



# Advanced Algorithms

## Assignment 2: Traveling Salesman Problem

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# 1 Introduction

For this assignment, we implemented and analyzed the running times and the errors of three algorithms for the Traveling Salesman Problem. The algorithms implemented are:

1. Nearest Neighbor Algorithm
2. Random Insertion Algorithm
3. 2-approximate Algorithm

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## 2 Nearest Neighbor Algorithm

For the first constructive heuristic algorithm we have chosen Nearest Neighbor. The goal of the algorithm is to start from a single-node path, and then add the neighbor with the minimum weight not in the existing path. The steps are the following:

1. **Initialization:** start from the single-node path 0;
2. **Selection:** let  $(v_0, \dots, v_k)$  be the current path. Find the vertex  $v_{k+1}$  not in the path with minimum distance from  $v_k$ ;
3. **Insertion:** insert  $v_{k+1}$  immediately after  $v_k$ ;
4. repeat from (2) until all vertices are inserted in the path.

### 2.1 Functions

The implementation doesn't require particular data structure so we created two functions for the algorithm:

- **NearestNeighbor(g,v):** method that builds the Hamiltonian circuit and returns its weight and the execution time;
- **minDist(p,g,l):** method that finds the neighbor with the minimum weight between the last vertex insert in the path p and its neighbors that are in some indexes of the list g.

### 2.2 Implementation

The final solution is the following:

- We start the time;
- We execute a deep copy unvisited of the original list v;
- We follow the algorithm and we start from the node 0 (*startingNode*) of the unvisited list.
- We remove the starting node from the unvisited list and we add always the starting node in our path (*is a list*);
- We initialize our solution (*dist*) to zero and we start our cycle until we have visited all vertices (*len(unvisited)>0*);
- In the while cycle:
  1. We find the neighbor with the minimum distance with the function minDist:
    - It initialize the distance to infinite, the vertex to none and it take the last node of the path (*p*);
    - In the cycle it checks all the neighbors of the node and returns the distance and the vertex with the minimum distance that is not just in p.

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2. We remove the vertex found from the list unvisited and we add the vertex in the path;
  3. We add the distance of the vertex found and we continue with the next iteration until we removed all the vertex from unvisited.
- When we finish the while cycle we add to dist the distance between the last node in the path and the startingNode and to close the Hamiltonian cycle we add the vertex to the path;
  - We end the time and we calculate the time cost;
  - We return the solution and the time.

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## 3 Random Insertion Algorithm

### 3.1 Input and Data Structures

### 3.2 Implementation

### 3.3 Complexity

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## 4 2-approximate Algorithm

The 2-approximate algorithm starts by defining a starting node from the list of vertices within a graph, and then a minimum spanning tree is constructed using Prim's algorithm with the starting node as the root. The vertices of the MST are then visited in a preorder walk/depth first search, and added to a list in the order of which they are visited. Finally, the starting node is added at the end of the preorder list in order to complete the Hamiltonian cycle in the MST.

