CS/ECE 374 A (Spring 2022) Homework 10 Solutions

Problem 10.1: Consider the following geometric matching problem: Given a set A of n points and a set B of n points in 2D, find a set of n pairs $S = \{(a_1, b_1), \ldots, (a_n, b_n)\}$, with $\{a_1, \ldots, a_n\} = A$ and $\{b_1, \ldots, b_n\} = B$, minimizing $f(S) = \sum_{i=1}^n d(a_i, b_i)$. Here, $d(a_i, b_i)$ denotes the Euclidean distance between a_i and b_i (which you may assume can be computed in O(1) time).

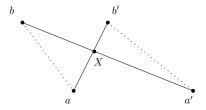
Assume that all points in A have y-coordinate equal to 0 and all points in B have y-coordinate equal to 1. (Thus, all points lie on two horizontal lines.) The points are not sorted. See the example below, which shows a solution that is definitely not optimal.

Assignment Project Exam Help

- (a) (20 pts) Consider the following greedy strategy: pick a pair $(a, b) \in A \times B$ minimizing d(a, b); then remove a from A and b from B, and repeat. Give a counterexample showing that this algorithm respectively a form A and B are the following that the same A and B are the following A are the following A and B are the following A are the following A and B are the following A and B are the following A are the following A and B are the following A are
- (b) (40 pts) Let a be the point in A with the smallest x-coordinate. Let b be the point in B with the smallest x-coordinate. Consider a solution S in which a is paired with some point b' with b' and the paired pair paired with b' with b' and b' with b' and b' with b' and b' with b' are b' with b
- (c) (40 pts) Now give a correct greedy algorithm to solve the problem. (The correctness should follow from (b).) Analyze the running time.

Solution:

- (a) One counterexample is $A = \{(0,0), (1,0)\}$ and $B = \{(1,1), (2,1)\}$.
 - This greedy strategy would pair (1,0) with (1,1), since the pair has the smallest distance 1. Then (0,0) would be paired with (2,1), of distance $\sqrt{5}$. The total cost is $1+\sqrt{5}>3.236$.
 - But the optimal solution is to pair (0,0) with (1,1), and (1,0) with (2,1), with cost $2\sqrt{2} < 2.829$.
- (b) By definition of a and b, we know that a is left of a' and b is right of b', as shown in the figure below. Let X be the intersection of the lines ab' and a'b.



By the triangle inequality, we have

$$d(a,b) + d(a',b') < (d(a,X) + d(X,b)) + (d(a',X) + d(X,b'))$$

= $(d(a,X) + d(X,b')) + (d(a',X) + d(X,b))$
= $d(a,b') + d(a',b).$

Create a new solution S' from S by deleting the pairs (a, b') and (a', b) and inserting the pairs (a,b) and (a',b'). Then

$$f(S') = f(S) + (d(a,b) + d(a',b')) - (d(a,b') + d(a',b))$$

$$< f(S)$$

by the above inequality d(a,b) $P_{1}^{d(a',b')} < d(a,b') + d(a',b)$. ASSIGNMENT Project Exam Help (c) The algorithm is simple: pick the smallest a in A and the smallest b in B; output the

pair (a, b); remove a from A and b from B; repeat.

Correctness follows from (b), because if the optimal solution does not pair a with b, then there would led to be in with strolly smalle cost a contradiction.

To bound the running time, note that the algorithm can be equivalently redescribed as follows: sort the points a_1, \ldots, a_n of A in increasing x-order and the points b_1, \ldots, b_n of B in decreasing perfect reprint the pairs (a_1, b_1) ; c.s. (a_n, b_n) .

Since sorting takes $O(n \log n)$ time, the total time is $O(n \log n)$.

Problem 10.2: We are given an unweighted undirected connected graph G = (V, E) with n vertices and m edges (with $m \ge n-1$), We are also given two vertices $s,t \in V$ and an ordering of the edges $e_1, \ldots, e_m \in E$. Suppose the edges e_1, \ldots, e_m are deleted one by one in that order. We want to determine the first time when s and t become disconnected. In other words, we want to find the smallest index j such that s and t are not connected in the graph $G_j = (V, E - \{e_1, \dots, e_j\}).$

A naive approach to solve this problem is to run BFS/DFS on G_j for each j = 1, ..., m, but this would require O(mn) time. You will investigate a more efficient algorithm:

- (a) (80 pts) Define a weighted graph G' with the same vertices and edges as G, where edge e_i is given weight -i. Let T be the minimum spanning tree of G'. Let π be the path from s to t in T. Let j^* be the smallest index such that e_{j^*} is in π . Prove that the answer to the above problem is exactly j^* .
- (b) (20 pts) Following the approach in (a), analyze the running time needed to compute j^* .

Solution:

(a) It suffices to prove the following two claims:

Claim 1. s and t are connected in G_{j^*-1} .

Proof: Every edge e_i in π has $i \geq j^*$. So, the path π from s to t uses only edges in $E - \{e_1, \ldots, e_{j^*-1}\}$ and remains a path in G_{j^*-1} .

