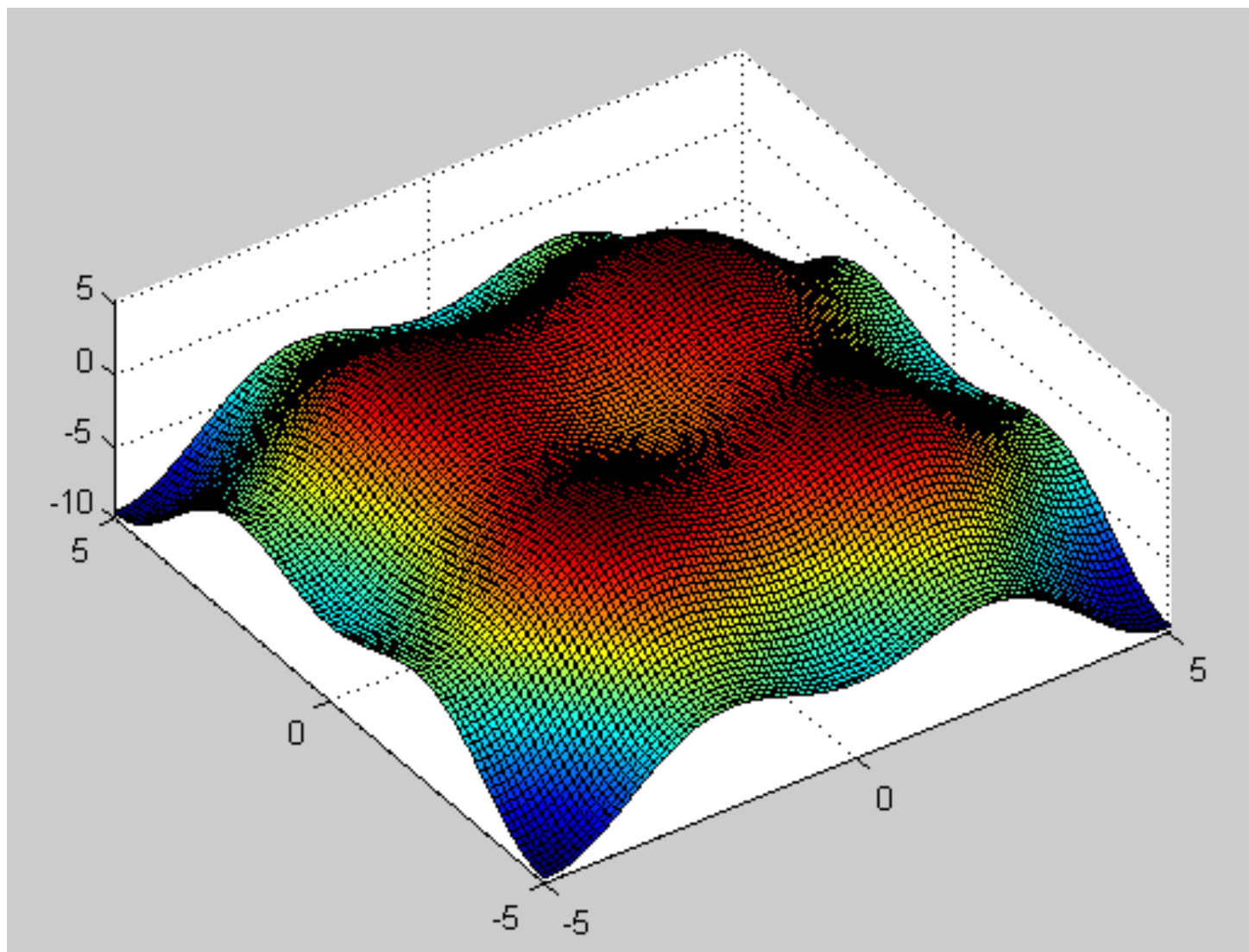


Practice 3

Maximization of the 3D function
with multiple solutions.

We must search for all of the 4 maximums of the
function $z = x \sin(|x|) + y \sin(|y|)$

With $x = [-5; 5]$ and $y = [-5; 5]$



To find all 4 solutions I'll try to use

1. Chromosomes with 22 binary genes
First 11 for X
And last 11 for Y
2. Population with 20 chromosomes
3. I'll try to calculate shared fitness function with
Manhattan distance d_{ij}
4. Trunked selection $\frac{1}{2}$

