# CMSC 27200 - Problem Set 4

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# Note: I am using the 3-day late pass for this assignment!

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# 1a

$$T(n) = T(\frac{n}{2}) + 2n, T(1) = 1$$
 
$$T(n) = (T(\frac{n}{4}) + 2(\frac{n}{2})) + 2n = T(\frac{n}{4}) + 2n + n$$
 
$$T(n) = (T(\frac{n}{8}) + 2(\frac{n}{4})) + 2n + n = T(\frac{n}{8}) + 2n + n + \frac{1}{2}n$$
 
$$T(n) = T(\frac{n}{2^k}) + 2n \sum_{j=0}^{k-1} (\frac{1}{2})^j$$
 We can simplify: 
$$\sum_{j=0}^{k-1} (\frac{1}{2})^j = \frac{1(1-(\frac{1}{2})^k)}{1-\frac{1}{2}} = 2(1-(\frac{1}{2})^k) = 2-2(\frac{1}{2})^k$$
 
$$T(n) = T(\frac{n}{2^k}) + 2n(2-2(\frac{1}{2})^k)$$
 
$$T(n) = T(\frac{n}{2^k}) + 4n(1-(\frac{1}{2})^k)$$
 Let  $k = \log_2 n$ . Then,  $2^k = n$  and  $\frac{n}{2^k} = 1$ . 
$$T(n) = T(1) + 4n(1-(\frac{1}{2})^{\log_2 n})$$
 
$$T(n) = T(1) + 4n(1-\frac{1}{n})$$
 
$$T(n) = 1 + 4n - 4$$
 
$$T(n) = 4n - 3$$
 Therefore,  $T(n)$  is  $O(n)$ .

# 1b

$$T(n) = 2T(\frac{n}{2}) + 3n^2, T(1) = 8$$

$$T(n) = 2(2T(\frac{n}{4}) + 3(\frac{n}{2})^2) + 3n^2 = 4T(\frac{n}{4}) + 3n^2 + \frac{3}{2}n^2$$

$$T(n) = 4(2T(\frac{n}{8}) + 3(\frac{n}{4})^4) + 3n^2 + \frac{3}{2}n^2 = 8T(\frac{n}{8}) + 3n^2 + \frac{3}{2}n^2 + \frac{3}{4}n^2$$

$$T(n) = 2^kT(\frac{n}{2^k}) + 3n^2\sum_{j=0}^{k-1}(\frac{1}{2})^j$$
We can simplify: 
$$\sum_{j=0}^{k-1}(\frac{1}{2})^j = \frac{1(1-(\frac{1}{2})^k)}{1-\frac{1}{2}} = 2(1-(\frac{1}{2})^k) = 2-2(\frac{1}{2})^k$$

$$T(n) = 2^kT(\frac{n}{2^k}) + 3n^2(2-2(\frac{1}{2})^k)$$

$$T(n) = 2^kT(\frac{n}{2^k}) + 6n^2(1-1(\frac{1}{2})^k)$$
Let  $k = \log_2 n$ . Then,  $2^k = n$  and  $\frac{n}{2^k} = 1$ .
$$T(n) = nT(1) + 6n^2(1-(\frac{1}{2})^{\log_2 n})$$

$$T(n) = 8n + 6n^2(1-\frac{1}{n})$$

$$T(n) = 8n + 6n^2 - 6n$$

$$T(n) = 6n^2 + 2n$$
Therefore,  $T(n)$  is  $O(n^2)$ .

# 1c

```
T(n) = 4T(\frac{n}{2}) + n^2 + 3, T(1) = 1
T(n) = 4(4T(\frac{n}{4}) + (\frac{n}{2})^2 + 3) + n^2 + 3 = 16T(\frac{n}{4}) + 2n^2 + 12 + 3
T(n) = 16(4T(\frac{n}{8}) + (\frac{n}{4})^2 + 3) + 2n^2 + 12 + 3 = 64T(\frac{n}{8}) + 3n^2 + 48 + 12 + 3
T(n) = 4^k T(\frac{n}{2^k}) + kn^2 + \sum_{j=0}^{k-1} 3 \cdot 4^j
We can simplify: \sum_{j=0}^{k-1} 3 \cdot 4^j = \frac{3(1-4^k)}{1-4} = 4^k - 1
T(n) = 4^k T(\frac{n}{2^k}) + kn^2 + (4^k - 1)
Let k = \log_2 n. Then, 2^k = n and \frac{n}{2^k} = 1. Also, n^2 = (2^k)^2 = (2^2)^k = 4^k.
T(n) = n^2 T(1) + n^2 \log_2 n + (4^{\log_2 n} - 1)
T(n) = n^2 + n^2 \log_2 n + (n^2 - 1)
 T(n) = n^2 \log_2 n + 2n^2 - 1
Therefore, T(n) is O(n^2 \log_2 n)
```