Claim 2. s and t are not connected in G_{j^*} .

Proof: $T - \{e_{j^*}\}$ has two connected components; call them S and V - S. We know that $s \in S$ and $t \in V - S$, or vice versa. By a known fact from class, the smallest-weight edge between S and V - S must be in the MST. Since e_{j^*} is the only edge between S and V - S in T, we know that e_{j^*} must be the smallest-weight edge between S and V - S. Thus, every edge e_i between S and V - S has weight at least as large as e_{j^*} , i.e., $i \leq j^*$, and so there is no edge between S and V - S in $E - \{e_1, \ldots, e_{j^*}\}$. So, S and S are in different components in S in S and S and S in S in S in S in S and S in S in

(b) We can compute the MST T in $O(n \log n + m)$ time by Prim's algorithm with Fibonacci heaps (or better with some of the prore advanced MST algorithms hat devered in class). The path π can be found in O(n) time by following parent pointers, assuming s is made the root). The index j^* can then be found in O(n) time by a linear scan over π . The total time is therefore $O(n \log n + m)$.

(Alternativell 1 veloc use Irlula 3 for the angle $O(m \log n)$ time, which is a little worse than Prim's unless the graph is sparse. But actually, in this application, the running time of Kruskal's algorithm can be improved to $O(m\alpha(m, n))$, which is better than as good as Prim's where $\alpha(\cdot)$ is the inverse Ackermann function; this is because the initial sorting step is trivial as the weights are just the negated indices from -m to -1, i.e., the edges are already given in decreasing order of weights.)

(*Note.* There is a more clever O(m)-time algorithm for this problem, which uses median finding and contractions to reduce the number of edges by a half in each round...)

Problem 10.3: Consider the following search problem:

MAX-DISJOINT-TRIPLES:

Input: a set S of n positive integers and an integer L.

Output: pairwise disjoint triples $\{a_1, b_1, c_1\}, \ldots, \{a_{k^*}, b_{k^*}, c_{k^*}\} \subseteq S$, maximizing the number of triples k^* , such that $a_i + b_i + c_i \leq L$ for each i.

For example, if $S = \{3, 10, 29, 30, 35, 55, 70, 83, 90\}$ and L = 100, an optimal solution is $\{3, 10, 83\}, \{29, 30, 35\}$, with two triples (there is no solution with three triples).

Consider the following decision problem:

DISJOINT-TRIPLES-DECISION:

Input: a set S of n positive integers, an integer L, and an integer k.

Output: True iff there exist k pairwise disjoint triples $\{a_1, b_1, c_1\}, \ldots, \{a_k, b_k, c_k\} \subseteq S$, such that $a_i + b_i + c_i \leq L$ for each i.

Prove that MAX-DISJOINT-TRIPLES has a polynomial-time algorithm iff DISJOINT-TRIPLES-DECISION has a polynomial-time algorithm.

(Note: One direction should be easy. For the other direction, see lab 12b for examples of this type of question. In MAX-DISJOINT-TRIPLES, the output is not the optimal value k^* but an optimal set of triples, although it may be helpful to give a subroutine to compute the optimal value k^* as a first step, as in the lab examples.)

Solution:

If MAX-DISJOINT-TRIPLES has a polynomial-time algorithm, then we can solve DISJOINT-TRIPLE-DECISION in polynomial time easily: find an optimal solution $\{\{a_1, b_1, c_1\}, \dots, \{a_{k^*}, b_{k^*}, c_{k^*}\}\}$, and then just return true iff $k \leq k^*$.

Conversely, suppose DISJOINT-TRIPLE-DECISION has a polynomial-time algorithm A. We first find the optimal value k^* . This can be done by calling A on the input (S, L, k) for $k = 1, 2, \dots, S$ and a suppose K. This requires O(n) calls to A and so takes polynomial time.

(Note: alternatively, with binary search, this requires just $O(\log n)$ calls to A.) After finding k^* , the triples and the finding k^* and the finding k^* are the finding k^* and the finding k^* and the finding k^* are the finding k^* and k^* are the finding k^* are the finding k^* and k^* are the finding k^* are the finding k^* are the finding k^* and k^* are the finding k^* are the finding k^* and k^* are the finding k^* and k^* are the finding k^* are the finding k^* and k^* are the finding k^* are the finding k^* and k^* are the finding k^* and

```
1. while k^* > 0 do
2. for every triple of three distinct elements a, b, c > 0 with a + b + c \le L do
3. call A on the input (S - \{a, b, c\}, L, k^* - 1)
4. if A returns true then
5. output \{a, b, c\}
6. S \leftarrow S - \{a, b, c\}, k^* \leftarrow k^* - 1
7. break (i.e., go back to line 1)
```

Analysis: There are $k^* \leq O(n)$ iterations of the outer while loop, and in each such iteration, we are looping through $O(n^3)$ triples a, b, c. Thus, the total number of calls to A is $O(n^4)$. So, if A runs in polynomial time, the whole algorithm runs in polynomial time.