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| <b>CS/Math 473 ♦ Fall 2022</b><br><b>Midterm I Solutions Problem 1</b> | Name: |
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Suppose we are given a sorted array  $A[1..n]$  of  $n$  distinct integers, which are not necessarily positive.

- (a) Describe a fast algorithm that either computes an index  $i$  such that  $A[i] = i$  or correctly reports that no such index exists.
- (b) Suppose we know in advance that  $A[1] > 0$ . Describe an even faster algorithm that either computes an index  $i$  such that  $A[i] = i$  or correctly reports that no such index exists.

**Solution (part (a)):** We can solve the problem in  $O(\log n)$  time using a variant of (or a reduction to) binary search. Here are two pseudocode descriptions of the algorithm, one recursive and one iterative.

⟨⟨Find index  $j$  such that  $i \leq j \leq k$  and  $A[j] = j$ ⟩⟩

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FINDINDEX( $i, k$ ):
  if  $i > k$ 
    return NONE
   $j \leftarrow \lceil (i + k) / 2 \rceil$ 
  if  $A[j] = j$ 
    return  $j$ 
  else if  $A[j] > j$ 
    return FINDINDEX( $i, j - 1$ )
  else
    return FINDINDEX( $j + 1, k$ )

```

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FINDINDEX( $A[1..n]$ ):
   $lo \leftarrow 1$ ;  $hi \leftarrow n$ 
  while  $lo \leq hi$ 
     $mid \leftarrow \lfloor (lo + hi) / 2 \rfloor$ 
    if  $A[mid] = mid$ 
      return  $mid$ 
    else if  $A[mid] > mid$ 
       $hi \leftarrow mid - 1$ 
    else
       $lo \leftarrow mid + 1$ 
  return NONE

```

The key observation is that because  $A$  is a sorted array of distinct integers, we have  $A[j] \geq A[i] + (j - i)$  for all indices  $i < j$ . In particular, if  $A[i] > i$ , then  $A[j] > j$  for all  $j > i$ . Equivalently, suppose we (implicitly) define a new array  $B[1..n]$  by setting  $B[i] = A[i] - i$  for all  $i$ . Then the elements of  $B$  are sorted in non-decreasing order (but they are not necessarily distinct), and we are looking for an index  $i$  such that  $B[i] = 0$ . ■

**Rubric:** 8 points max. For an explicit algorithm: 1 for binary search idea + 1 for base case + 4 for recursive cases + 2 for time analysis. -1 for each off-by-one error. -1 for returning TRUE/FALSE instead of index. -1 for stating running time as a recurrence without solving it. A reduction to binary search is worth full credit. Max 3 points for a  $\Theta(n)$ -time algorithm; max 2 points for anything slower; scale partial credit.

**Solution (part (b)):** If  $A[1] = 1$ , we can clearly return 1 immediately. On the other hand, if  $A[1] > 1$  then  $A[i] > i$  for all  $i$ , so we can return None immediately. So we can solve this problem in  $O(1)$  time! ■

**Rubric:** 2 points: 1 for algorithm + 1 for running time

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| <b>CS/Math 473 ♦ Fall 2022</b><br><b>Midterm 1 Solutions Problem 2</b> | Name: |
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Describe and analyze an algorithm to find a matching with maximum total weight, in a binary tree with weighted edges.

**Solution:** Let  $T$  be the input tree, and let  $r$  denote its root node. For any vertex  $v$ , we define two functions:

- $MWM(v)$  is the weight of a maximum weight matching in the subtree rooted at  $v$ .
- $MWM^*(v)$  is the weight of a maximum weight matching in the subtree rooted at  $v$ , where  $v$  is not incident to a matching edge.

We need to compute  $MWM(r)$ . These functions satisfy the following mutual recurrences. Here  $v.l$  and  $v.r$  denote the left and right children of  $v$ , respectively.

$$MWM(v) = \begin{cases} 0 & \text{if } v = \text{NULL} \\ 0 & \text{if } v \text{ is a leaf} \\ \max \begin{cases} w(v, v.l) + MWM^*(v.l) + MWM(v.r) & \text{if } v.l \neq \text{NULL} \\ w(v, v.r) + MWM(v.l) + MWM^*(v.r) & \text{if } v.r \neq \text{NULL} \\ MWM^*(v) & \text{otherwise} \end{cases} & \text{otherwise} \end{cases}$$

$$MWM^*(v) = \begin{cases} 0 & \text{if } v \text{ is a leaf} \\ MWM(v.l) + MWM(v.r) & \text{otherwise} \end{cases}$$

We can memoize these functions into two new fields at each node of  $T$ , and we can evaluate the functions in postorder.

