

Source code:

```
import java.util.*;

class Attempts
{
    public int value;
    public boolean changed;
    public Attempts()
    {
        value = 0;
        changed = false;
    }
}

class Chromosome
{
    private static int numberLearningAttempts = 1000;

    private final boolean fixGenomeWhenSuccess = false;
    private final double fitnessBoost = 19.;

    private int[] genome;
    private int ID;

    public static void setNumberOfLearningAttempts(int numberLearningAttempts)
    {
        Chromosome.numberLearningAttempts = numberLearningAttempts;
    }

    public Chromosome(int ID, int[] genome)
    {
        this.ID = ID;
        this.genome = genome;
    }

    public Chromosome(int ID, int size, int fixedElements, Random random)
    {
        this.ID = ID;
        genome = new int[size];
        ArrayList<Integer> positions = new ArrayList<>();
        for(int i = 0; i < size; i++)
        {
            genome[i] = 2; // init as ?
            positions.add(i);
        }
        Collections.shuffle(positions, random);
        for(int i = 0; i < fixedElements; i++)
        {
            int pos = positions.get(i);
            genome[pos] = random.nextInt(2);
        }
    }

    @Override
    public String toString()
    {
        if(genome == null)
            return "";
        return String.valueOf(ID) + '\t' + sequenceToString(genome);
    }

    public int[] getSequence()
```

```
{
    return this.genome;
}
```

```
public static String sequenceToString(int[] genome)
{
    if(genome == null)
        return "";
    StringBuilder s = new StringBuilder();
    for(int aGenome : genome)
        if(aGenome == 2)
            s.append('?');
        else
            s.append(aGenome);
    return s.toString();
}
```

```
public static Chromosome mate(int ID, Chromosome ind1, Chromosome ind2, Random rand)
{
    int recPos = 1 + rand.nextInt(ind1.genome.length - 1);
    int[] genomeNew = new int[ind1.genome.length];
    System.arraycopy(ind1.genome, 0, genomeNew, 0, recPos);
    System.arraycopy(ind2.genome, recPos, genomeNew, recPos, genomeNew.length - recPos);
    return new Chromosome(ID, genomeNew);
}
```

```
public static int[] getCounts(int[] genome)
{
    int[] cnts = new int[3];
    for(int aGenome : genome)
        cnts[aGenome]++;
    return cnts;
}
```

```
public double liveLearnAndReturnFitness(Random random, Attempts att)
{
    int len = genome.length;
    if(getCounts(genome)[1] == len)
        // this individual has correct sequence, no learning will happen
        return 1 + fitnessBoost;
    else if(getCounts(genome)[0] > 0)
        // this individual has 0s in the sequence, will not learn right solution
        return 1;
    else
    {
        // do learning
        int[] sequence = genome.clone();
        ArrayList<Integer> positionOfQuestionMarks = new ArrayList<>();
        for(int i = 0; i < sequence.length; i++)
            if(sequence[i] == 2)
                positionOfQuestionMarks.add(i);
        int attempts = 1;
        for(; attempts <= numberLearningAttempts; attempts++)
        {
            int[] learnedBits = getRandomBinaryString(positionOfQuestionMarks.size(), random);
            for(int i = 0; i < positionOfQuestionMarks.size(); i++)
                sequence[positionOfQuestionMarks.get(i)] = learnedBits[i];
            if(getCounts(sequence)[1] == len)
            {
                if(fixGenomeWhenSuccess)
                    genome = sequence;
                att.value = attempts;
            }
        }
    }
}
```

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        att.changed = true;
        return 1 + fitnessBoost * (numberLearningAttempts - attempts) / 1000.;
    }
}
// could not learn
return 1;
}
}

public static int[] getRandomBinaryString(int len, Random random)
{
    int[] s = new int[len];
    for(int i = 0; i < len; i++)
        s[i] = random.nextBoolean() ? 1 : 0;
    return s;
}

public static Chromosome sampleIndividualPropToFitness(Random random, HashMap<Chromosome, Double>
largeFitness,
                ArrayList<Chromosome> fitnessOne,
                double totFitnessGreaterThanOrEqualTo)
{
    double totFitness = fitnessOne.size() + totFitnessGreaterThanOrEqualTo;
    if(random.nextDouble() < fitnessOne.size() / totFitness)
        // sample from those with fitness = 1
        return fitnessOne.get(random.nextInt(fitnessOne.size()));
    else
    {
        // sample from those with fitness > 1
        double fitnessMark = random.nextDouble() * totFitnessGreaterThanOrEqualTo;
        double cumulativeFitness = 0.;
        Chromosome sampledInd = null;
        for(Chromosome ind : largeFitness.keySet())
        {
            cumulativeFitness += largeFitness.get(ind);
            if(cumulativeFitness >= fitnessMark)
            {
                sampledInd = ind;
                break;
            }
        }
        return sampledInd;
    }
}

