

### Libraries

- 1. **GeneAL** for genetic algorithm implementation
- Matplotlib to plot and analyze the performance of combination of genetic operators.

## Deliverables

- 1. Analysing the convergence rate by using a varied combination of genetic operators.
- 2. Graphical Representation of comparisons.
- 3. Details report in addition to current literature reviewed by us.(given in report)

## Final Presentation

### **Constraint Handling**

- 1. In order to prevent a single position from exceeding the boards limit. (Say 20X20 has 400 positions but after mutation the string returned is 11011110 then it exceeds 400), we have coded the mutations after an intervals such that it does not exceed the board limit.
- 2. Same problem occurred after crossmutations also, hence we coded the position in sets of different strings to avoid such an occurrence.
- 3. Eg ['01101101', '00001110', '10001101', '10000010', '00111001', '00101010']
- 4. Hence we did not have to attach any penalty to any solution.

#### **Fitness Function**

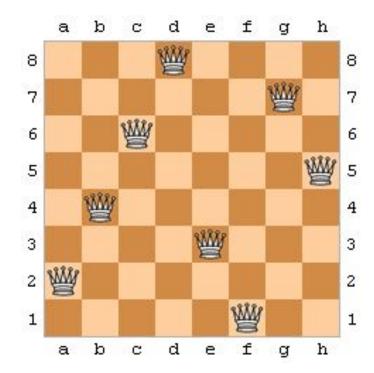
For each solution a fitness is defined as the number of blocks all the queens can cover on the board. The more the number of blocks a solution covers, the better it suits the problem statement.

In code:

def fitness(sol\_list, n)

sol\_list - a particular solution

n - board size



#### Roulette Wheel

We took the best ten solutions and assigned them the parents role on the basis of the roulette wheel which was biased towards allotting fitter parents.

```
# 8. Runs the roulette wheel once on a set of ten solutions
# Solutions must be ordered in decreasing order of fitness
def run roulette wheel(best ten):
    r = random.randint(0,360) #degeree of pointer after each spin
    chosen two = []
    for i in range(15):
        if(0<=r<=140):
            chosen two =[best ten[0],best ten[1]]
        elif(141<=r<=220):
            chosen two =[best ten[2],best ten[3]]
        elif(221<=r<=286):
            chosen two =[best ten[4],best ten[5]]
        elif(287<=r<=338):
            chosen two =[best_ten[6],best_ten[7]]
        elif(339<=r<=360):
            chosen two =[best ten[8],best ten[9]]
    return chosen two
```

#### **Elitist Selection**

The best two chromosomes are preserved in the next generation.
This is done so that the best two solutions are never lost due to crossmutaions.

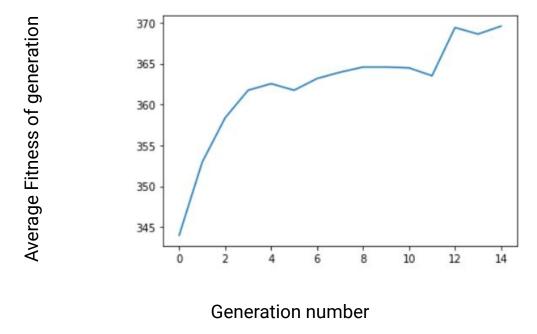
```
#12. Flitist Selection
# a,b are best solution of current generation
def elitist selection(a,b):
   offsprings=[]
    #Preserving the parents(parents fed to this function are the best two of the generation)
   offsprings.append(a)
   offsprings.append(b)
    #single point crossover
   for i in range(5):
        x = deepcopy(b)
        y = deepcopy(a)
        r = random.randint(0,len(a)-1)
        x[:r] = a[:r]
       y[:r] = b[:r]
        #print(r)
       offsprings.append(x)
       offsprings.append(v)
    #double-point crossover
   for i in range(9):
        c = deepcopy(b)
        d = deepcopy(a)
        p = random.randint(1,len(b)-2)
        q = random.randint(1,len(b)-2)
        c[p] = a[q]
       d[q] = b[p]
       offsprings.append(c)
        offsprings.append(d)
   return offsprings
```