# 1d

$$T(n) = 10T(\frac{n}{2}) - 16T(\frac{n}{4}) + 6n + 7, T(1) = 1, T(2) = 9$$
  
Let  $10T(\frac{n}{2}) - 16T(\frac{n}{4})$  be the homogeneous part of the recurrence and  $6n + 7$  be the inhomogeneous part of the recurrence.

First, we will solve the homogeneous part of the recurrence.

```
T_0(n) = n^p = 10(\frac{n}{2})^p - 16(\frac{n}{4})^p
n^{p} = 10 \frac{n^{p}}{2^{p}} - 16 \frac{n^{p}}{4^{p}}
\frac{10}{2^{p}} - \frac{16}{(2^{p})^{2}} = 1
(2^{p})^{2} - 10(2^{p}) + 16 = 0
We can let x = 2^p.
x^2 - 10x + 16 = 0
(x-8)(x-2) = 0
x = 8, 2
This means 2^p = 8 so p = 3 and 2^p = 2 so p = 1.
```

Therefore, the homogeneous part of the recurrence is  $T_0(n) = c_1 n + c_2 n^3$ .

Now, we will solve the inhomogeneous part of the recurrence.

```
T_1(n) = c_3 n \log_2 n + c_4
c_3 n \log_2 n + c_4 = 10(c_3(\frac{n}{2})\log_2(\frac{n}{2}) + c_4) - 16(c_3(\frac{n}{4})\log_2(\frac{n}{4}) + c_4) + 6n + 7
c_3 n \log_2 n + c_4 = 5c_3 n \log_2(\frac{n}{2}) + 10c_4 - 4c_3 n \log_2(\frac{n}{4}) - 16c_4 + 6n + 7
c_3 n \log_2 n + c_4 = 5c_3 n(\log_2 n - 1) + 10c_4 - 4c_3 n(\log_2 n - 2) - 16c_4 + 6n + 7
c_3 n \log_2 n + c_4 = c_3 n \log_2 n + 3c_3 n - 6c_4 + 6n + 7
This means 0 = (3c_3 + 6)n so c_3 = -2 and -6c_4 + 7 = c_4 so c_4 = 1.
Therefore, the inhomogeneous part of the recurrence is T_1(n) = -2n \log_2 n + 1.
```

Now, we will use the initial conditions to solve for  $c_1$  and  $c_2$ .

```
T(n) = c_1 n + c_2 n^3 - 2n \log_2 n + 1
T(1) = 1, so c_1 + c_2 + 1 = 1, so c_1 = -c_2.
T(2) = 9, so 2c_1 + 8c^2 - 4\log_2 2 + 1 = 2c_1 - 8c_1 - 4 + 1 = -6c_1 - 3 = 9, so c_1 = -2 and c_2 = 2.
T(n) = -2n + 2n^3 - 2n\log_2 n + 1
Therefore, T(n) is O(n^3)
```

### 2

Note, we will use K to denote the index K in 0-based indexing and k to denote the kth element in the sorted merged array. For example, if k=5, then K=4, such that we find the 5th element, which appears at index 4 of the sorted merged array. Thus, K = k - 1 for the rest of this problem. In general, we will apply a binary search algorithm on both arrays to narrow down where index K is located.

