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CMPSCI 611 Advanced Algorithms Midterm Exam Fall 2010

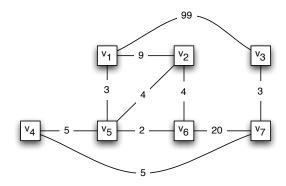
A. McGregor 19 October 2010

DIRECTIONS:

- Do not turn over the page until you are told to do so.
- This is a *closed book exam*. No communicating with other students, or looking at notes, or using electronic devices. You may ask the professor to clarify the meaning of a question but do so in a way that causes minimal disruption.
- If you finish early, you may leave early but do so as quietly as possible. The exam script should be given to the professor.
- There are five questions. All carry the same number of marks but some questions may be easier than others. Don't spend too long on a problem if you're stuck you may find that there are other easier questions.
- The front and back of the pages can be used for solutions. There are also a couple of blank pages at the end that can be used. If you are using these pages, clearly indicate which question you're answering. Further paper can be requested if required.
- The exam will finish at 3:45 pm.

1	/10
2	/10
3	/10
4	/10
5	/10
Total	/50
5	/10

Question 1: In the first part of this question, consider the following undirected graph where the value of each edge represents the length of that edge:



1. What is the total length of the shortest path between v_1 and v_3 ?

ANSWER: 3+5+5+3=16

2. What is the total length of the edges in a minimum spanning tree?

ANSWER: Edges of a MST are $(v_5, v_6), (v_1, v_5), (v_3, v_7), (v_5, v_2), (v_4, v_5), (v_4, v_7)$. Total weight is 2 + 3 + 3 + 4 + 5 + 5 = 22.

The next part of this question concerns an arbitrary weighted, undirected graph. For any two nodes u and v let $\delta_G(u, v)$ denote the length of the shortest path between u and v in the graph G. For each of the following statements, write whether they are true or false (no proofs required although including good reasoning may get partial credit even if you get the final answer wrong):

3. $\delta_T(u,v) = \delta_G(u,v)$ if T is a minimum spanning tree of G.

FALSE. Consider the graph G with nodes $\{v_1, v_2, v_3\}$ and edges $\{(v_1, v_2), (v_2, v_3), (v_3, v_1)\}$ where each edge has weight 1. All distances in G are 1 but in the minimum spanning tree, there is at least once pair of nodes that are distance 2 apart.

4. $\delta_G(u,v) \leq \delta_G(u,w)$ implies $\delta_{G'}(u,v) \leq \delta_{G'}(u,w)$ where G' is the graph formed by adding 1 to each of the edge lengths in G.

FALSE. Consider the graph with nodes $\{v, x, u, w\}$ and edges $\{(v, x), (x, u), (u, w)\}$ with weights 1,1, and 2.5 respectively.

5. $\delta_G(u,v) \leq \delta_G(u,w)$ implies $\delta_{G''}(u,v) \leq \delta_{G''}(u,w)$ where G'' is the graph formed by doubling each of the edge lengths in G.

TRUE.

Question 2: This question is about the subset system (E, \mathcal{I}) where:

$$E = \{a, b, c, d\}$$

$$\mathcal{I} = \{\emptyset, \{a\}, \{b\}, \{c\}, \{d\}, \{a, b\}, \{a, c\}, \{b, c\}, \{c, d\}, \{a, b, c\}\}\}$$

1. List all the maximal subsets in \mathcal{I} . Why can you conclude that (E,\mathcal{I}) is not a matroid?

ANSWER: The maximal subsets are $\{c,d\}$ and $\{a,b,c\}$. If the subset system was a matroid, all the maximal subsets would have the same cardinality. So the subset system is not a matroid.

2. Consider the weighting function w(a) = 1, w(b) = 2, w(c) = 3, and w(d) = 4. What solution is returned by the greedy algorithm? How does this compare to the optimal solution?

ANSWER: The greedy solution is $\{c,d\}$ and this has weight 7. This is the optimal solution!

3. Specify a weight function w on E such that the greedy algorithm doesn't return an optimal solution. Include the greedy solution and the optimal solution in your answer.

ANSWER: Consider the weight function w(a) = w(b) = w(c) = 2 and w(d) = 3. The greedy solution is $\{c,d\}$ and this has weight 5. The optimal solution is $\{a,b,c\}$ and this has weight 6.

4. Identify two subsets $i, j \subset E$ such that $(E, \mathcal{I} + i + j)$ is a matroid.

ANSWER: Add the subsets $\{b.d\}$ and $\{d,b,c\}$.

Question 3: Give a linear-time algorithm that takes as input a tree T and determines whether it has a perfect matching, i.e., whether there exists a subset of edges that touches each node exactly once.

ANSWER: Let V and E be the vertices and edges of T and let n = |V|. Consider the following algorithm:

- 1. $M \leftarrow \emptyset$
- 2. While $E \neq \emptyset$:
 - (a) Let $v \in V$ be a leaf (i.e., a node of degree 1) and let $(u, v) \in E$ be the unique edge incident on v
 - (b) $M \leftarrow M + (u, v)$
 - (c) $V \leftarrow V u v$ and remove all edges from E that are incident on u or v
- 3. If |M| = n/2 return "perfect matching" and if not, return "not perfect matching"

We first note that M is always a matching because whenever an edge e is added to M, all the edges sharing an endpoint with e are removed from the graph and therefore cannot subsequently be added to M. Hence if |M| = n/2, then we indeed have a perfect matching. Consequently we never return "perfect matching" if T does not have a perfect matching.

