LAB : 3

OBJECTIVE :

WAP to implement the genetic algorithm.

Requirements :

- Windows/ Mac / Linux Pc
- JDK installed (here using JDK 15)
- Text Editor / IDE (here using VS Code)

Problem Statement :

Create a random pouulation of size 'n' , where each individual (technically a chromosome, which is a solution also) is represented by a binary string of 0's and 1's (initialized randomly). In other words each gene can have two values 0 and 1 , and string of 'n' genes represents an individual. We nedd to apply Algorithm (GA) to get a solution with all 1's.

Implementation :

Individual.java :

```
package Lab3;
import java.lang.Math;

public class Individual {
    private int[] chromosome;
    private double fitness = -1;

    public Individual(int[] chromosome) {
        this.chromosome = chromosome;
    }

    public Individual(int chromosomeLength) {
        this.chromosome = new int[chromosomeLength];
```

```
        for (int gene = 0; gene < chromosomeLength; gene++) {
            if (0.5 < Math.random()) {
                this.setGene(gene, 1);
            } else {
                this.setGene(gene, 0);
            }
        }
    }

    public int[] getChromosome() {
        return this.chromosome;
    }
    public int getChromosomeLength() {
        return this.chromosome.length;
    }

    public void setGene(int offset, int gene) {
        this.chromosome[offset] = gene;
    }

    public int getGene(int offset) {
        return this.chromosome[offset];
    }

    public void setFitness(double fitness) {
        this.fitness = fitness;
    }

    public double getFitness() {
        return this.fitness;
    }

    public String toString() {
        String output = "";
        for (int gene = 0; gene < this.chromosome.length; gene++) {
            output += this.chromosome[gene];
        }
```

```
        return output;
    }
}
```

Populaton.java :

```
package Lab3;

import java.util.Arrays;
import java.util.Comparator;
import java.util.Random;

public class Population {
    private Individual population[];
    private double populationFitness = -1;

    public Population(int populationSize) {
        this.population = new Individual[populationSize];
    }

    public Population(int populationSize, int chromosomeLength) {
        this.population = new Individual[populationSize];

        for (int individualCount = 0; individualCount < populationSize;
individualCount++) {
            Individual individual = new Individual(chromosomeLength);
            this.population[individualCount] = individual;
        }
    }

    public Individual[] getIndividuals() {
        return this.population;
    }

    public Individual getFittest(int offset) {
        Arrays.sort(this.population, new Comparator<Individual>() {
            @Override
            public int compare(Individual o1, Individual o2) {
                if (o1.getFitness() > o2.getFitness()) {
                    return -1;
                } else if (o1.getFitness() < o2.getFitness()) {
```

```
                return 1;
            }
            return 0;
        }
    });

    return this.population[offset];
}

public void setPopulationFitness(double fitness) {
    this.populationFitness = fitness;
}
public double getPopulationFitness() {
    return this.populationFitness;
}
public int size() {
    return this.population.length;
}

public Individual setIndividual(int offset, Individual individual) {
    return population[offset] = individual;
}

public Individual getIndividual(int offset) {
    return population[offset];
}

public void shuffle() {
    Random rnd = new Random();
    for (int i = population.length - 1; i > 0; i--) {
        int index = rnd.nextInt(i + 1);
        Individual a = population[index];
        population[index] = population[i];
        population[i] = a;
    }
}
}
```