5. Without mutations

6. Repeat until maximum shared fitness of last 10 population are equals

Code

Chromosom.java

```
import java.util.ArrayList;
import java.util.List;
import java.util.Random;

public class Chromosom {
    private List<Boolean> genes = new ArrayList<Boolean>();

    private static Random random = new Random();

    private Double sharedFitness;

    public Double getSharedFitness() {
        return sharedFitness;
    }

    public void setSharedFitness(Double sharedFitness) {
        this.sharedFitness = sharedFitness;
    }

    public Chromosom(){
        for(int i = 0; i < 22; i++){
            genes.add(random.nextBoolean());
        }
    }

    public Chromosom(List<Boolean> genes){
        this.genes = genes;
    }

    public static List<Chromosom> crossover(Chromosom papa, Chromosom mama){
        List<Chromosom> children = new ArrayList<Chromosom>();

        List<Boolean> firstChildGenes = new ArrayList<>();
        List<Boolean> secondChildGenes = new ArrayList<>();

        for(int i = 0; i < 22; i++){
            if(random.nextBoolean()){
                firstChildGenes.add(papa.getGenes().get(i));
                secondChildGenes.add(mama.getGenes().get(i));
            }else{
                firstChildGenes.add(mama.getGenes().get(i));
                secondChildGenes.add(papa.getGenes().get(i));
            }
        }

        children.add(new Chromosom(firstChildGenes));
        children.add(new Chromosom(secondChildGenes));

        return children;
    }
}
```

```

public List<Boolean> getGenes() {
    return genes;
}

public void setGenes(List<Boolean> genes) {
    this.genes = genes;
}

public double getFitness(){
    double x = 0;
    double y = 0;

    for(int i = 1; i < 11; i++){
        x += genes.get(i) ? Math.pow(2.0, i - 1)/1023.0*5.0 : 0;
        y += genes.get(i + 11) ? Math.pow(2.0, i - 1)/1023.0*5.0 : 0;
    }

    x *= genes.get(0) ? -1.0 : 1.0;
    y *= genes.get(11) ? -1.0 : 1.0;

    return x * Math.sin(Math.abs(x)) + y * Math.sin(Math.abs(y));
}

public double getX(){
    double x = 0;

    for(int i = 1; i < 11; i++){
        x += genes.get(i) ? Math.pow(2.0, i - 1)/1023.0*5.0 : 0;
    }

    x *= genes.get(0) ? -1.0 : 1.0;

    return x;
}

public double getY(){
    double y = 0;

    for(int i = 12; i < 22; i++){
        y += genes.get(i) ? Math.pow(2.0, i - 1)/1023.0*5.0 : 0;
    }

    y *= genes.get(0) ? -1.0 : 1.0;

    return y;
}
}

```

Population.java

```

import java.util.ArrayList;
import java.util.Collections;
import java.util.List;
import java.util.Random;

public class Population {

    private List<Chromosom> chromosoms = new ArrayList<Chromosom>();

    private double sigma = 9.0;

    private static Random random = new Random();

```

```

public Population() {
    for (int i = 0; i < 20; i++) {
        chromosomes.add(new Chromosom());
    }
}

public Population(List<Chromosom> chromosomes) {
    this.chromosomes = chromosomes;
}

public List<Chromosom> getChromosomes() {
    return chromosomes;
}

public void setChromosomes(List<Chromosom> chromosomes) {
    this.chromosomes = chromosomes;
}

public Population nextGeneration() {
    List<Chromosom> nextGen = new ArrayList<>();

    Collections.sort(chromosomes, (c1, c2) ->
c1.getSharedFitness().compareTo(c2.getSharedFitness()));

    for (int i = 0; i < 10; i++) {
        nextGen.addAll(Chromosom.crossover(chromosomes.get(random.nextInt(10)),
chromosomes.get(random.nextInt(10))));
    }

    return new Population(nextGen);
}

public double getSharedFitness(int index) {
    double result = 0;

    for (int i = 0; i < 20; i++) {
        double x = Math.abs(getChromosomes().get(index).getX() -
getChromosomes().get(i).getX());
        double y = Math.abs(getChromosomes().get(index).getY() -
getChromosomes().get(i).getY());

        double d = x + y;

        result += s(d);
    }

    return getChromosomes().get(index).getFitness() / result;
}

private double s(double d) {
    return d < sigma ? 1 - (d / sigma) : 0;
}

public void calcSharedFitnessList() {
    for (int i = 0; i < 20; i++) {
        getChromosomes().get(i).setSharedFitness(getSharedFitness(i));
    }
}

public double getMax() {
    List<Double> fitnessList = new ArrayList<>();

    for (int i = 0; i < 20; i++) {

```

```

        fitnessList.add(chromosoms.get(i).getSharedFitness());
    }

    return Collections.max(fitnessList);
}

public double getAverage() {
    double sum = 0;
    for (int i = 0; i < 20; i++) {
        sum += chromosoms.get(i).getSharedFitness();
    }

    return sum/20.0;
}
}

```

plotGraph.m (Matlab code for graphic plot)

```

x = (-5 : 0.1 : 5);
y = (-5 : 0.1 : 5);
z = zeros(101, 101);

for i = 1:101
    for j = 1 : 101
        z(i, j) = x(i) * sin(x(i)) + y(j) * sin(y(j));
    end
end

surf(x, y, z);

```