

## Problem 1

1. Give pseudo code for an algorithm that takes as input [a directed, weighted graph, a source node  $s$  and a destination node  $d$ ] and returns the average of the lengths of all of the paths leading from  $s$  to  $d$ .

```
AVERAGE_DISTANCE(g, s, d)
```

```
v = array(0...g.length)
```

```
for i = 0...g.length
```

```
    v[i] = false
```

```
l = list()
```

```
DFS(g, s, d, v, l, 0)
```

```
return GET_AVERAGE(l)
```

```
DFS(g, s, d, v, l, c)
```

```
v[s] = true
```

```
if s == d
```

```
    l.add(c)
```

```
    v[s] = false
```

```
    return
```

```
for i...g.length
```

```
    if g[s][d] != 0
```

```
        if v[i] == false
```

```
            DFS(g, i, d, v, l, c + 1)
```

```
v[s] = false
```

```
GET_AVERAGE(l)
```

```
su = 0
```

```
ss = l.size
```

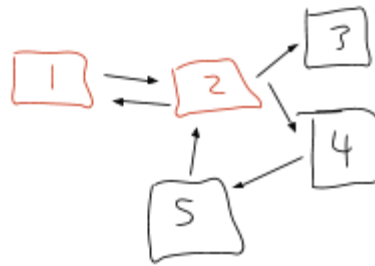
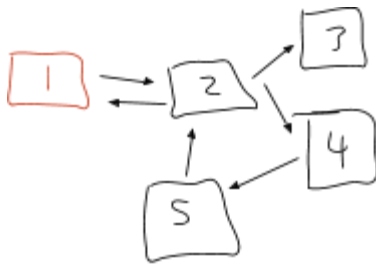
```
for i = 0...ss
```

```
    su = su + l[i]
```

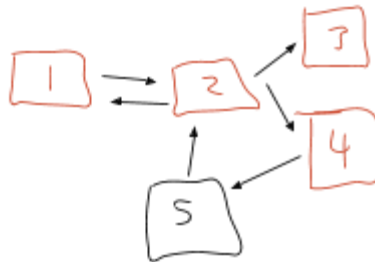
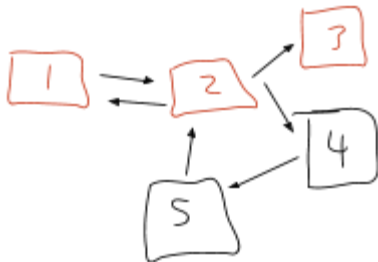
```
a = su / ss
```

```
return a
```

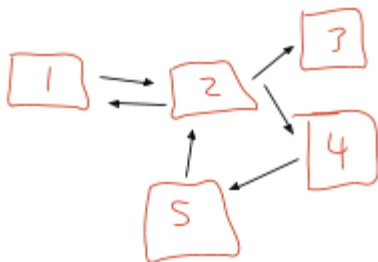
2. Give an illustration of your algorithm



In DFS, I go deep down a path until I reach a "dead-end", then I retrace back only as far as to find another valid path to go down.



In my algorithm, whenever I reach the destination I was looking for, I snapshot how many edges I passed to get there, to be later aggregated then used to find the average.



3. Do a time complexity analysis (big oh) of your algorithm.

$O(N!)$ . The time complexity here will be the same time complexity as counting the number of possible paths, which is shown to be  $O(N!)$  here: [Count all possible paths between two vertices](#).

4. Describe the choice(s) that you have made to make your algorithm run faster.

DFS makes the most sense for this algorithm.

I keep track of the amount of edges we are processing in the recursion with the "c + 1". When the recursion has found, if it exists, the destination, then I store the amount of edges it took to get there in a dynamic list. You cannot do averages along the way, you must first have all the data items. Once I found all the lengths, I simply summed them up, then used that number to divide by the amount of paths found. This returns the average.

5. Write Java code that implements your algorithm. The code should be added to the base that we have in the sample key for Asgn3.

See included files.

## Problem 2

1. Consider the hard problem of determining all the shortest round trips from a source node to every other node in a directed, weighted graph. Give pseudo code for an algorithm that takes as input [a directed, weighted graph and a source node s] and prints all the shortest round trips from the source node to every other node in a graph. Your algorithm must first pre-process the graph by removing the edges with highest weight (e.g, if the edge weights range from 1 to 7, then all the edges with a weight of 7 must be first removed from the graph before the round trips are computed). Assume that a round trip will still exist from the source to every other node after these edges are removed.