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```
1 APPROX_METRIC_TSP (G)
2   V = { v1, v2, ..., vn }
3   r ← v1
4   T* ← PRIM (G, r)
5   < vi1, vi2, ..., vin > = H' ← PREORDER (T*, r)
6   return < H', vi1 > = H
```

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### 4.1 Input and Data Structure

The 2-approximate algorithm requires a minimum spanning tree to be constructed using Prim's algorithm, and in order to be efficient, Prim's algorithm requires a heap data structure. We reused Prim's algorithm and the min heap data structure that were implemented in Assignment 1 for this course; however, the input files for Assignment 1 provided a list of connected nodes, whereas the input files for Assignment 2 only provide a list of distinct vertices and their x and y coordinates. Consequently, we made the following modifications to the Graph and Node classes, and added two new functions:

- **Node:** three new instance variables are initialized:
  - xcoord: x-coordinate of vertex
  - ycoord: y-coordinate of vertex
  - weighttype: weight type of the vertex, either Euclidean or Geographic
- **Graph:** the buildGraph function restructures the input file to resemble the input file format of Assignment 1, before calling the createNodes and makeNodes functions. Specifically, the buildGraph function takes each vertex and connects it to all other vertices in the input file.
- **Weight:** new function that takes the weight type, x-coordinate, and y-coordinate for two vertices as input, and returns the weight between the two vertices. For the Euclidean weight type, there is no coordinate conversion and we simply calculate the Euclidean distance rounded to the nearest integer. For the Geographic weight type, the x- and y-coordinates are converted to radians and then the geographic distance is calculated using the instructions in the FAQ site provided by the assignment.
- **Preorder:** new function that takes a MST graph, starting node, and empty set as input. The vertices of the MST are visited in a preorder walk/depth first search and

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added to the list in that order. The list is returned once all vertices of the MST have been visited and added to the list.

## 4.2 Implementation

The solution to the cost of the Hamiltonian cycle in the MST is performed in the following steps:

1. Create the Graph object through the Graph() class and call the buildGraph method
2. Calculate the MST of the graph using Prim's algorithm
3. Call the Preorder function, passing the inputs of the MST and the same starting node used as the root in Prim's algorithm.
4. Append the starting node to the end of the list returned by the Preorder function in order to complete the Hamiltonian cycle in the MST
5. Call the Weight function to calculate the distance between each node in the Hamiltonian cycle, sum all of the edge weights to calculate the total cost of the cycle



## 5 Results

### 5.1 Table of results

| Instance                  | Nearest Neighbor |          |           | Random Insertion |          |           | 2-Approximation |          |           |
|---------------------------|------------------|----------|-----------|------------------|----------|-----------|-----------------|----------|-----------|
|                           | Solution         | Time (s) | Error (%) | Solution         | Time (s) | Error (%) | Solution        | Time (s) | Error (%) |
| tsp_dataset/burma14.tsp   | 4048             | 0.00005  | 21.82%    | 4035             | 0.00033  | 21.43%    | 4003            | 0.00011  | 20.46%    |