public static boolean isFixated(ArrayList<Chromosome> inds)
{
    int[] firstGenome = inds.get(0).genome;
    for(Chromosome ind : inds)
        if(!Arrays.equals(firstGenome, ind.genome))
            return false;
    return true;
}

public static double[] getGenomeProb(Collection<Chromosome> inds)
{
    double[] res = new double[3];
    for(Chromosome ind : inds)
    {
        int[] thisInd = getCounts(ind.genome);
        res[0] += thisInd[0];
        res[1] += thisInd[1];
        res[2] += thisInd[2];
    }
}

```

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    }
    double sum = res[0] + res[1] + res[2];
    res[0] /= sum;
    res[1] /= sum;
    res[2] /= sum;
    return res;
}

}

public class Main
{
    static long seed = System.currentTimeMillis();

    public static void main(String[] args) throws InterruptedException
    {
        Random random = new Random(seed);
        int numberOfSamples = 1000;
        int len = 20;
        int fixed = 10;

        Chromosome.setNumberOfLearningAttempts(numberOfSamples);

        ArrayList<Chromosome> inds = new ArrayList<>();
        for(int i = 0; i < numberOfSamples; i++)
            inds.add(new Chromosome(i, len, fixed, random));

        int generation = 0;
        double[] populationStats = Chromosome.getGenomeProb(inds);

        while(!Chromosome.isFixated(inds))
        {
            HashMap<Chromosome, Double> largeFitness = new HashMap<>();
            ArrayList<Chromosome> fitnessOne = new ArrayList<>();
            double totFitnessGreaterThanOne = 0., totalAttempts = 0, bestAttempts = 1000;
            int cnt = 0;
            for(Chromosome ind : inds)
            {
                Attempts attempts = new Attempts();
                double fitness = ind.liveLearnAndReturnFitness(random, attempts);
                totalAttempts += attempts.value;
                if(attempts.changed)
                    cnt++;
                if(attempts.changed && attempts.value < bestAttempts)
                    bestAttempts = attempts.value;
                if(fitness > 1)
                {
                    largeFitness.put(ind, fitness);
                    totFitnessGreaterThanOne += fitness;
                } else
                    fitnessOne.add(ind);
            }
            System.out.printf(
                "GENERATION %4d\t\t0 --> %.6f\t\t\t1 --> %.6f\t\t\t? --> %.6f\t\t\taverage --> %3.6f\t\t\tbest --> %3.6f\n",
                generation, populationStats[0], populationStats[1], populationStats[2], totalAttempts / cnt,
                bestAttempts);

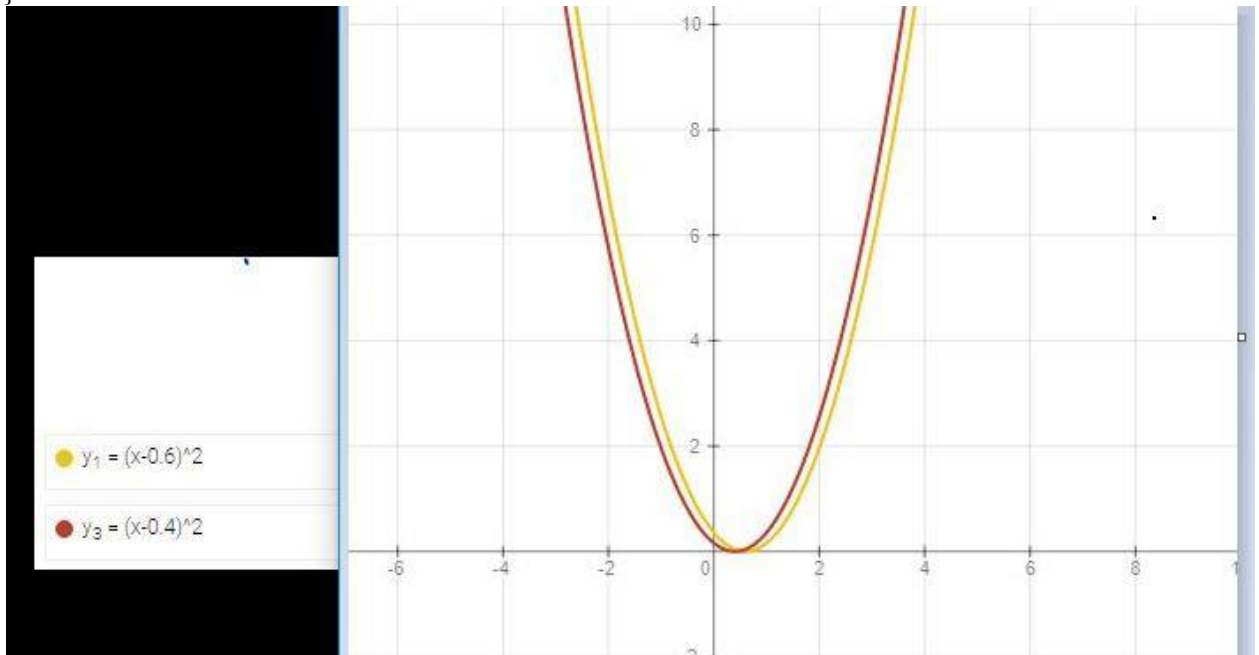
            ArrayList<Chromosome> newGen = new ArrayList<>();
            for(int i = 0; i < numberOfSamples; i++)
            {
                Chromosome samp1 = Chromosome.sampleIndividualPropToFitness(random, largeFitness, fitnessOne,
                    totFitnessGreaterThanOne);
            }
        }
    }
}