The resulting dynamic programming algorithm runs in  $O(n)$  time, where  $n$  is the number of vertices in  $T$ . ■

**Rubric:** 10 points: standard dynamic programming rubric. These are not the only correct solution. -1 for assuming every interior node has both left and right children.

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| <b>CS/Math 473 ♦ Fall 2022</b><br><b>Midterm I Solutions Problem 3</b> | Name: |
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Suppose we are given two bit strings  $P[1..m]$  (the “pattern”) and  $T[1..n]$  (the “text”), where  $m \leq n$ . Describe and analyze an algorithm to find the minimum Hamming distance between  $P$  and a substring of  $T$  of length  $m$ . For full credit, your algorithm should run in  $O(n \log n)$  time.

**Solution (consider 0s and 1s separately):** For any integer  $0 \leq s \leq n - m$  (“shift”), let  $HD(s)$  denote the Hamming distance between  $P[1..m]$  and  $T[s + 1..s + m]$ ; we need to compute  $\min_s HD(s)$ .

Let us write  $HD(s) = m - Both1(s) - Both0(s)$ , where

- $Both1(s)$  is the number of indices  $i$  such that  $P[i] = T[i + s] = 1$ .
- $Both0(s)$  is the number of indices  $i$  such that  $P[i] = T[i + s] = 0$ .

More formally, we have

$$Both1(s) = \sum_{i=1}^m P[i] \cdot T[s + i] \quad Both0(s) = \sum_{i=1}^m (1 - P[i]) \cdot (1 - T[s + i])$$

We can evaluate  $Both1(s)$  for every index  $s$  using convolution as follows. Define two sequences  $p$  and  $t$  by setting  $p_i = P[m - i]$  and  $t_i = T[i]$  for each index  $i$ . Then for all  $s$  we have

$$Both1(s) = \sum_i p_{m-i} \cdot t_{s+i} = (p \star t)_{m+s}$$

We can construct the sequences  $p$  and  $t$  in  $O(n)$  time, and then compute their convolution in  $O(n \log n)$  time using fast Fourier transforms.

We can similarly express  $Both0(s)$  as a convolution by defining  $p'_i = 1 - P[m - i]$  and  $t'_i = 1 - T[i]$  for each index  $i$ . Then for all  $s$  we have

$$Both0(s) = \sum_i p'_{m-i} \cdot t'_{s+i} = (p' \star t')_{m+s}$$

We can construct the sequences  $p'$  and  $t'$  in  $O(n)$  time, and then compute their convolution in  $O(n \log n)$  time using fast Fourier transforms.

After computing both convolutions, we can compute  $\min_s (m - Both0(s) - Both1(s))$  in  $O(n)$  time. The entire algorithm runs in  **$O(n \log n)$  time.** ■

**Solution (powers of -1):** For any integer  $0 \leq s \leq n - m$  (“shift”), let  $HD(s)$  denote the Hamming distance between  $P[1..m]$  and  $T[s + 1..s + m]$ ; we need to compute  $\min_s HD(s)$ .

First we modify the arrays to make Hamming distance behave more like a vector dot-product. For any index  $i$ , define

$$P'[i] = \begin{cases} 1 & \text{if } P[i] = 1 \\ -1 & \text{if } P[i] = 0 \end{cases} \quad T'[i] = \begin{cases} 1 & \text{if } T[i] = 1 \\ -1 & \text{if } T[i] = 0 \end{cases}$$

Then for any shift  $0 \leq s \leq n - m$ , we have

$$\sum_{i=1}^m P'[i] \cdot T'[s+i] = \sum_{i=1}^m (1 - 2[P[i] \neq T[s+i]]) = m - 2 \cdot HD(s)$$

Now define two sequences  $p$  and  $t$  by setting  $p_i = P'[m-i]$  and  $t_i = T'[i]$  for each index  $i$ . Then for all  $s$  we have

$$\sum_{i=1}^m P'[i] \cdot T'[s+i] = \sum_i p_{m-i} \cdot t_{s+i} = (p \star t)_{m+s}$$

and thus  $HD(s) = (m - (p \star t)_{m+s})/2$ .

We can construct the sequences  $p$  and  $t$  in  $O(n)$  time, compute their convolution in  $O(n \log n)$  time using fast Fourier transforms, and finally compute  $\max_s (m - (p \star t)_{m+s})/2$  in  $O(n)$  time. The entire algorithm runs in  **$O(n \log n)$  time**. ■

**Solution (squared differences):** For any integer  $0 \leq s \leq n - m$  (“shift”), let  $HD(s)$  denote the Hamming distance between  $P[1..m]$  and  $T[s+1..s+m]$ ; we need to compute  $\min_s HD(s)$ .