| tsp_dataset/ulysses16.tsp | 9988             | 0.00006  | 45.62%    | 8427             | 0.00044  | 22.86%    | 7788            | 0.00013  | 13.54%    |
| tsp_dataset/ulysses22.tsp | 10586            | 0.00012  | 50.95%    | 9396             | 0.00110  | 33.98%    | 8308            | 0.00022  | 18.47%    |
| tsp_dataset/eil51.tsp     | 511              | 0.00044  | 19.95%    | 734              | 0.01191  | 72.30%    | 605             | 0.00076  | 42.02%    |
| tsp_dataset/berlin52.tsp  | 8980             | 0.00045  | 19.07%    | 13593            | 0.01071  | 80.23%    | 10402           | 0.00080  | 37.92%    |
| tsp_dataset/kroA100.tsp   | 27807            | 0.00125  | 30.66%    | 45040            | 0.06388  | 111.63%   | 30516           | 0.00251  | 43.39%    |
| tsp_dataset/kroD100.tsp   | 26947            | 0.00141  | 26.55%    | 45199            | 0.06452  | 112.26%   | 28599           | 0.00257  | 34.31%    |
| tsp_dataset/ch150.tsp     | 8191             | 0.00272  | 25.47%    | 12567            | 0.21607  | 92.51%    | 9126            | 0.00595  | 39.80%    |
| tsp_dataset/gr202.tsp     | 49336            | 0.00613  | 22.85%    | 75119            | 0.52650  | 87.05%    | 52615           | 0.01018  | 31.01%    |
| tsp_dataset/gr229.tsp     | 162430           | 0.01025  | 20.67%    | 264776           | 0.76313  | 96.71%    | 179335          | 0.01320  | 33.23%    |
| tsp_dataset/pcb442.tsp    | 61979            | 0.04577  | 22.06%    | 108418           | 5.63636  | 113.51%   | 72853           | 0.04758  | 43.47%    |
| tsp_dataset/d493.tsp      | 41660            | 0.05979  | 19.02%    | 66608            | 7.83201  | 90.30%    | 45595           | 0.06093  | 30.26%    |
| tsp_dataset/dsj1000.tsp   | 24630960         | 0.15553  | 32.00%    | 39530984         | 65.98835 | 111.85%   | 25526005        | 0.32737  | 36.80%    |

Figure 1: Table of results

### 5.2 Graph of the execution time comparison

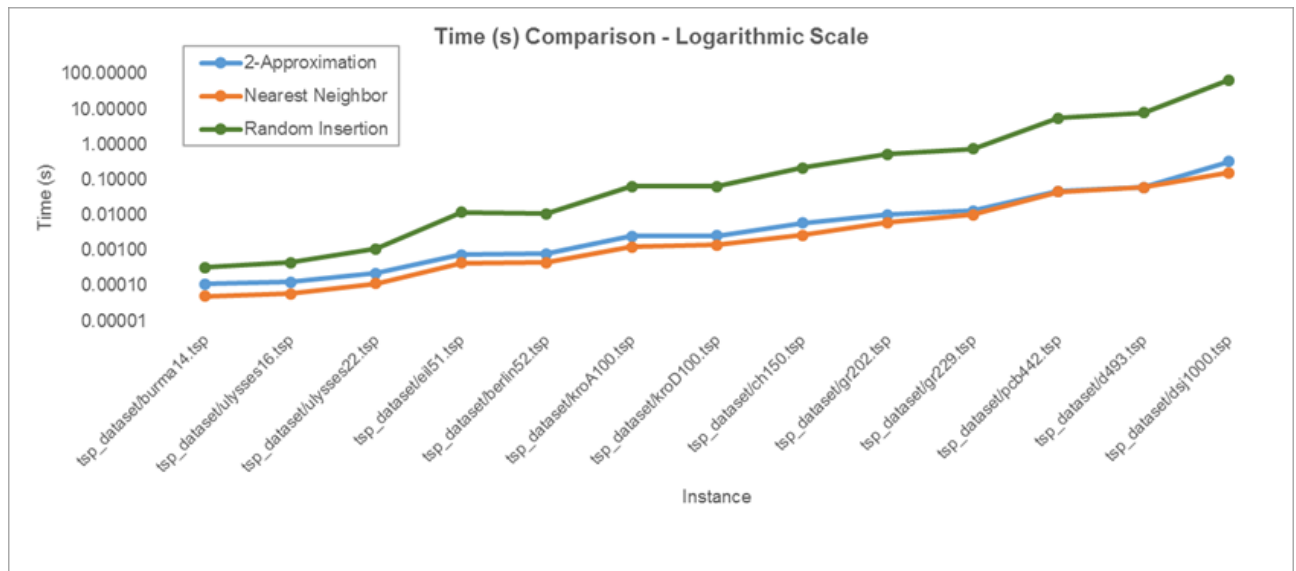


Figure 2: Execution time comparison

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### 5.3 Graph of the error comparison

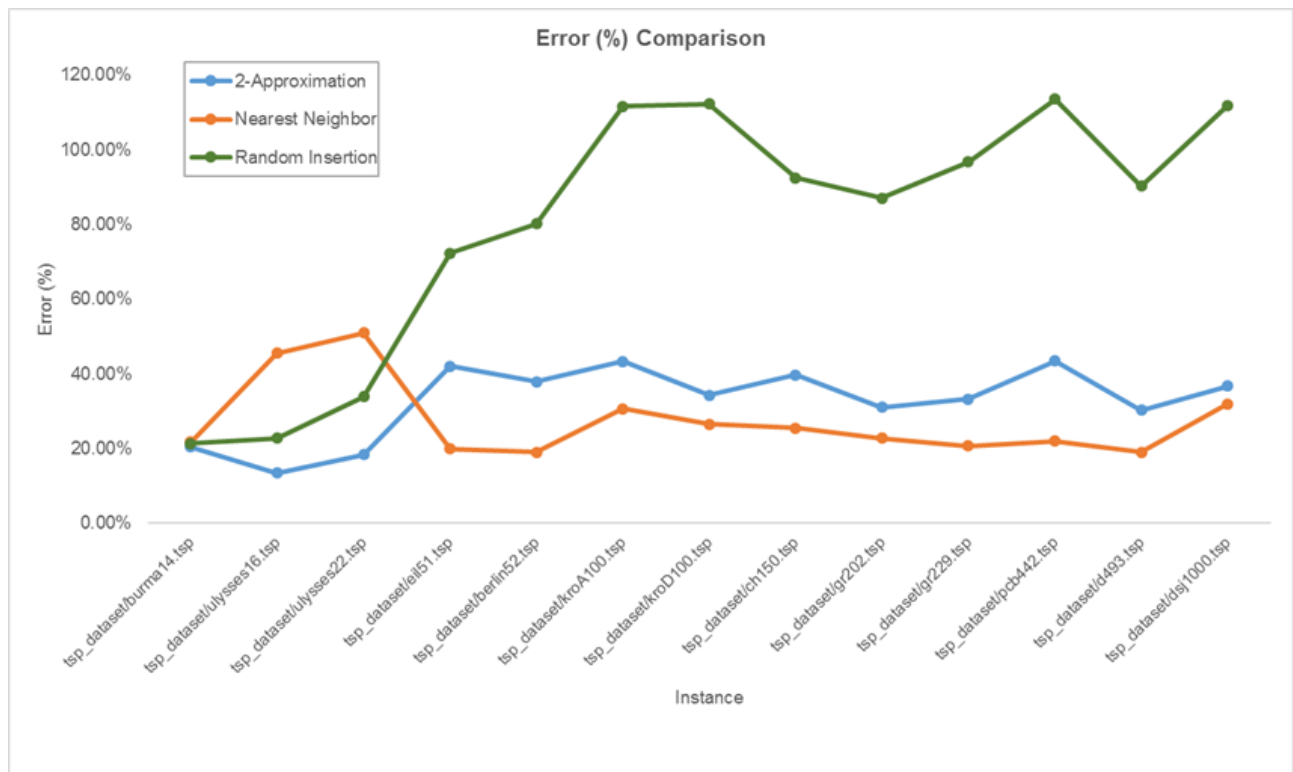


Figure 3: Error comparison

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## 6 Conclusion

From the Error Comparison graph and Results table, we can observe that for instances with less than 50 coordinates the approximation error is below or just at 50% for all three algorithms. However, as the number of coordinates grow past 50, the approximation error of the random insertion algorithm continues to grow past 100%, while the approximation error of the 2-approximate and nearest neighbor algorithms remain at or below 50%. From the Time Comparison graph and Results table, we can observe that for instances less than 442 coordinates, all three algorithms have an execution time below 1 second, although there is a clear distinction with the random insertion algorithm time increasing more quickly than the nearest neighbor and 2-approximate algorithms. After 442 coordinates, the random insertion algorithm time grows rapidly, taking over a minute for the largest instance with 1000 coordinates, where the nearest neighbor and 2-approximate algorithms always remain below half a second.

With respect to the approximation error, the nearest neighbor algorithm does better for all but three of the instances. Interestingly, the three instances for which nearest neighbor is outperformed on are the three lowest coordinate instances. For the three lowest coordinate instances, the 2 approximate algorithm outperforms the other two.

Considering both the approximation error and the execution time, we observe that the nearest neighbor algorithm is the most efficient.