Homework # 4

Due in class on Wednesday, November 15.

Reminder: you should put at the TOP LEFT corner of your homework: your name (LAST name first), your student ID number, and finally your tutorial section. Put each of these on a separate line. BELOW this information, include information on your collaborations, if any. **Points will be taken off for students who do not provide this information in the above manner.** Your cooperation with this will be greatly appreciated by the TA's.

Some of the following problems ask you to find an algorithm for a problem and explain its running time. For such problems, try to find an algorithm with as fast a running time as possible, ignoring constant factors.

- 1. Cormen, Leiserson and Rivest, problem 16-4, page 326: Professor McKenzie is consulting for the president of A.-B. Corporation, which is planning a company party. The company has a hierarchical structure; that is, the supervisor forms a tree rooted at the president. The personnel office has ranked each employee with a conviviality rating, which is a real number. In order to make the party fun for all attendees, the president does not want both an employee and his or her immediate supervisor to attend
 - (a) Describe an algorithm to make up the guest list. The goal shuld be to maximize the sum of the conviviality ratings of the guests.
 - (b) What is the running time of your algorithm? (Use of big-Oh notation is fine in expressing the running time). Explain your answer.
 - (c) How can the professor modify the algorithm to ensure that the president gets invited to the party, while maximizing the sum of the convivality ratings subject to this constraint?
- 2. You are given two strings of n characters each and an additional parameter k. In each string there are n k + 1 substrings of length k, and so there are $\Theta(n^2)$ pairs of substrings, where one substring is from one string and one is from the other. For a pair of substrings, define the *match count* as the number of opposing characters that match when the two substrings of length k are aligned. The problem is to compute the match-count for each of the $\Theta(n^2)$ pairs of substrings from the two strings.

Find an algorithm that solves this problem. Describe your algorithm and explain what is its running time.

3. Let x and y be binary strings of length n. The Hamming Distance between x and y, denoted by H(x, y), is defined to be the number of places where x and y disagree. That is, if $x = x_1 x_2 \dots x_n$ and $y = y_1 y_2 \dots y_n$ then H(x, y) is the number of positions i for which $x_i \neq y_i$.

A S set of binary strings of length n is called a (n, d)-Hamming set if the Hamming distance between any pair of strings in S is at least d.

(a) Give an example of a maximum-size Hamming set, with n = 3 and d = 2.

(b) Following is a greedy algorithm for generating a (n, d)-Hamming set. Give an example for n = 3 and d = 2 of an ordering of the list M (containing 8 strings) such that the Hamming set returned by this greedy algorithm is *not* of maximum size.

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Hamming-Set(n,d) {returns a (n,d)-Hamming set}
initialize M to be a list of the 2^n binary strings
of length n (in arbitrary order)
initialize S to be the empty set
repeat
let s be the first string in M
remove s from M and add s to the set S
for each string s' in M do
    if H(s,s') < d then remove s' from M
until M is empty
return S</pre>
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(c) For the algorithm of part (b), show that in the worst case $O(2^n m)$ operations are done (where the test H(s,s') < d counts as one operation).

4. I recently found the following variant of the greedy algorithm of the previous problem in a 1997 U.S. patent (patent number 5,604,097, "Methods for Sorting Polynucleotides Using Oligonucleotide Tags" by S. Brenner):

Describe an input (n, d) and an initial list M on which this variant outputs a set which is *not* a (n, d)-Hamming set?