CSOR W4231 Analysis of Algorithms I - Spring 2021 Homework #3

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Problem 1

At a high level, two modified versions of the DFS algorithm are constructed and used to traverse the input tree, iteratively matching unmatched nodes. The first modified DFS algorithm (DFS-Leaves), traverses the tree to match all leaf nodes and the neighbor of each leaf node provided that they have not yet been matched. The second modified version of DFS algorithm (DFS-Match) traverses the tree to match the remaining internal nodes that have not yet been matched. The output of the algorithm is either True, indicating that the tree has a perfect matching, or False, indicating that no perfect matching exists.

The main procedure, PerfectMatchTree, initializes an array matched that records which nodes have been matched, using a binary $\{0,1\}$ indicator (0 if node is not matched, 1 if matched). Next, the algorithm evaluates the tree and its attributes to determine whether it has a perfect matching. If |V| = 0, the algorithm returns True, since a graph on zero nodes vacuously contains a perfect matching. If |V| is odd, the algorithm returns False since a perfect matching can only exist if the graph has an even number of vertices. Otherwise, the algorithm runs DFS-Leaves first and then DFS-Match. If the sum of all entries in the matched array is |V| then a perfect matching exists and the algorithm returns True; otherwise, a perfect matching does not exist and the algorithm returns False.

Pseudocode and proof of correctness provided on the following two pages.

Running Time: Referring to the run-time comments in the pseudocode (in blue), lines 2 and $\overline{3}$ require constant time, the for-loop to initialize the matched array (lines 4-6) requires O(n) time, and lines 7 through 10 requires constant time. The DFS-Leaves procedure requires the usual O(n+m) run time for the original DFS procedure and O(n) time to match all leaves, for a total run time of O(n+m) for DFS-Leaves. The DFS-Match procedure also requires the usual O(n+m) run time for the original DFS procedure and O(m) time for the for-loop at lines 37-42 to run for each node; indeed, for each node u, the for-loop iterates deg(u) times. The total run time of DFS-Match is therefore O(n+m). Therefore, the total run time of the entire algorithm is

$$O(1) + O(n) + O(n+m) + O(n+m) = O(n+m)$$

```
1: PerfectMatchTree(T = (V, E))
       n = size(V)
                                    // Define number of nodes in T. Running time: O(1)
 2:
       matched = []
 3:
                                           // Initialize empty array. Running time: O(1)
       for u \in V do
                            // Initialize each entry in array to False. Running time: O(n)
 4:
          matched[u] = 0
 5:
       end for
 6:
      if n == 0 then
                                                                  // Running time: O(1)
 7:
          return 'True'
 8:
       else if n is odd then
                                                                  // Running time: O(1)
 9:
          return 'False'
10:
11:
       else
          Run DFS-Leaves(T)
                                                        // Total running time: O(n+m)
12:
                                                        // Total running time: O(n+m)
13:
          Run DFS-Match(T)
          if sum(matched) == n then
                                                                  // Running time: O(n)
14:
             return 'True'
15:
          else
16:
             return 'False'
17:
          end if
18:
       end if
19:
20: end
21:
    DFS-Leaves(T = (V, E))
                                                     // Only the modified portion shown
22:
23: //Add the following to the beginning of the Search procedure in the DFS pseudocode
      if degree(u) == 1 and matched[u] == 0 then
                                                                  // Running time: O(1)
24:
          if matched[neighbor(u)] == 0 then
25:
             matched[u] = 1
26:
27:
             matched[neighbor(u)] = 1
28:
          else
             return 'False'
29:
          end if
30:
       end if
31:
       return matched
                                              // Add to the end of the Search procedure
32:
33: end
34:
35: DFS-Match(T = (V, E))
                                                     // Only the modified portion shown
36: //Add the following to the beginning of the Search procedure in the DFS pseudocode
       for (u, v) \in E do
                                                  // Iterates deg(u) times for each u \in V
37:
          if matched[u] == 0 and matched[v] == 0 then
38:
             matched[u] = 1
39:
             matched[v] = 1
40:
41:
          end if
       end for
42:
      return matched
                                              // Add to the end of the Search procedure
43:
44: end
```

<u>Correctness</u>: The algorithm first matches all leaf nodes and their neighbors, provided that they have not yet been matched. Because leaf nodes have exactly one neighbor, a perfect matching necessarily includes all edges incident on leaf nodes. If the neighbor of any unmatched leaf node has already been matched then a perfect matching does not exist.