### Algorithm

- Stored Data: Keep track of the left and right bounds of the A and B subarray.
- Initialization: Let leftA = 0 and rightA = n 1 be the bounds of the A subarray. Similarly, let leftB = 0 and rightB = m 1 be the bounds of the B subarray.
- Recursive Step:
  - Base Case: If rightA < leftA, return the (K leftA)th element in the B subarray. Similarly, if rightB < leftB, return the (K leftB)th element in the A subarray.
  - Otherwise, calculate the midpoint of the A and B subarray, midA and midB.
    - \* If A[midA] < B[midB]:</li>
      · If K ≤ midA + midB: set rightB = midB 1 (removes upper half of B subarray)
      · K > midA + midB: set leftA = midA + 1 (removes lower half of A subarray)
      \* If A[midA] > B[midB]:
      · K ≤ midA + midB: set rightA = midA 1 (removes upper half of A subarray)
      · K > midA + midB: set leftB = midB + 1 (removes lower half of B subarray)
  - Call the recursive function with the updated leftA, rightA, leftB, and rightB subarray bounds.

#### Pseudocode

```
def Kth(leftA, rightA, leftB, rightB):
    if leftA > rightA, do:
        ret B[K - leftA]
    if leftB > rightB, do:
        ret A[K - leftB]
    set midA = (leftA + rightA) // 2
    set midB = (leftB + rightB) // 2
    if A[midA] < B[midB], do:
        if K \leq midA + midB, do:
            ret Kth(leftA, rightA, leftB, midB - 1)
        else, do:
            ret Kth(midA + 1, rightA, leftB, rightB)
        if K \leq midA + midB, do:
            ret Kth(leftA, midA - 1, leftB, rightB)
        else, do:
            ret Kth(leftA, rightA, midB + 1, rightB)
```

# **Explanation**

At each step, we aim to decrease the size of one subarray by half. We determine the midpoint of each subarray, midA and midB. We make the following observations:

- midA is the number of elements that lie to the left of midA in subarray A
- midB is the number of elements that lie to the left of midB in subarray B
- T = midA + midB is the number of elements that lie to the left of midA in subarray A and midB in subarray B

We know that either  $K \leq T$  or K > T, and either A[midA] < B[midB] or A[midA] > B[midB], so we can outline 4 cases below:

- If A[midA] < B[midB]: This means A[midA] comes before B[midB] in the sorted merged array.
  - There are at least T+1 elements before B[midB] in the sorted merged array. This includes A[midA] since A[midA] < B[midB], midA number of elements before A[midA], and midB number of elements before B[midB], for a total of at least T+1 elements. This means there is index 0 to at least T before B[midB] in the sorted merged array.
  - There are at most T+1 elements before and including A[midA] in the sorted merged array. This includes A[midA], midA number of elements before A[midA], and midB number of elements before B[midB], for a total of at most T+1 elements. This means there is index 0 to at most T before and including A[midA] in the sorted merged array.
  - Based on these observations, we narrow our search for the following cases:
    - \*  $K \leq T$ : Since there is index 0 to at least T before B[midB] in the sorted merged array, we eliminate midB and everything after in subarray B by setting rightB = midB 1.
    - \* K > T: Since there is index 0 to at most T before and including A[midA] in the sorted merged array, we eliminate midA and everything before in subarray A by setting leftA = midA + 1.
- If A[midA] > B[midB]: This means B[midB] comes before A[midA] in the sorted merged array.
  - There are at least T+1 elements before A[midA] in the sorted merged array. This includes B[midB] since B[midB] < A[midA], midA number of elements before A[midA], and midB number of elements before B[midB], for a total of at least T+1 elements. This means there is index 0 to at least T before A[midA] in the sorted merged array.
  - There are at most T+1 elements before and including B[midB] in the sorted merged array. This includes B[midB], midA number of elements before A[midA], and midB number of elements before B[midB], for a total of at most T+1 elements. This means there is index 0 to at most T before and including B[midB] in the sorted merged array.
  - Based on these observations, we narrow our search for the following cases:
    - \*  $K \leq T$ : Since there is index 0 to at least T before A[midA] in the sorted merged array, we eliminate midA and everything after in subarray A by setting rightA = midA 1.
    - \* K > T: Since there is index 0 to at most T before and including B[midB] in the sorted merged array, we eliminate midB and everything before in subarray B by setting leftB = midB + 1.

Finally, we consider the termination case. If leftA > rightA, we it means we have exhausted all possibilities for the Kth element in array A. Therefore, it must lie in array B. In particular, we return the (K - leftA)th element in B since leftA represents the number of elements smaller than the overall Kth element. The same argument applies for leftB > rightB.