### **SHORTEST\_ROUNDTRIP\_PRE\_PROCESSED(g, s)**

```
p = array(0...g.length)(0...g.length)
```

```
h = 0
```

```
l = {}
```

```
for i = 0...g.length
```

```
  for j = 0...g.length
```

```
    w = g[i][j]
```

```
    p[i][j] = w
```

```
    if w >= h
```

```
      h = w
```

```
      l[h].add([i,j])
```

```
for i = 0...l[h].length
```

```
  pa = l[h]
```

```
  p[pa[0]][pa[1]] = 0
```

```
for j = 1...p.length
```

```
  SHORTEST_ROUND_TRIP(p, s, j)
```

### **SHORTEST\_ROUND\_TRIP(G, s, d)**

```
DIJKSTRA(G, s, d)
```

DIJKSTRA(G, d, s)

**DIJKSTRA(G, S, D)**

```
for each vertex V in G
  distance[V] <- infinite
  parents[V] <- -1
  visited[V] <- false
```

```
distance[S] <- 0
parents[S] <- -1
```

```
for i in 0...G.length
  u = FIND_NEXT_LEAST_WEIGHT(distance, visited)
```

```
visited[u] = true
```

```
for v in 0...G.length
  p <- distance[u] + G[u][v]
  if !visited[v] and p < distance[v]
    distance[v] = p
    parents[v] = u
```

```
PRINT_SOLUTION(distance, visited)
```

**PRINT\_SOLUTION(distance, visited)**

// Not provided here

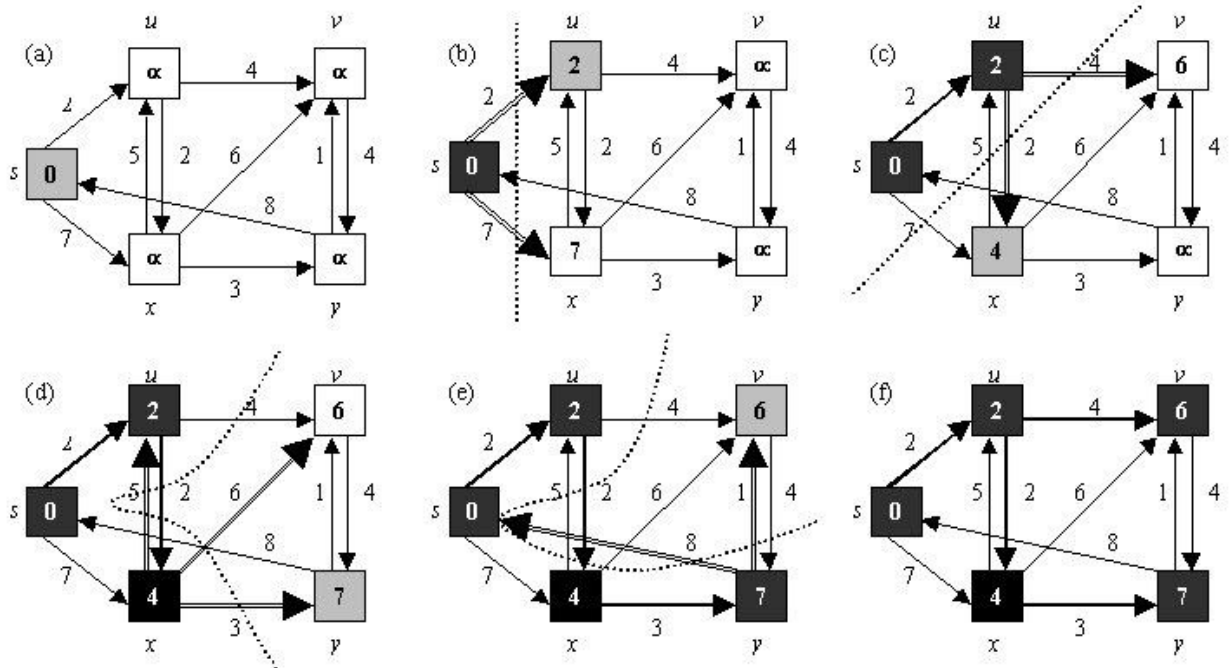
2. Give a pictorial illustration of your algorithm

```
{
  ...
  7: [[1, 1], [2, 3], [4, 2]]
}
```

1	2	3	4	5
5	7	3	4	2
2	1	3	7	1
4	1	3	0	5
0	0	7	1	2



1	2	3	4	5
5	0	3	4	2
2	1	3	0	1
4	1	3	0	5
0	0	0	1	2



(Source of Dijkstra illustration: [Dijkstra's Algorithm](#))

### 3. Do a time complexity analysis (big oh) of your algorithm.

The time complexity should be 2 Dijkstra algorithms, with the advantage that a number of max edges ( $M$ ) have been removed. In order to find that max edge, I have to iterate through all the edges ( $E$ ). The amount of work this makes is  $O(E(E-M)\log V)$ .

### 4. Describe the choice(s) that you have made to make your algorithm run faster.

I did not work on the dijkstra implementation already provided. I can speak to the optimization piece of removing the largest edges.

I made a copy of the original graph, only because it was unclear whether or not we could mutate the original input. Assuming that we couldn't (which seems like good practice), I went through and did a basic copy of the 2D array. I also paid attention to the highest weight found so far. Now, this next part is the part that I am proud of. I wanted to keep track of all the locations of the highest weights, so that we didn't have to iterate through the entire graph again to find them. I did this by using a HashMap in Java (noted as `{}` in pseudocode). I kept track of where the highest values were in this map, so that I could simply go back and just assign those locations to be "0". I didn't need to keep track of all the weight's locations, just the locations that are currently thought to be the highest. Once another highest weight was found, I only added to that list. This reduced an iteration through the entire graph.

5. **Write Java code that implements your algorithm. The code should be added to the base that we have in the sample key for Asgn3.**

See included files