A DP approach can be used to solve this problem. At a high-level, the algorithm counts the number of symbols in S' that sequentially match S (in order). If the number of sequentially matched symbols is equal to the number of symbols in S' (= m), the algorithm returns True. Otherwise, the algorithm returns False. The subproblems consist of comparing the symbol S'[k] to the symbol S[t] for fixed k and t, where $k \leq t$. The subproblems are iteratively solved one by one, in increasing order. If S'[k] = S[t] for some fixed values k and t, then we add 1 to the count, which is stored in an array M of size n + 1 (includes the boundary condition), from the bottom up. In particular,

$$M[i] = \begin{cases} M[i-1] + 1 , & \text{if } S[i] = S'[j] \\ M[i-1] , & \text{if } S[i] \neq S'[j] \end{cases}$$

Boundary condition: M[0] = 0. Pseudocode is provided below.

```
SubSequence(S, S')
       n = length(S)
2:
       m = length(S')
3:
       if m > n then
4:
          return 'False'
5:
       end if
6:
       if m == 0 then
7:
          return 'True'
8:
       end if
9:
       Let M = array(n+1)
10:
       M[0] = 0
11:
       i = 1
12:
       i = 1
13:
       while i \leq n and j \leq m do
14:
          if S[i] == S'[j] then
15:
              M[i] = M[i-1] + 1
16:
17:
              i = i + 1
              j = j + 1
18:
          else if S[i] \neq S'[j] then
19:
              M[i] = M[i-1]
20:
              i = i + 1
21:
          end if
22:
       end while
23:
       if M[n] == m then
24:
          return 'True'
25:
       else return 'False'
26:
       end if
27:
28: end
```

Running Time: The symbols in the sequence S' are compared to the symbols in S over a maximum of n iterations. Indeed, there are n symbols in S, and if m > n then no comparison is made since then S' cannot be a subsequence of S. Therefore, the running time is O(n).

Space: The algorithm requires an additional $\Theta(n)$ space for the array M. The array M is filled from the bottom up.

<u>Correctness</u>: If m=0, the algorithm returns True regardless of the size of S since an empty sequence is a subsequence of all sequences. Otherwise, the algorithm sequentially and iteratively compares the symbols in S' to the symbols in S, adding 1 to the number of matched symbols if a given subproblem has a truth value of True. If S'[k] = S[t] for some fixed values k and t, the algorithm increments both S' and S to their next symbols so that S'[k+1] is compared to S[t+1]; otherwise, it increments to the subproblem comparing S'[k] to S[t+1]. Therefore, if elements S' are all present in the same order in S, the algorithm will account for it.

Construct a modified version of Dijkstra's algorithm such that it accounts for both the edge weights and the vertex costs. In particular, for each $(u, v) \in E$ where $u \in S$ and $v \in V - S$, pick v such that $dist[u] + w_{uv} + c_v$ is a minimum among all nodes in V - S. In particular, modify the Update procedure such that dist[v] maintains the minimum weight and cost from s to u, for all $u \in V - S$. Additionally, modify the Initialize procedure to initialize $dist[s] = c_s$. Last, Dijkstra is modified to return the dist array.