We can express the Hamming distance  $HD(s)$  as follows:

$$\begin{aligned} HD(s) &= \sum_{i=1}^m (P[i] - T[i+s])^2 \\ &= \underbrace{\sum_{i=1}^m P[i]^2}_{SumP} - 2 \cdot \underbrace{\sum_{i=1}^m P[i] \cdot T[i+s]}_{Both1(s)} + \underbrace{\sum_{i=1}^s T[i+s]^2}_{SumT(s)} \end{aligned}$$

(We can remove the squaring because  $0^2 = 0$  and  $1^2 = 1$ !) We compute each of the terms  $SumP$ ,  $Both1(s)$ , and  $SumT(s)$  for all  $s$  as follows:

- We can compute  $SumP$  in  $O(m)$  time by brute force, once for all  $s$ .
- We can compute the third term  $SumT(s)$  for all  $s$  in  $O(n)$  time using the recurrence  $SumT(s) = SumT(s-1) - T[s] + T[s+m]$ .
- Finally, we compute  $Both1(s)$  for all  $s$  using convolution. Specifically, we define two sequences  $p$  and  $t$  by setting  $p_i = P[m-i]$  and  $t_i = T[i]$  for each index  $i$ . Then for all  $s$  we have

$$Both1(s) = \sum_{i=1}^m P[i] \cdot T[s+i] = \sum_i p_{m-i} \cdot t_{s+i} = (p \star t)_{m+s}$$

We can construct the sequences  $p$  and  $t$  in  $O(n)$  time, and then compute their convolution in  $O(n \log n)$  time using fast Fourier transforms.

Finally, we compute  $\max_s (SumP - 2 \cdot Both1(s) + SumT(s))$  in  $O(n)$  time by brute force. The entire algorithm runs in  **$O(n \log n)$  time**. ■

**Rubric:** 10 points. These are not the only correct solutions.

- 1 for using FFT/convolution at all
- 2 for correctly dealing with both 0s and 1s (separately considering 00 and 11 matches, separately considering 01 and 10 matches, mapping (0, 1) to (-1, 1), squared differences, etc.)
- 3 for correctly setting up convolutions (reversing either  $T$  or  $P$ )
- 3 for correctly reading the minimum Hamming distance from the convolution(s)
- 1 for time analysis (if the algorithm is mostly correct)

A correct algorithm that runs in  $O(mn)$  or  $O(mn \log n)$  time is worth at most 4 points.

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| <b>CS/Math 473 ♦ Fall 2022</b><br><b>Midterm I Solutions Problem 4</b> | Name: |
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Describe and analyze an algorithm to compute, given a sequence of integers separated by @ (average) signs, the **smallest** possible value the expression can take by adding parentheses.

**Solution:** Let  $A[1..n]$  be the input array. For any indices  $i \leq k$ , let  $MinAve(i, k)$  denote the smallest possible value that can be obtained from the interval  $A[i..k]$  by adding parentheses. We need to compute  $MinAve(1, n)$ . This function satisfies the following recurrence:

$$MinAve(i, k) = \begin{cases} A[i] & \text{if } i = k \\ \min \{ MinAve(i, j) @ MinAve(j + 1, k) \mid i \leq j < k \} & \text{otherwise} \end{cases}$$

The resulting dynamic programming algorithm runs in  $O(n^3)$  time. ■

**Rubric:** 10 points: standard dynamic programming rubric. This is not the only correct evaluation order. The memoization arrays don't appear to have the right structure for a monotonicity speedup via SMAWK. As far as I know, this is the fastest algorithm for this problem (up to logarithmic factors).

**Non-solution:** Consider the following greedy algorithm: Merge the adjacent pair of numbers with the largest average (breaking ties arbitrarily), replace them with their average, and recurse. For example:

$$\begin{array}{c}
 \underline{8 @ 6} @ 7 @ 5 @ 3 @ 0 @ 9 \\
 \underline{7 @ 7} @ 5 @ 3 @ 0 @ 9 \\
 \underline{7 @ 5} @ 3 @ 0 @ 9 \\
 \underline{6 @ 3} @ 0 @ 9 \\
 \underline{6 @ 3} @ 4.5 \\
 \underline{4.5 @ 4.5} \\
 4.5
 \end{array}$$

With the right data structures, this algorithm can be implemented to run in  $O(n \log n)$  time; the only real bottleneck is maintaining a priority queue of adjacent pairs.

Unfortunately, this greedy algorithm does **not** always compute the optimal expression. Consider the input  $2 @ 5 @ 0 @ 6$ . The greedy algorithm outputs  $(2 @ 5) @ (0 @ 6) = 3.25$ , but the optimal expression is  $2 @ (5 @ (0 @ 6)) = 3$ . ♣