### Runtime

At each step, we are reducing our search space by half. So, for array A, there can be at most  $\log n$  recursive calls and similarly, for array B, there can be at most  $\log m$  recursive calls. This means that the total runtime is  $O(\log n + \log m)$ .

# 3

Suppose we have an array A of size n that stores  $x_1, ..., x_n$ , such that  $A[0] = x_1, ..., A[n-1] = x_n$ . Our algorithm is to calculate all possible pair sums and use a two pointer strategy to find 2 pair sums that add to our expected sum S.

### Algorithm

**Step 1**: Calculate all possible sums. Store this in *sums*.

- Stored Data: Store all possible sums in a list sums.
- ullet Initialization: Let sums be an empty list.

• Iterative Step: For each  $0 \le i < n$  and  $0 \le j < n$  such that i < j, add the object  $(s = x_i + x_j, i, j)$  to sums. After this, sort sums by s.

**Step 2**: Find all unique pair sums and maximum i and minimum j for each.

- Stored Data: Store the new sums in a list new\_sums.
- Initialization: Let new\_sums be an empty list.
- Iterative Step: For each sum in sums, find the maximum index i such that there exists a j where i < j and  $x_i + x_j = s$  and find the minimum index j such that there exists a i where i < j and  $x_i + x_j = s$ . Store the sum, maximum index i, and minimum index j as an object in  $new\_sums$ . This can be done by iterating through sums since duplicate pair sums are adjacent since sums is in sorted order.

**Step 3**: Use two pointers to find sums in  $new\_sums$  that add up to S.

- Stored Data: Keep track of left and right pointers of the new\_sums array.
- Initialization: Let left = 0 and  $right = len(new\_sums) 1$ .
- Iterative Step: If left > right, return False. Otherwise, calculate the current total sum  $current\_S$ .
  - If  $current_S < S$ , increment left.
  - If  $current\_S > S$ , decrement right.
  - If  $current_S = S$ :
    - \* If minimum j of left sum is less than maximum i of right sum OR if minimum j of right sum is less than maximum i of the left sum, return True.
    - \* Otherwise, increment left and decrement right.

#### Pseudocode

```
def does_sum_exist(A, S):
    n = len(A)
    sums = []
    for i in range(n):
        for j in range(i + 1, n):
            s = A[i] + A[j]
            sums.append((s, i, j))
    sums.sort()
    new_sums = []
    index = 0
    while index < len(sums):
        next_index = index + 1
        while next_index < len(sums) and sums[index][0] == sums[next_index][0]:
            new_s = sums[index][0]
            new_max_i = max(sums[index][1], sums[next_index][1])
            new_min_j = min(sums[index][2], sums[next_index][2])
            sums[index] = (new_s, new_max_i, new_min_j)
            next_index += 1
        new_sums.append(sums[index])
        index = next_index
    left, right = 0, len(new_sums) - 1
    while left <= right:</pre>
        current_S = new_sums[left][0] + new_sums[right][0]
        if current_S < S:
```

```
left += 1
if current_S > S:
    right -= 1
if current_S == S:
    if new_sums[left][2] < new_sums[right][1] or new_sums[right][2] < new_sums[left][1]:
        return True
    else:
        left += 1
        right -= 1</pre>
```

return False

# Explanation

The correctness of the algorithm is determined at step 3. At this point, we have all unique sums of 2 numbers and stored the maximum i and minimum j for which there exists another index to produce the sum. Given this, we reduce our problem to finding 2 sums of 2 numbers that add up to S.

Since  $new\_sums$  is in sorted order of sums, we can initialize left and right pointers at opposite ends of the array. From here, we can get the following conditions:

