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Submission notes

- Due at 11:59 pm on Friday, November 13.
- Solutions must be typeset using one of the template files. For each problem, your answer must fit in the space provided (e.g. not spill onto the next page) **without** space-saving tricks like font/margin/line spacing changes.
- Upload a PDF version of your completed problem set to Gradescope.
- Teaching staff reserve the right to request original source/tex files during the grading process, so please retain these until an assignment has been returned.
- Please remember that for problem sets, collaboration with other students must be limited to a high-level discussion of solution strategies. If you do collaborate with other students in this way, you must identify the students and describe the nature of the collaboration. You are not allowed to create a group solution, and all work that you hand in must be written in your own words. Do not base your solution on any other written solution, regardless of the source.

1. (Candy Corn Walks)

My proposed algorithm has the following features:

- The algorithm takes input of (1) the adjacency list representing the graph G and (2) the starting vertex s . The adjacency list input should include the color of the incident edge for each neighbor as an integer: 1 for yellow, 2 for orange, and 3 for white.
- The vertices and edges in this algorithm represent the vertices and edges in the given graph G .
- The edges are directed, and each edge has an associated integer value according to its color in graph G .
- Each vertex will have an associated array of three boolean values, indexed 1, 2, and 3. The boolean at index k records whether the vertex has been “ k -visited”, which we’ll describe later.
- The algorithm will output a set containing the vertices that can be reached from the starting vertex via some candy corn walk.
- The algorithm makes use of a Breadth-First Search with some modifications:
 - The queue used in the BFS will hold not just the next vertex, but also the color of the incident edge. For example, a pair $(a, 2)$ at the front of the queue indicates that a is our next vertex and we’ll take an orange edge to get to it.
 - The queue will have the pair $(s, 0)$ as a start, where s is the input starting vertex.
 - The search will go on until the queue is empty. In each iteration, let’s say the algorithm pops the queue and gets the pair (x, k) . If $k = 3$, our algorithm will add x into the output set.
 - Next, unlike a regular BFS who’d add all x ’s unvisited neighbors into the queue, our algorithm will add a neighbor y only if (1) the color of xy is equal to either k or $k + 1$, and (2) y hasn’t been marked l -visited, where l is the color of xy . If the algorithm ends up adding y to the queue, it’ll then immediately mark y as l -visited.
- When the search is over, the algorithm returns the output set constructed during the search.

Let n and m be the number of vertices and edges in G , respectively. To analyze the run-time behavior of this algorithm, let’s break it down into the initialization phase and the search phase:

- The initialization phase assigns each vertex an array of three booleans filled with **false**. Since there are n vertices, this part will take $O(n)$.
- The search phase will visit every edge in G at most once, just like in a regular BFS, and it’ll visit every vertex at most three times. This is because a vertex is marked every time it’s added to the queue, but it only gets three marks maximum, after which it’d never get added anymore. The only exception is the starting vertex, which may appear up to four times in the queue, but that’s a constant. So, the search phase will take $O(3n + m) = O(n + m)$.

Therefore, since the most time-consuming part of the algorithm takes $O(n + m)$, the entire algorithm takes $O(n + m)$.

2. (Graph DP)

My proposed algorithm is as follows, whose first argument is an adjacency list representing the graph G .

Algorithm: max_st_weight

Input: hash_map<vertex, hash_set<vertex>> A , int l , vertex s , vertex t ,
function(vertex, vertex):number w

Output: number

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1 int  $n \leftarrow$  number of keys in  $A$ 
2 vertex  $L[1..n] \leftarrow$  topological sort  $A$ 
3 hash_map<vertex, number>  $M[0..l] \leftarrow$  filled with  $\{\}$ 
4 for int  $k = 0$  to  $l$  do
5    $M[k][t] \leftarrow 0$ 
6 for each vertex  $u$  in  $A$ 's keys do
7    $M[l][u] \leftarrow -\infty$ 
8 for int  $k = l - 1$  to  $0$  do
9   for int  $i = n - 1$  to  $1$  do
10    vertex  $u \leftarrow L[i]$ 
11     $M[k][u] \leftarrow -\infty$ 
12    for each vertex  $v$  in  $A[u]$  do
13       $M[k][u] \leftarrow \max(M[k][u], w(u, v) + M[k + 1][v])$ 
14 return  $M[0][s]$ 