It remains to show that if T has a perfect matching then we return "perfect matching". Assume M^* is such a matching. If v is a leaf and $(u,v) \in E$ then (u,v) must be in M^* since otherwise v will not be covered. Hence, without loss of generality we may add (u,v) to M and remove all other edges incident on u. After removing u and v from V, the resulting subgraph has a perfect matching and we may recurse on this graph. In this way we find all the edges of M^* and hence $|M| = |M^*| = n/2$.

Question 4: A subsequence is palindromic if it is the same whether read left to right or right to left. For instance, the sequence

$$A, C, G, T, G, T, C, A, A, A, A, A, T, C, G$$

has many palindromic subsequences, including A, C, G, C, A and A, A, A, A. Devise an algorithm that takes a sequence x_1, x_2, \ldots, x_n and returns the length of the longest palindromic subsequence. For full marks the running time should be $O(n^2)$ but partial marks are available for less efficient solutions.

Hint: Let A[i,k] be the length of the longest palindromic subsequence of $x_i, x_{i+1}, \dots x_{i+k-1}$ and consider computing A[1,n] via dynamic programming.

ANSWER: Since a subsequence of length 1 is palindromic, A[i,1] = 1 for all i. For $i \in [n]$ and $k \in \{2, ..., n-i+1\}$

$$A[i,k] = \begin{cases} \max(A[i,k-1], A[i+1,k-1]) & \text{if } x_i \neq x_{i+k-1} \\ 2 + A[i+1,k-2] & \text{if } x_i = x_{i+k-1} \end{cases}$$

The rationale for the formula is that the longest palindromic subsequence of $x_i, x_{i+1}, \ldots, x_{i+k-1}$ either includes x_i and x_{i+k-1} (in which case they must be equal) or it doesn't. If it doesn't the longest palindromic subsequence of $x_i, x_{i+1}, \ldots, x_{i+k-1}$ is also a palindromic subsequence of $x_{i+1}, \ldots, x_{i+k-1}$ or x_i, \ldots, x_{i+k-2} . It it does then A[i, k] is 2 + A[i+1, k-2].

Hence A[1,n] can be constructed by the following algorithm:

- 1. For all $i \in [n]$, let A[i, 1] = 1
- 2. For all $i \in [n]$, let A[i,2] = 2 if $x_i = x_{i+1}$ and A[i,2] = 1 otherwise
- 3. For k = 3 to n:
 - (a) For i = 1 to n k + 1:

$$A[i,k] = \begin{cases} \max(A[i,k-1], A[i+1,k-1]) & \text{if } x_i \neq x_j \\ 2 + A[i+1,k-2] & \text{if } x_i = x_j \end{cases}$$

4. Return A[1,n]

The running time is clearly $O(n^2)$ since there are $O(n^2)$ values to be computed and each requires O(1) time.

Question 5: Given a sorted list of distinct integers $A[1], A[2], \ldots, A[n]$, you want to find out whether there is an index i for which A[i] = i. Give an algorithm that runs in time $O(\log n)$. Remember to prove that your algorithm is correct and analyze the running time.

For example, A = [-1, 1, 3, 5, 6, 7] does have such an index, i.e., A[3] = 3. But A = [0, 1, 2, 5, 6, 7] doesn't have such an index.

ANSWER: Consider the following algorithm.

- 1. $a \leftarrow 1$
- 2. $b \leftarrow n$
- 3. While $b a + 1 \ge 3$
 - (a) $k \leftarrow \lceil (a+b)/2 \rceil$
 - (b) If A[k] = k then return YES!
 - (c) If A[k] < k then $a \leftarrow k + 1$
 - (d) If A[k] > k then $b \leftarrow k 1$
- 4. If A[a] = a, A[a+1] = a+1, or A[a+2] = a+2 then return YES! Otherwise NO!

The running time is $O(\log n)$. This follows because the value of b-a+1 decreases by a factor 2 in each iteration. Let a_i and b_i be the values of a and b at the start of the i-th iteration. Then

$$b_{i+1} - a_{i+1} + 1 \le \max(b_i - k, k - a_i) \le (b_i - a_i + 1)/2$$

Hence, there are at most $O(\log n)$ iterations. Because each iteration takes O(1) time, the total running time is as claimed.

We now prove correctness. Note that we can never report YES when the answer is NO because we only answer YES when a test A[i] = i is passed for some i. It remains to prove that we never answer NO when the answer should be YES. To do this we prove that if there exists $i \in [n]$ such that A[i] = i, then at any point of the algorithm the values of a and b satisfy $a \le i \le b$. We show this by induction on the number of times the while loop is performed. Before the loop is performed, a = 1 and b = n and so the base case is trivially true. Assume that the claim is true before the jth time the loop is performed. If A[k] < k then we know all $\ell \le k$ satisfy $A[\ell] \le A[k] - (k - \ell) < \ell$ and $i \ge k + 1$ as required. Similarly, if A[k] > k then we know all $\ell \ge k$ satisfy $A[\ell] \ge A[k] + (\ell - k) > \ell$ and hence $i \le k - 1$ as required. Therefore, either we find the fixed point during some iteration or in the fourth line. Either way, we output YES.