- $new\_sums[left] + new\_sums[right] < S$ : In this case, we have traversed all elements of  $new\_sums$  that lie after right, meaning to increase our current total sum, we must increment left because  $new\_sums[left+1] > new\_sums[left]$ .
- $new\_sums[left] + new\_sums[right] > S$ : In this case, we have traversed all elements of  $new\_sums$  that lie before left, meaning to decrease our current total sum, we must decrement right because  $new\_sums[right 1] < new\_sums[right]$ .
- $new\_sums[left] + new\_sums[right] = S$ : In this case, we know there exists 2 pairs of sums that add up to S. However, we have to verify if they are legitimate. Define the following:
  - Let  $s_1$  denote the first sum and  $s_2$  denote the second sum.
    - \* Let  $i_1$  be the maximum index i such that there exists some j where  $i_1 < j$  and  $x_{i_1} + x_j = s_1$ .
    - \* Let  $i_2$  be the maximum index i such that there exists some j where  $i_2 < j$  and  $x_{i_2} + x_j = s_2$ .
    - \* Let  $j_1$  be the minimum index j such that there exists some i where  $i < j_1$  and  $x_i + x_{j_1} = s_1$ .
    - \* Let  $j_2$  be the minimum index j such that there exists some i where  $i < j_2$  and  $x_i + x_{j_2} = s_2$ .
  - For this current solution to be legitimate, one of the following has to hold:
    - \*  $j_1 < i_2$ : In this case, there is some  $a < j_1$  where  $x_a + x_{j_1} = s_1$  and  $i_2 < b$  where  $x_{i_2} + x_b = s_2$ . Thus,  $a < j_1 < i_2 < b$  and  $x_a + x_{j_1} + x_{i_2} + x_b = s_1 + s_2 = S$ , so a valid solution exists.
    - \*  $j_2 < i_1$ : In this case, there is some  $a < j_2$  where  $x_a + x_{j_2} = s_2$  and  $i_1 < b$  where  $x_{i_1} + x_b = s_1$ . Thus,  $a < j_2 < i_1 < b$  where  $x_a + x_{j_2} + x_{i_1} + b = s_2 + s_1$ , so a valid solution exists.
  - In particular, we must check both of these conditions because whether each pair sum is produced before or after the other is arbitrary, as  $new\_sums$  is sorted by sum, not index. If none of these conditions hold, we increment left and decrement right. The reason we have to modify both left and right is because each pair sum in  $new\_sums$  is unique and  $new\_sums[left] + new\_sums[right] = S$ , so incrementing or decrementing only one of left or right cannot ever produce S since only one number is changing.

Finally, note that we terminate once left has passed right. The reason we can allow left and right to be equal is that can represent two sets of identical sums at four different indices. For example, suppose that left = right and S[left][1] = i and S[left][2] = j. This means there exists an a such that i < a and  $x_i + x_a = s$  and there exists a b such that b < j and  $x_b + x_j = s$ . If S[left][2] < S[left][1], this means that j < i, so b < j < i < a, so  $x_b + x_j + x_i + x_a = s + s = S$ .

### Runtime

- Step 1: We iterate through at most n elements in each loop since there a total of  $n^2$  possible pair sums. Then, we sort the  $n^2$  sums, which takes  $n^2 \log n$  time. Therefore, this step is  $O(n^2 \log n)$ .
- Step 2: We iterate through  $n^2$  sums in one loop. Although there is an internal loop, we update index to match the end of the internal loop, so each element in sums is explored once. Therefore, this step is  $O(n^2)$ .
- Step 3: Using two pointers for  $new\_sums$ , of which there are at most  $n^2$ , we hit each sum exactly once. Therefore, this step is O(n).

Combining all the steps, this algorithm takes  $O(n^2 \log n)$  time.

# 4

We will apply a divide and conquer algorithm to find all the visible lines. The idea is that we find the visible lines among lower sloped lines and visible lines among higher sloped lines. Them, we combine these solutions appropriately to get the visible lines among both these sets of lines.