#### **Self-Made Selection**

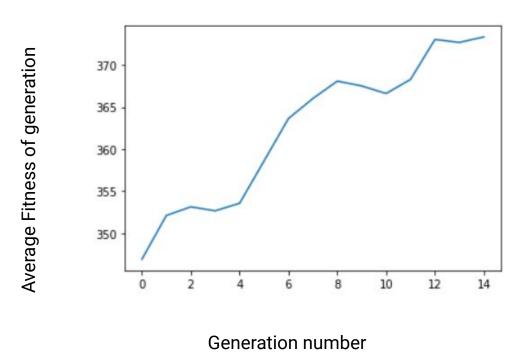
Very similar to elitist selection. However, here the two best are not preserved, they are used to derive the whole of the next generation with a mix of single point and double point cross-overs.

```
#6. Crossmutation(a,b) - returns generation of size 30 with a,b as parent
    - chosen parent chromosome
    - chosen parent chromosome
def crossmutate only two parents(a,b):
    offsprings=[]
    #single point crossover
    for i in range(5):
        x = deepcopy(b)
        y = deepcopy(a)
        r = random.randint(1,len(a)-1)
        x[:r] = a[:r]
        y[:r] = b[:r]
        #print(r)
        offsprings.append(x)
        offsprings.append(y)
    #double-point crossover
   for i in range(10):
        c = deepcopy(b)
        d = deepcopy(a)
        p = random.randint(0,len(b)-1)
        q = random.randint(0,len(b)-1)
        c[p] = a[q]
        d[q] = b[p]
        offsprings.append(c)
        offsprings.append(d)
   return offsprings
```

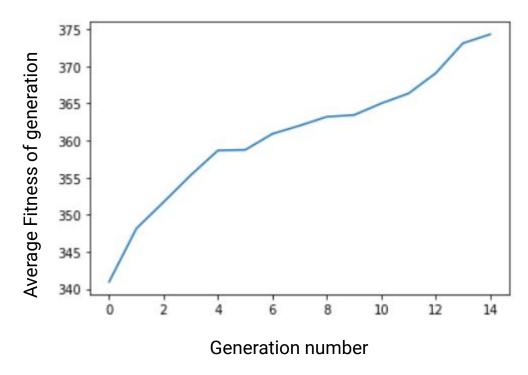
#### **Elitist Selection**



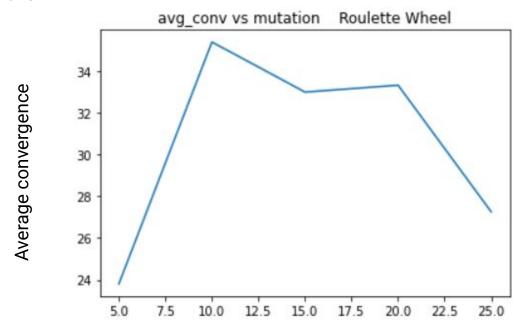
#### **Self-Made Selection**



#### **Roulette-Wheel Selection**

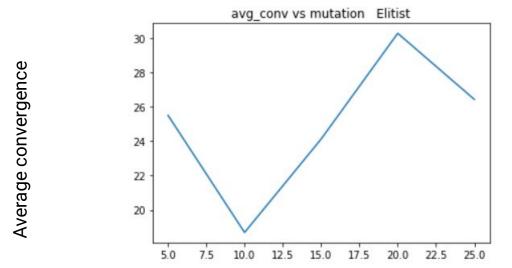


#### **Roulette-Wheel Mutation**



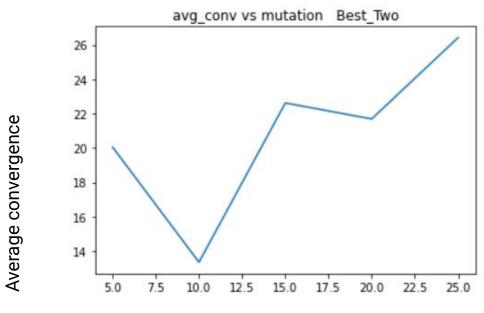
Number of mutations after each generation

#### **Elitist Selection**



Number of Mutations per generation

#### **Self-Made Selection**



Number of mutation in each generation

#### References

#### Slide 2(image)

https://www.semanticscholar.org/paper/A-Genetic-Algorithm-Based-Approach-for-Solving-the-Alharbi-Venkat/216290df1c920d8a68bcbda7600ae93add6ac4a5

#### Slide 4(image)

https://www.researchgate.net/publication/309770246\_A\_Study\_on\_Genetic\_Algorithm\_and\_its\_Applications)

#### Slide 6-10(image)

https://www.tutorialspoint.com/genetic\_algorithms

# Thank you