GeneticAlgorithm.java :

```
package Lab3;

public class GeneticAlgorithm {
    private int populationSize;
    private double mutationRate;
    private double crossoverRate;
    private int elitismCount;

    public GeneticAlgorithm(int populationSize, double mutationRate, double crossoverRate, int elitismCount) {
        this.populationSize = populationSize;
        this.mutationRate = mutationRate;
        this.crossoverRate = crossoverRate;
        this.elitismCount = elitismCount;
    }

   Population population = new Population(this.populationSize, chromosomeLength);
        return population;
    }

    public double calcFitness(Individual individual) {

        int correctGenes = 0;
        for (int geneIndex = 0; geneIndex < individual.getChromosomeLength();
geneIndex++) {
            if (individual.getGene(geneIndex) == 1) {
                correctGenes += 1;
            }
        }
        double fitness = (double) correctGenes / individual.getChromosomeLength();
        individual.setFitness(fitness);
        return fitness;
    }

    public void evalPopulation(Population population) {
        double populationFitness = 0;
        for (Individual individual : population.getIndividuals()) {
            populationFitness += calcFitness(individual);
        }
        population.setPopulationFitness(populationFitness);
    }
```

```
    public boolean isTerminationConditionMet(Population population) {
        for (Individual individual : population.getIndividuals()) {
            if (individual.getFitness() == 1) {
                return true;
            }
        }

        return false;
    }

    public Individual selectParent(Population population) {
        Individual individuals[] = population.getIndividuals();
        // Spin roulette wheel
        double populationFitness = population.getPopulationFitness();
        double rouletteWheelPosition = Math.random() * populationFitness;
        // Find parent
        double spinWheel = 0;
        for (Individual individual : individuals) {
            spinWheel += individual.getFitness();
            if (spinWheel >= rouletteWheelPosition)
                return individual;
         }
        return individuals[population.size() - 1];
    }

    public Population crossoverPopulation(Population population) {
        Population newPopulation = new Population(population.size());
        for (int populationIndex = 0; populationIndex < population.size();
populationIndex++) {
            Individual parent1 = population.getFittest(populationIndex);
            if (this.crossoverRate > Math.random() && populationIndex >=
                this.elitismCount) {

             Individual offspring = new Individual(parent1.getChromosomeLength(       ));

                Individual parent2 = selectParent(population);
                for (int geneIndex = 0; geneIndex < parent1.getChromosomeLength();
geneIndex++) {
                    if (0.5 > Math.random()) {
                        offspring.setGene(geneIndex, parent1.getGene(geneIndex));
                    } else {
                        offspring.setGene(geneIndex, parent2.getGene(geneIndex));
                    }
                }
```

```
                newPopulation.setIndividual(populationIndex, offspring);
            } else {
                newPopulation.setIndividual(populationIndex, parent1);
            }
        }

        return newPopulation;
    }

    public Population mutatePopulation(Population population) {
        Population newPopulation = new Population(this.populationSize);
        for (int populationIndex = 0; populationIndex < population.size(); populationIndex++) {
            Individual individual = population.getFittest(populationIndex);
            for (int geneIndex = 0; geneIndex < individual.getChromosomeLength(); geneIndex++)
{
                // Skip mutation if this is an elite individual
                if (populationIndex > this.elitismCount) {
                    // Does this gene need mutation?
                    if (this.mutationRate > Math.random()) {
                        // Get new gene
                        int newGene = 1;
                        if (individual.getGene(geneIndex) == 1) {
                            newGene = 0;
                        }
                        // Mutate gene
                        individual.setGene(geneIndex, newGene);
                    }
                }
            }
            newPopulation.setIndividual(populationIndex, individual);
        }

        return newPopulation;
    }

}
```