```

```

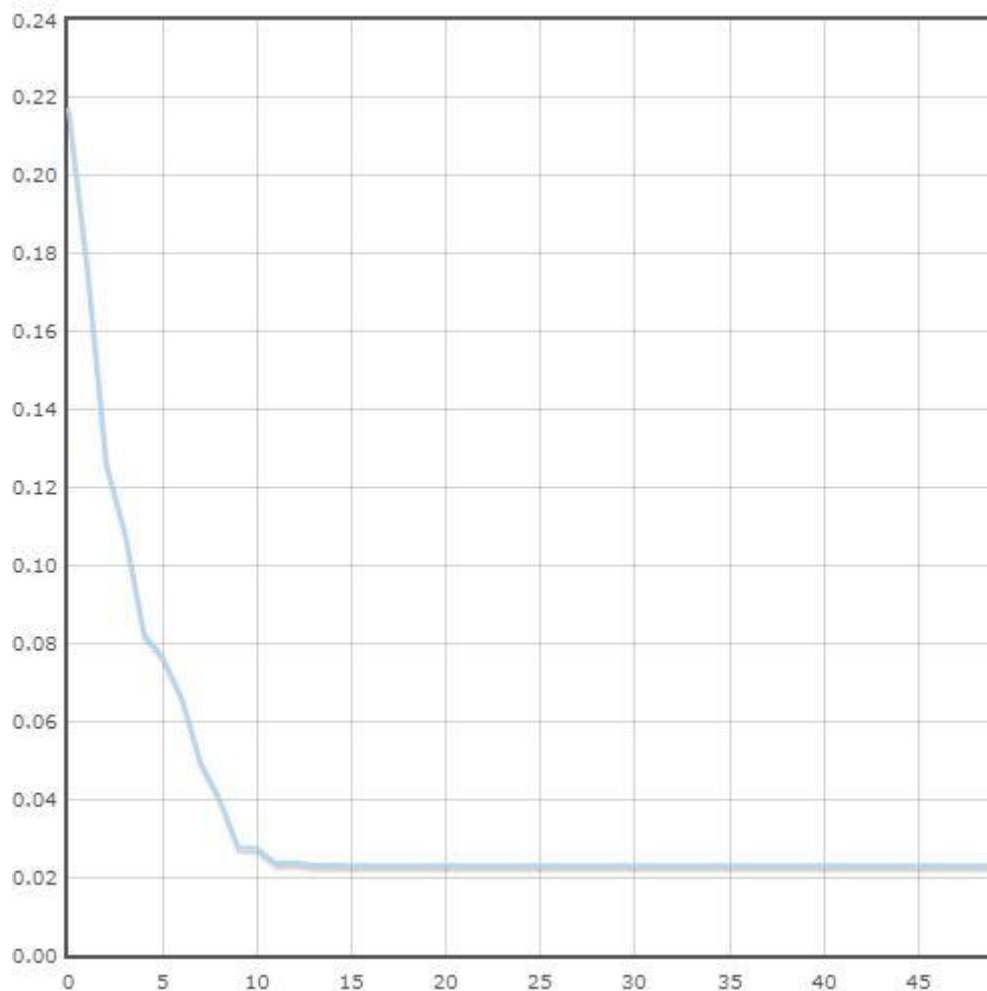
        Chromosome samp2 = Chromosome.sampleIndividualPropToFitness(random, largeFitness, fitnessOne,
                                                                    totFitnessGreaterThanOrEqualToOne);
        Chromosome newSamp = Chromosome.mate(i, samp1, samp2, random);
        newGen.add(newSamp);
    }
    inds = newGen;
    populationStats = Chromosome.getGenomeProb(inds);
    generation++;
}
System.out.println("Done:\t" + Chromosome.sequenceToString(inds.get(0).getSequence()) + "\t" +
populationStats[0] + "\t" + populationStats[1] + "\t" + populationStats[2]);
Thread.sleep(1000);
}
}

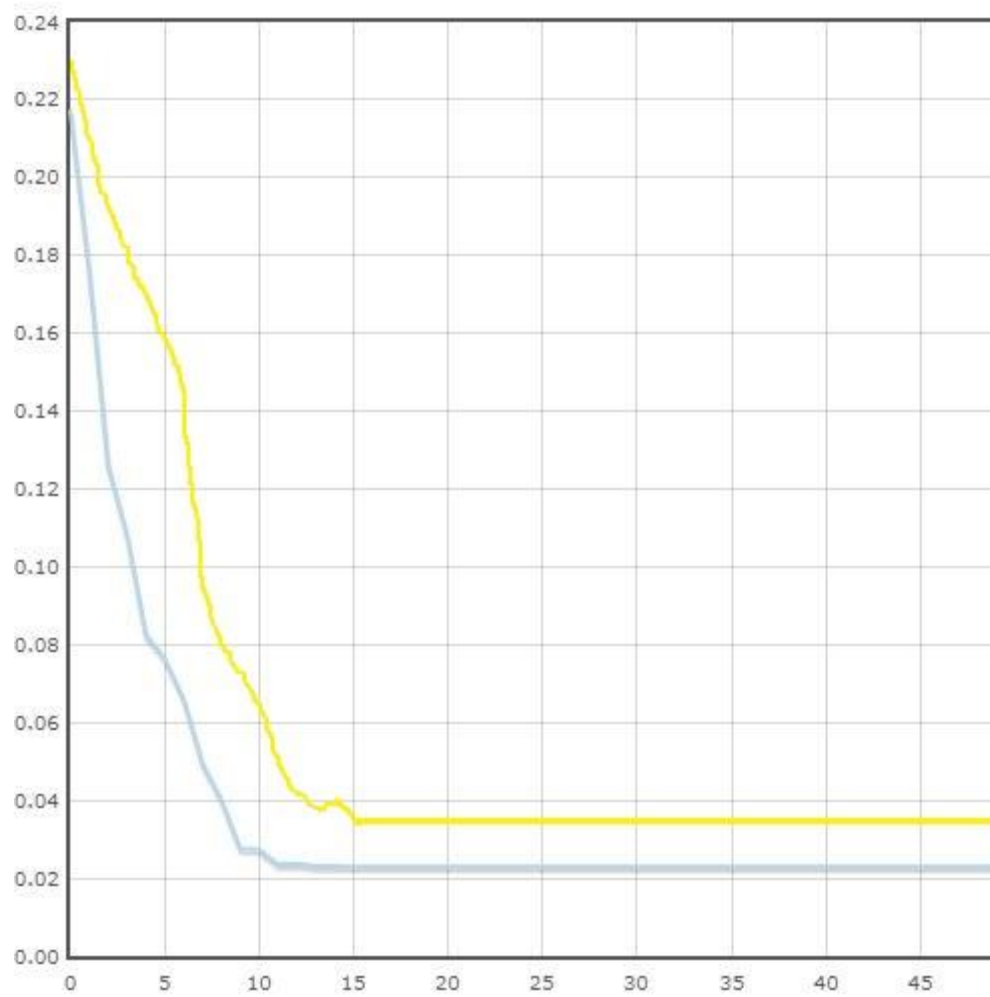
```



	A	B	C	D	E	F	G
1	0:0.21747751					0:0.51747932	
2	1:0.17643043					1:0.47643043	
3	2:0.12644100					2:0.42644254	
4	3:0.10857654					3:0.40857936	
5	4:0.08274021					4:0.38274184	
6	5:0.07661301					5:0.37661706	
7	6:0.0663210					6:0.3665887	
8	7:0.0496021					7:0.34961778	
9	8:0.04036321					8:0.34037143	
10	9:0.027723256					9:0.327727623	
11	10:0.027446589					10:0.327442953	
12	11:0.02361369					11:0.32363541	
13	12:0.02371857					12:0.32371857	
14	13:0.0231231					13:0.3232465	
15	14:0.023100145					14:0.323100017	
16	15:0.023047152					15:0.323059485	
17	16:0.023047152					16:0.323051761	
18	17:0.023047152					17:0.323051761	

СОХРАНЕНО В ONEDRIVE





Blue line- $y = (x - 0.6)^2$

Yellow line- $y = (x - 0.4)^2$