Pseudocode:

```
1: Note: Only the modified portion of each procedure is shown
    Dijkstra-Modified(G = (V, E, w, c), s):
       Initialize(G, s)
4:
       \dots Update(u, v)
6:
7:
       return dist
8: end
9:
    Initialize(G, s)
10:
11:
       dist[s] = c_s
12:
13: end
14:
    Update(u, v):
15:
       if dist[v] > dist[u] + w_{uv} + c_v then
16:
          dist[v] = dist[u] + w_{uv} + c_v
17:
          prev[v] = u
18:
       end if
19:
20: end
```

Running Time: The running time is the same as the running time of Dijkstra's algorithm. That is, the run time is $O(n \log n + m \log n)$.

<u>Correctness</u>: Correctness follows from the validity of Dijkstra's Algorithm. In particular, the modified algorithm returns the shortest

This can be solved using a greedy approach. In particular, we can progressively choose a locally optimal position for each of the guards. In doing so, the position of all guards will be globally optimal. That is, the minimum number of guards will be used to protect all paintings (to be shown in proof of correctness). The first guard should be placed at either $x_1 + 1$ or $x_n - 1$ so that paintings on both his left and his right side, within a distance of 1, are protected (if any paintings exist in those positions); this ensures that as many paintings as possible are being protected by the first guard. Without loss of generality, the first guard will be placed at position $x_1 + 1$ on the line L. If any paintings are positioned before $x_1 + 2$ on the line, we do not need to place any additional guards to protect those paintings since they are already being protected by the first guard. Suppose the positions of paintings x_2, \ldots, x_k are less than $x_1 + 2$ (i.e., protected by the first guard), with $k \in \{2, \ldots, n\}$. The second guard should be placed using a similar approach. That is, place the second guard at position $x_{k+1} + 1$ so that he protects all paintings positioned on the interval $[x_{k+1}, x_{k+1} + 2]$, on the line L. Continue in this way until all paintings are guarded.

```
MinNumGuards(\{x_1, x_2, \ldots, x_n\})
       X = array(n)
 2:
       for i = 1 to n do
 3:
          X[i] = x_i
 4:
       end for
 5:
       quardPositions = MinGuardPlacement(X)
 6:
 7:
      minNumGuards = length(guardPositions)
       return quardPositions, minNumGuards
                                                    // Returns positions and # of guards
 8:
 9: end
10:
    MinGuardPlacement(X)
11:
                                            // Initialize empty array for guard positions.
       quard = []
12:
       guard[1] = X[1] + 1
13:
       i = 1
                                         //j used to used to iterate through each guard.
14:
       for k = 2 to n do
15:
          if X[k] > guard[j] + 1 then
16:
             guard[j+1] = X[k] + 1
                                          // Place next guard at locally optimal position.
17:
                                                  // Increment j for next guard position.
             j = j + 1
18:
          end if
19:
       end for
20:
21:
      return guard
                                                               // Returns guard positions
22: end
```

Running Time: Constructing the array X in the MinNumGuards procedure requires O(n) time. The for-loop in the MinGuardPlacement procedure iterates n-1 times and each of the entries of X and guard are evaluated exactly one time. Thus, the MinGuardPlacement procedure requires O(n) time. Therefore, the total run-time is O(n).

<u>Correctness</u>: Let g_i denote the position of guard i for $i \in \{1, ..., n\}$. We induct on n.

Base Case (n = 1): Exactly one guard is used to protect the painting at position x_1 , and is placed at position $g_1 = x_1 + 1$. Clearly, the minimum number of guards required to protect the one painting is 1.

Induction Hypothesis: Assume the algorithm outputs the minimum number of guards required to protect n-1 paintings at positions $x_1, x_2, \ldots, x_{n-1}$.