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This algorithm uses the idea of the following backtracking algorithm, which returns the maximum weight from any vertex u to the only sink t using at most l edges when k edges are used so far:

$$\text{BT}(k, u) = \begin{cases} -\infty, & \text{if } k = l \\ 0, & \text{else if } u = t \\ \max[w(u \rightarrow v) + \text{BT}(k + 1, v) \text{ for } v \text{ in out-neighborhood of } u], & \text{otherwise} \end{cases}$$

Since the graph is acyclic, every step we take from u will be closer to t in terms of the number of steps left, which is advantageous because then any out-neighbor of u would not depend on u for its maximum weight to t . This paved the way for our dynamic programming algorithm:

- The algorithm can use memoization to avoid calculating the same value twice. I used a hash-map for its fast lookup when vertices are the keys.
- The topological sort conveniently gives us the order we need to calculate in (line 2).
- The two base cases are when l edges are used already (lines 6-7) and when the current vertex is t (lines 4-5).
- I filled the memoization table in *reverse* order (lines 8-9) since calculating the answer for (k, u) depends on $k + 1$ and u 's out-neighbors who come later in the topological order.
- The final answer is given by “when the current vertex is s where 0 edges are used so far” (line 14).

Let's analyze the dynamic programming algorithm's run-time complexity. First, we know that topological sort runs in $O(n + m)$, where n and m are the number of vertices and edges in G . Assuming that the weight-function w and a vertex's hashing both have constant running time, the “table-filling” portion takes $O(l \cdot (n + \sum_{v \in V} \text{outdegree}(v))) = O(l \cdot (n + m))$. Therefore, the entire algorithm runs in $O(l \cdot (n + m))$.

3. (MST Properties)

a. Suppose that G is cyclic, and consider a cycle $v_1v_2, v_2v_3, \dots, v_nv_1$, where v_1, \dots, v_n are the vertices involved. Let v_1v_2 be the (unique) maximum-weight edge in this cycle. Assume that v_1v_2 is in a minimum spanning tree T of G , and let's see why this cannot happen:

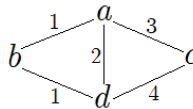
- **Claim.** We could replace the edge v_1v_2 in T with some other edge from the cycle, and T would still be a spanning tree of G .
- **Proof.** Since T is a spanning tree, we know that T is connected, but removing any edge from T disconnects it and results in exactly two components.

Let's remove v_1v_2 from T . After that, every vertex in G is in one of two components: the one v_1 is in, which we'll call C_1 , or the one with v_2 , called C_2 . Since then, there is at least one place in $v_1..n$ where two neighboring vertices v_i and v_j are in different components, *other than v_1 and v_2* . We can guarantee this because we know that v_1 and v_2 are in different components, and every other vertex in the cycle must fall into either C_1 or C_2 . If there are multiple instances of neighboring v_i and v_j where they're in different components, just pick any of them as long as they're not v_1 and v_2 .

Now, since v_i and v_j are neighbors and are in different components, if we add the edge v_iv_j to T , we'll for sure end up with a tree. Moreover, we know that the new T is still a spanning tree because it has the same number of edges as before, which implies that it still includes every vertex in G , by the characteristic that any tree has $|V| - 1$ edges. \square

We've just shown that, if v_1v_2 were in an MST, we could always replace it with some other edge from the same cycle and get a spanning tree with less total weight, because v_1v_2 was the maximum-weight edge in the cycle. T wasn't even minimum to begin with, so our assumption in the beginning must've been wrong. Therefore, the minimum spanning tree of G cannot contain a maximum-weight edge from any cycle. \square

b. The following graph makes a counterexample to the statement:



The minimum spanning tree of the above graph would include edges $\{ab, bd, ac\}$. The cycle $\{ac, cd, ad\}$ has ad being the minimum-weight edge, which is not included in the MST.

