

CSC-20043 Computational and Artificial Intelligence I

Evolutionary Algorithms Practical 2

Steady-State GA for the Travelling Salesman Problem

The code outline below is available as file `adcTSP-outline.java`. The missing parts are for you to complete, to check your understanding. (This is not assessed, and you do not need to submit anything.)

Instructions:

- A) If you do not know what the Travelling Salesman Problem is then look it up
- B) Read through the code outline and understand what each part should do
- C) Write code to replace the `CCCC`, to complete the code that calculates fitnesses
- D) Write code to replace the `DDDDs`, to complete the code that selects parents for reproduction
- E) Write code to replace the `EEEEs`, to produce the child genotype using mutation and crossover.
Your mutation function should swap two towns around in a route. Your crossover function should copy the towns before a cut point from parent 1 and take the remaining towns in the order they appear in parent 2.
- F) Run the completed program several times. Has it evolved routes of length approx. 424?

```
import java.util.Random;
class adcTSP { // Travelling Salesman Problem

    final int numTowns          = 30;
    final int numGenotypes       = 1000;
    final int genotypeLength     = numTowns;
    final int numReproductions   = 1000000;
    final int numMutationsPerReproduction = 1;

    final int towns[/*numTowns*/][/*2*/]=
    { {82, 7} , {91,38} , {62,32} , {71,44} , {83,69} , {68,58} ,
      {54,67} , {87,76} , {13,40} , {71,71} , {44,35} , {18,54} ,
      {64,60} , {37,84} , {41,94} , { 2,99} , { 7,64} , {22,60} ,
      {25,62} , {54,62} , { 4,50} , {74,78} , {18,40} , {24,42} ,
      {25,38} , {41,26} , {45,21} , {58,69} , {58,35} , {83,46} };
    double distances[][] = new double[numTowns][numTowns];

    int genotypes[][] = new int[numGenotypes][genotypeLength];
    double fitness[] = new double[numGenotypes]; // fitness = -1 * route length
    int fittestIndividual = 0;

    Random rng = new Random();
    int random(int n) { return rng.nextInt(n); } // random integer between 0 and n-1

    void calculateFitness(int individual) // fitness = -1 * route length
    {
        CCCC
    }

    void initialise()
    { /* calculate distances between towns */
        for (int i=0;i<numTowns;i++) for (int j=0;j<=i;j++)
        { int dx=towns[j][0]-towns[i][0], dy=towns[j][1]-towns[i][1];
          distances[i][j] = distances[j][i] = Math.sqrt((double)(dx*dx+dy*dy));
        }

        /* generate initial (random) population */
        for (int individual=0;individual<numGenotypes;individual++)
        { int remainingTowns[] = new int[numTowns];
          for (int i=0;i<numTowns;i++) remainingTowns[i] = i;
          for (int g=0;g<genotypeLength;g++)
          { int index = random(numTowns-g);
            genotypes[individual][g] = remainingTowns[index];
            remainingTowns[index] = remainingTowns[numTowns-g-1];
          }
        }
        /* calculate fitness array for initial population */
        for (int individual=0;individual<numGenotypes;individual++)
            calculateFitness(individual);
    }
}
```

```

void mutate(int individual) // swap two towns in individual's route
{
    EEEE
}

void crossover(int parentA, int parentB, int child)
{
    EEEE
}

void steadyStateGaMainStep()
{ /* pick three individuals a,b,c at random */
    DDDD

    /* reorder such that c is the least fit */
    DDDD

    /* but don't loose fittestIndividual */
    if (c==fittestIndividual) {int temp=b;b=c;c=temp;}

    /* crossover a,b (in random order) to create child that replaces c */
    if (random(2)==1) {int temp=a;a=b;b=temp;}
    crossover(a,b,c);
    /* mutate child */
    for (int m=0;m<numMutationsPerReproduction;m++) mutate(c);
    /* calculate fitness of child */
    calculateFitness(c);
}

void outputStatistics(int numRreproductionsSoFar)
{ if (numRreproductionsSoFar==0) System.out.println("#repros\tbest\t: route");
  System.out.print(""+numRreproductionsSoFar+"\t"+(-fitness[fittestIndividual])+"\t:");
  for (int g=0;g<genotypeLength;g++)
      System.out.print(" "+(genotypes[fittestIndividual][g]+1));
  System.out.println();
}

void run()
{ initialise();
  outputStatistics(0);

  for (int reproduction=1;reproduction<=numReproductions;reproduction++)
  { steadyStateGaMainStep();
    if (reproduction%20000==0) outputStatistics(reproduction);
  }
}

public static void main(String args[])
{ new adcTSP().run();
}

}

```