# Algorithm

- Stored Data: Keep track of the visible lines with lower slopes lower\_visible\_lines and higher slopes higher\_visible\_lines. Note that due to the implementation of the recursive step, every line in lower\_visible\_lines will be visible with each other and in sorted order by slope. Similarly, every line in higher\_visible\_lines will be visible with each other and in sorted order by slope.
- Initialization: Sort all the lines by slope. For all lines with the same slope, remove the line with the smaller y-intercept until every line has a distinct slope. Let these set of lines be  $L = [L_1, ..., L_m]$ .
- Recursive Step:
  - If there are fewer than 3 lines, all must be visible. Thus, we return all the lines.
  - If there are exactly 3 lines, either 2 or 3 are visible. If the first and second line intersect before the first and third, then all three lines are visible and we return all the lines. Otherwise, we return the first and third lines.
  - If there are more than 3 lines, we split the lines into lower\_lines and higher\_lines based on slope. Then, we call visible\_lines for lower\_lines, which gives lower\_visible\_lines, and for higher\_lines, which gives higher\_visible\_lines. lower\_visible\_lines and higher\_visible\_lines are visible among each other and sorted by slope. We call combine\_visible\_lines with lower\_visible\_lines and higher\_visible\_lines, which returns the lines visible among all lines in lower\_visible\_lines and higher\_visible\_lines. It works as follows:
    - \* If there are fewer than 3 lines total in lower\_visible\_lines and higher\_visible\_lines, return the concatenation of lower\_visible\_lines and higher\_visible\_lines.
    - \* Otherwise, initialize all\_visible\_lines as lower\_visible\_lines. As we consider each line in higher\_visible\_lines, let the line under consideration be called next\_highest\_line. Consider the following for next\_highest\_line:
      - · If there are fewer than 2 lines in all\_visible\_lines, append next\_highest\_line to all\_visible\_lines.
      - · Otherwise, let the second most recent line added to all\_visible\_lines be called penult and most recent line added to all\_visible\_lines be called ult. Until penult and ult intersect before penult and next\_highest\_line, remove ult (and update penult and ult with second most recent and recent line added to all\_visible\_lines). Then, add next\_highest\_line to all\_visible\_lines.
    - \* Finally, return  $all\_visible\_lines.$

# Pseudocode

```
def delete_middle(first_line , second_line , third_line):
   if intersectX(first_line , second_line) > intersectX(first_line , third_line):
        return True
```

```
def visible_lines(L):
    if size(L) < 3, do:
        return L
    else if size (L) = 3, do:
        set first_line = L[0]
        set second_line = L[1]
        set third_line = L[2]
        if delete_middle(first_line, second_line, third_line), do:
            L. removes (second_line)
        return L
    else:
        set lower_lines, higher_lines = split_by_slope(L)
        set lower_visible_lines = visible_lines (lower_lines)
        set higher_visible_lines = visible_lines (higher_lines)
        return combine_visible_lines(lower_visible_lines, higher_visible_lines)
def combine_visible_lines(lower_visible_lines, higher_visible_lines):
    if size(lower_visible_lines) + size(higher_visible_lines) < 3, do:
        return concatenate (lower_visible_lines, higher_visible_lines)
    set all_visible_lines = lower_visible_lines
    for next_highest_line in higher_visible_line:
        if size (all_visible_lines) < 2, do:
            all_visible_lines.append(next_highest_line)
        else, do:
            while size(all_visible_lines) >= 2 and delete_middle(all_visible_lines
               [-2], all_visible_lines[-1], next_highest_line), do:
                all_visible_lines.pop()
            all_visible_lines.append(next_highest_line)
    return all_visible_lines
```

### **Explanation**

#### Eliminate duplicate slopes

First, we eliminate all lines of duplicate slopes because among multiple parallel lines, keeping only the one with the highest y-intercept since that can be seen over others. Assume from now on, all our lines have different slopes. Now, sort all the lines in order of slope.

#### Case: less than 3 lines

Consider the case of fewer than 3 lines, showing all lines are visible.

- 0 lines: trivial
- 1 line: visible throughout domain
- 2 lines: one line visible before intersection (single point because there are no parallel lines) and other line visible after intersection

#### Case: exactly 3 lines

Now, consider the base case of 3 lines,  $L_1$ ,  $L_2$ , and  $L_3$ . Assume these lines are sorted in order of slope.  $L_1$  (minimum slope) is always visible and  $L_3$  (maximum slope) is always visible, since their range can extend from a point to  $-\infty$  (minimum slope) or a point to  $\infty$  (maximum slope).  $L_2$  visible only if  $L_1$  and  $L_2$  intersect before  $L_1$  and  $L_3$ . This is because if  $L_1$  and  $L_2$  intersect after  $L_1$  and  $L_3$  intersect, then  $L_3$  will have taken over  $L_1$  after they intersect, and since  $L_3$  has a higher slope than  $L_2$ ,  $L_2$  will never get a chance to take over  $L_1$  since  $L_3$  has already taken over.