4. (Light-st Walk)

My proposed algorithm has the following features:

- The algorithm takes input of (1) the adjacency list representing the graph G , (2) the starting vertex s , (3) the destination vertex t , and (4) the weight function w .
- The vertices and edges in this algorithm represent the vertices and edges in the given graph G .
- The edges are undirected and do not have associated values.
- Each vertex will have three associated values: an integer recording the minimum weight from the starting vertex, a boolean recording whether the vertex has been visited, and a place to hold the vertex's predecessor. The initial minimum weight will be set to infinite, except if the vertex is s , which would be set to 0.
- The algorithm will output a list of vertices representing the path with the minimum total weight possible to get from s to t .
- The algorithm makes use of the Nonnegative Dijkstra's Algorithm with just one small modification:
 - The priority queue will hold a pair: the next vertex and the current weight, just like in a regular Nonnegative Dijkstra's Algorithm. *We'll use a binary min-heap as the priority queue.*
 - The priority queue will have the pair $(s, 0)$ as a start, where s is the input starting vertex.
 - The search will go on until the queue is empty, by default. In each iteration, let's say the algorithm pops the queue and gets the pair (x, k) . The algorithm will skip processing x if it's already visited, or otherwise it'll now mark x as visited. If $x = t$, the search ends immediately, and the algorithm returns the path by going over x 's predecessors.
 - Next, for each x 's neighbor y , the algorithm will add $(y, k + w(y))$ to the priority queue only if (1) $k + w(y)$ is less than y 's recorded minimum weight, and (2) y hasn't been marked visited. If the algorithm ends up adding y to the queue, it'll then record x as y 's predecessor. The modification here is that the weight function is called on the next vertex instead of on the incident edge.
- If the algorithm makes it to when the Nonnegative Dijkstra's Algorithm is over, it'll report that there isn't a path from s to t , which might be in the form of an error.

Let n and m be the number of vertices and edges in G , respectively. To analyze the run-time behavior of this algorithm, let's break it down into the initialization phase and the search phase:

- The initialization phase assigns each vertex its three associated values. Since there are n vertices, this part will take $O(n)$.
- The search phase will run in $O(n + m \log m)$, just like in a regular Nonnegative Dijkstra's Algorithm, because the only modification we did was changing how each edge weight is calculated by calling a different function (the vertex-weight function, in this case). Each vertex and each edge will still be visited the same maximum number of times just like in a Nonnegative Dijkstra's Algorithm.

Therefore, since the most time-consuming part of the algorithm takes $O(n + m \log m)$, the entire algorithm takes $O(n + m \log m)$.

5. (MST Updates)

a. My proposed algorithm, upon receiving the input, runs as follows:

- If $e \in T$, since T was a minimum spanning tree to start with, no further actions are needed, and the algorithm ends here.
- Otherwise, add e to T .
- Now, T should contain a cycle, and we can find that cycle using a Depth-First Search with the starting vertex being an endpoint of e . Let S be the set of edges in that cycle.
- Find the maximum-weight edge in S and remove it from T .
- T is now the updated MST.

b. My proposed algorithm, upon receiving the input, runs as follows:

- If $e \notin T$, since T was a minimum spanning tree to start with, no further actions are needed, and the algorithm ends here.
- Otherwise, remove e from T .
- Now, T should be divided into two connected components. Find all the vertices that are reachable from one endpoint of e with a Depth-First Search, and call that set of vertices R .
- Let S be the set of edges that have exactly one endpoint in R .
- Find the minimum-weight edge in S and add it to T .
- T is now the updated MST.