Inductive Step: Let K denote the minimum number of guard to protect the n-1 paintings. That is, suppose that paintings at positions x_i, \ldots, x_{n-1} are protected by the K^{th} guard, for $i \in \{2, \ldots, n-1\}$. If $x_n \leq g_K + 1$, then the number of guards required to protect all n paintings output by the algorithm is K, which is the minimum number of guards required to protect the first n-1 paintings, from the inductive hypothesis. Thus, the minimum number of guards required for all n paintings when $x_n \leq g_K + 1$ is K. Therefore, the algorithm outputs the minimum number of guards required for all n paintings when $x_n \leq g_K + 1$. On the other hand, if $x_n > g_K + 1$, the algorithm places a guard at position $x_n + 1$ since the painting at position x_n is too far from the guard at position g_k to be able to protect it. That is, an additional guard must be used to protect the n^{th} painting. Then the algorithm outputs K+1 and the positions of each of the guards. Because K is the minimum number of guards required for the first n-1 paintings, then K+1 is the minimum number required for all n paintings when $x_n > g_K + 1$. Therefore, the algorithm outputs the minimum number of guards in all cases.

This can be solved using a DP approach. Let OPT(j) be the minimum total cost to store books in some libraries. We know that a copy is always stored at L_n . Suppose i is the largest index less than n such that a copy is stored at L_i in the optimal solution, then $OPT(n) = c_n + OPT(n-1)$.

Let the subproblems be such that we must choose L_1, \ldots, L_j to store copies of the book at. There are two cases: one in which we incur a storage cost c_j for storing at library L_j , or there is a user delay penalty (k-j) if a copy is not stored at L_i . That is,

$$OPT(j) = \min \begin{cases} c_j + OPT(j-1) &, & \text{if a copy is stored at } L_k \\ (i-j) + OPT(j-1) &, & \text{if a copy is not stored at } L_k & \text{for some } i > j \end{cases}$$

Let $d_{j,i} = i - j$ denote the user delay when a book is not available at L_j , but available at L_i , for i > j. We then have

$$OPT(j) = \min\{c_j + OPT(j-1), d_{j,i} + OPT(j-1)\}\$$

Set a boundary condition OPT(0) = 0. Use an array M to store the values of OPT(j).

Pseudocode provided on the next page - does not fit on this page.

Running Time: There are n subproblems. The bottleneck of the algorithm are the two nested for-loops, each of which iterates n times, in the MinStorageCost procedure. Therefore, the run time is $O(n^2)$.

Space: Additional space requirement is $\Theta(n)$ for the array M.

<u>Correctness</u>: The recurrence provided in the algorithm computes the minimum cost between storing a copy at library L_j or not storing a copy at L_j , for $j \in \{1, ..., n\}$. The minimal cost at iteration j depends on both the cost at L_j , or the user delay at server j and on the minimum cost at iteration j-1. Because each M[j-1] returns the minimum cost, M[j] must also return the minimum cost. To show this, induct on j:

Base case (j = 0): M[0] = 0 is trivially the minimum cost for j = 0.

Induction Hypothesis: Suppose M[j-1] returns the minimum cost for L_1, \ldots, L_{j-1} .

Inductive Step: $M[j] = \min\{c_j + M[j-1], \ d_{j,i} + M[j-1]\}$. From the induction hypothesis, M[j-1] returns the minimum cost, then computing the minimum reduces to $\min\{c_j, d_{j,i}\}$ which will return the minimum. Thus, M[j] returns the minimum total cost for L_1, \ldots, L_j . Similarly, M[n] returns the minimum total cost for L_1, \ldots, L_n

```
MinStorageCost(n):
       Initialize array M[0, \ldots, n]
 2:
       Set M[0] = 0
 3:
       for j = 1 to n do
 4:
          for i \geq j do
 5:
             d_{j,i} = i - j
                                        // Compute and store all possible user delay values
 6:
          end for
 7:
       end for
 8:
       for j = 1 to n do
 9:
          M[j] = \min\{c_j + M[j-1], d_{j,i} + M[j-1]\}
10:
       end for
11:
       return M[n]
12:
13: end
14:
    OptLibraries(j):
15:
       if j == 0 then return
16:
17:
          if M[j] == c_j + M[j-1] then
18:
              Output L_i
19:
              OptLibraries(j-1)
20:
          else if M[j] == d_{j,i} + M[j-1] then
21:
              OptLibraries(j-1)
22:
          end if
23:
       end if
24:
25: end
```