### Divide and conquer strategy

We use a divide and conquer algorithm. When we have at most 3 lines, we can find the visible lines using the procedure above. However, if there are more than 3 lines, we split them into two groups <code>lower\_lines</code> and <code>higher\_lines</code>, where <code>lower\_lines</code> is the set of lines with the lower half of slopes and <code>higher\_lines</code> is the set of lines with the upper half of slopes. Since the lines are sorted by slope, every line in <code>lower\_lines</code> has a smaller slope than every line in <code>higher\_lines</code>. Then, aim to find the visible lines for <code>lower\_lines</code> and <code>higher\_lines</code>, calling the results <code>lower\_visible\_lines</code> and <code>higher\_visible\_lines</code>, respectively.

At this point, we must combine the results lower\_visible\_lines and higher\_visible\_lines. Since lower\_visible\_lines are visible among each other, and higher\_visible\_lines are visible among each other, the correct solution to this problem involves finding all the lines among lower\_visible\_lines and higher\_visible\_lines that are visible altogether.

### Combining solutions to subproblems

We must combine subproblem solutions, lower\_visible\_lines and higher\_visible\_lines, to get the solution to the overall problem. Due to our base case, we return visible lines such that they are in order of slope. So, we know that lower\_visible\_lines and higher\_visible\_lines are in order of slope.

From here, we must find the final combination of lines between lower\_visible\_lines and higher\_visible\_lines. First, we can let all\_visible\_lines be lower\_visible\_lines, so the initial set of lines in all\_visible\_lines are all visible and in sorted order by slope. Now, as we consider each new line in higher\_visible\_lines, we need to make sure it is visible among all lines in all\_visible\_lines.

Call each new line in higher\_visible\_lines being considered next\_highest\_line, since at this point, it will have the highest slopes among all of all\_visible\_lines. We consider next\_highest\_line against the second most recent and most recent line added to all\_visible\_lines. Call the second most recent line penult and most recent line ult. The idea is that when adding next\_highest\_line, it will always be visible since it is the maximum sloped line, so we must remove other lines if necessary in all\_visible\_lines such that they can all be seen with next\_highest\_line. Furthermore, we know that if next\_highest\_line can be added to all\_visible\_lines with comparing to penult and ult, the updated all\_visible\_lines must be visible because the intersections of previous lines occurred before next\_highest\_line intersected with ult.

- Suppose these 3 lines can be visible together (checked via base case of 3 lines). This means next\_highest\_line takes over ult as the latest visible line (based on x coordinate intersection takeover) after ult takes over penult. Since all previous intersections occur before that of ult and next\_highest\_line, it means next\_highest\_line can be added to all\_visible\_lines.
- Suppose these 3 lines cannot be visible together (checked via base case of 3 lines). Since next\_highest\_line currently has the highest slope of lines considered, it must be visible. Therefore, ult must be removed because penult and ult intersect after penult and next\_highest\_line, so ult cannot be seen. We continuously remove the most recent line in all\_visible\_lines until we can safely add next\_highest\_line (meaning the second most recent and recent are visible with it, or there is less than 2 lines left in all\_visible\_lines).

#### Runtime

- Sorting the lines takes  $O(n \log n)$  time using mergesort.
- Removing the lines with duplicate slopes and preserving the one with highest y-intercept takes O(n) time since we can simply iterate through the sorted lines and only consider lines without a duplicate slope (as the duplicate is always present right after in the sorted lines).
- Our algorithm replicates mergesort. Since we are splitting the lines in half and combining the solutions, the recursion depth is  $\log_2 n$ . At each stage, we have to merge two sets of lines together. The time complexity of merging a set of lines A and B is O(len(A) + len(B)). Furthermore, we know that each stage, each line can only be considered in a single merge, meaning each line is considered once per stage, so there are O(n) considerations total for merging (across different merges in the same stage). Therefore, the recurrence relation is  $T(n) = 2T(\frac{n}{2}) + n$ , which has a time complexity of  $n \log n$ .

The overall time complexity of this algorithm is the number of recursions,  $\log n$ , by merging time complexity at each stage of recursion, n, so we get an  $O(n \log n)$  algorithm.