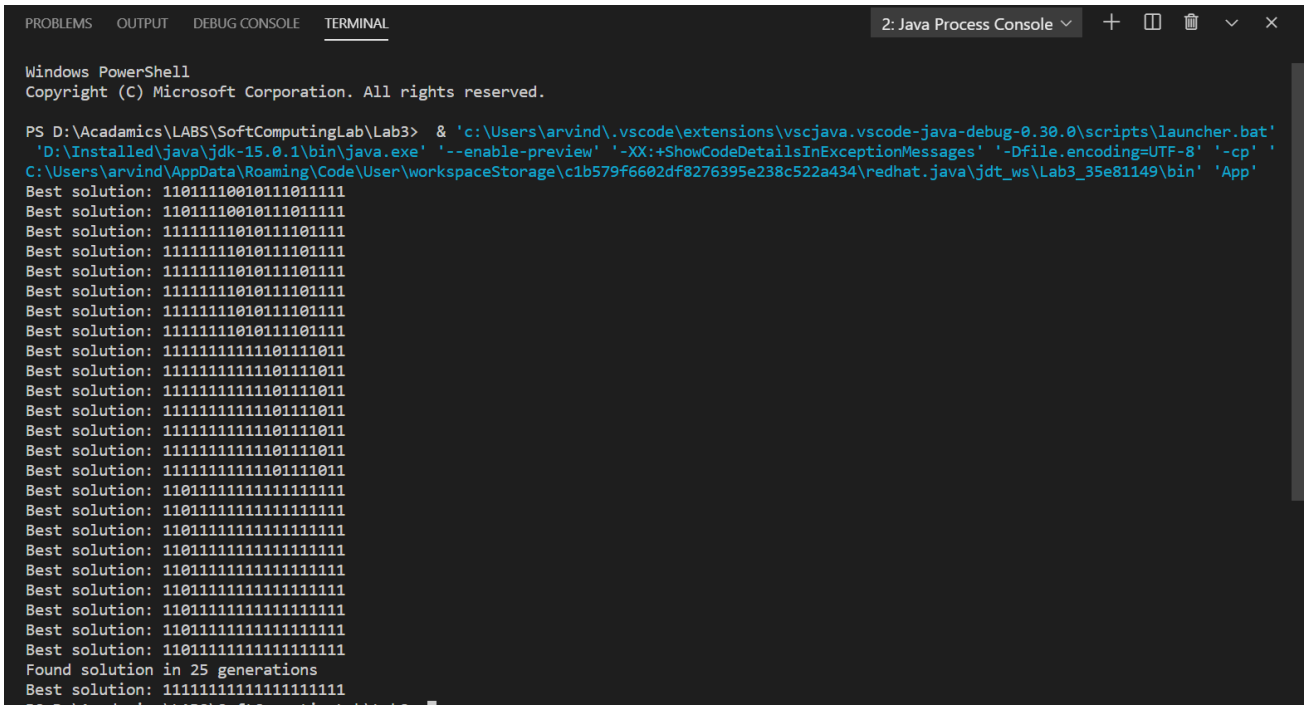
App.java :

```
//import Lab3.Individual;
import Lab3.Population;
import Lab3.GeneticAlgorithm;

public class App {
    private static final int numberOfBits = 20;

    public static void main(String[] args) {
    GeneticAlgorithm ga = new GeneticAlgorithm(100, 0.001, 0.95, 2);
    Population population = ga.initPopulation(numberOfBits);
    ga.evalPopulation(population);
    int generation = 1;
    while (ga.isTerminationConditionMet(population) == false) {
        System.out.println("Best solution: " + population.getFittest(0).toString());
        population = ga.crossoverPopulation(population);
        population = ga.mutatePopulation(population);
        ga.evalPopulation(population);
        generation++;
    }
    System.out.println("Found solution in " + generation + " generations");
    System.out.println("Best solution: " + population.getFittest(0).toString());
  }
}
```

Output :


PROBLEMS
OUTPUT
DEBUG CONSOLE
TERMINAL
2: Java Process Console
Windows PowerShell
Copyright (C) Microsoft Corporation. All rights reserved.
PS D:\Acadamics\LABS\SoftComputingLab\Lab3> & 'c:\Users\arvind\.vscode\extensions\vscjava.vscode-java-debug-0.30.0\scripts\launcher.bat' 'D:\Installed\java\jdk-15.0.1\bin\java.exe' '--enable-preview' '-XX:+ShowCodeDetailsInExceptionMessages' '-Dfile.encoding=UTF-8' '-cp' 'C:\Users\arvind\AppData\Roaming\Code\User\workspaceStorage\c1b579f6602df8276395e238c522a434\redhat.java\jdt_ws\Lab3_35e81149\bin' 'App'
Best solution: 1101111001011011111
Best solution: 1101111001011011111
Best solution: 1111111101011101111
Best solution: 1111111101011101111
Best solution: 1111111101011101111
Best solution: 1111111101011101111
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Best solution: 1101111111111111111
Best solution: 1101111111111111111
Found solution in 25 generations
Best solution: 1111111111111111111