## Day 5: Adam Gasienica-Samek Contest 1 Tutorial

 $Adam_{-}GS$ 

1st OCPC, Winter 2023

February 22, 2023

Given two sequences  $a_0, a_1, \ldots, a_{n-1}$  and  $a_0, a_1, \ldots, a_{n-1}$ , define a sequence

$$c_0,c_1,\ldots,c_{n+m-2}$$
 as

$$c_k = \sum_{i+j=k} a_i b_j$$

Find the sum of c.

$$n, m \leq 10^5$$
.

#### Consider the polynomials:

- $A(x) = a_0 x^0 + a_1 x^1 + \dots + a_{n-1} x^{n-1}$
- $B(x) = b_0 x^0 + b_1 x^1 + \dots + b_{m-1} x^{m-1}$
- $C(x) = c_0 x^0 + c_1 x^1 + \dots + c_{n+m-2} x^{n+m-2}$

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$$C(x) = c_0 x^0 + c_1 x^1 + \dots + c_{n+m-2} x^{n+m-2}$$

Then  $C(x) = A(x)B(x) \Rightarrow$  You can solve this with FFT.

Easier solution:

$$\sum_{i=0}^{n+m-2} c_i = \left(\sum_{i=0}^{n-1} a_i\right) \left(\sum_{i=0}^{m-1} b_i\right)$$

Proof: Expand it, or use the fact that C(1) = A(1)B(1).

## G - Guessing by Divisibility

There is a secret positive integer n. You can ask questions "Is n divisible by x?". Find n using 1500 questions.  $n < 10^4$ .

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There are 1229 primes numbers below 10000. The sum of  $\lfloor \log_p 10^4 \rfloor$  over these primes is 1280.

There are n people. Each person either always tells the truth or always lie. There are more truth tellers than liars.

You can ask person a the question "How long is b's nose?". Whenever a person lie, their nose length increases. Determine all the liars with 10001 questions.  $n < 10^4$ .

Let's ask two people x and y about z in two consecutive queries. If they reply with the same number, then they are either both liars or both truth tellers.

Ask people  $2, 3, \ldots, n$  about person 1. There are two cases:

- ▶ If n = 2k, then there are at least k + 1 truth tellers. A strict majority of  $2, 3, \ldots, n$  must give the same answer.
- ▶ If n = 2k + 1, then there are also at least k + 1 truth tellers. At least half of 2, 3, ..., n must give the same answer.

Case 1: A majority of  $2, 3, \ldots, n$  answer the same.

Then the majority are all truth tellers, and the rest are liars. To check person 1, ask 1 about anyone and then ask any truth teller about 1 to check if the nose of 1 changed.

Case 2: No majority of  $2, 3, \ldots, n$  answer the same. Then exactly half of them answer the same.

If the other half doesn't answer the same, this is the same as case 1.

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If the other half doesn't answer the same, this is the same as case 1.

Otherwise, one half is truth tellers and the other is liars. Then person 1 must be a truth teller. We ask both 1 and 2 about 3 to check if 2 is a liar.

Both cases require n+1 questions.

The limit is set to 10001 so that an alternative solution which requires n+2 questions when n is odd can pass.

A tournament with n vertices is generated by randomly selecting each edge: from the smaller vertex to the larger with probability  $\frac{a}{b}$  and from larger to smaller otherwise. Find the probability that the generated graph is a DAG.  $n < 10^6$ .

There are exactly n! tournaments that are DAGs, and each of them corresponds to a permutation of  $1, 2, \ldots, n$ .

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$$p^{m-k}(1-p)^k$$

where m is the number of edges  $\frac{n(n-1)}{2}$ .

We want to find the sum over all permutations. Let c(n,k) denote the number of permutations with n elements and k inversions.

Then the answer to the problem is:

$$\sum_{k=0}^{m} c(n,k) p^{m-k} (1-p)^k$$

Let's find a recurrence for c(n, k).

$$c(1,0) = 1$$

$$c(n,k) = \sum_{i=0}^{n-1} c(n-1, k-i)$$

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Doing this directly takes  ${\cal O}(n^3)$  time.

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Therefore  $f_n(x) = 1(1+x)(1+x+x^2)\cdots(1+x+\cdots+x^{n-1}).$ 

We can rewrite the answer to

$$\sum_{i=0}^{m} c(n,i)p^{m-i}(1-p)^{i} = p^{m} \sum_{i=0}^{m} c(n,i) \left(\frac{1-p}{p}\right)^{i}$$

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Notice that the right side is just evaluating the polynomial  $f_n$  at  $\frac{1-p}{p}$ . We may calculate it in O(n).

Given a directed, weighted graph, find a walk with length at least k that has the highest possible median weight.

 $n \le 10^5$ ,  $m \le 2 \times 10^5$ ,  $k \le 50$ .

Binary search the answer by checking if the median can be at least x.

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Claim: We only have to check walks with length at most 2k.

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Let dp[x][l] be the maximal sum of values of a walk with length l ending at vertex x.

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For  $l=1,2,\dots,2k$  and each edge  $u\to v$ , we can update dp[v][l] with dp[u][l-1].

Time complexity:  $O((n+m)k \log C)$  where C is the maximum weight.

#### E - Erase the Primes

There is an array  $a_1, a_2, \ldots, a_n$  and there are q queries. For each query, you have to find the cost of the subarray  $a_l, a_{l+1}, \ldots, a_r$ .

The cost of an array is the minimum number of the operations needed to make every number divisible by two different primes ("good"), or -1 if it's impossible. An operation is replacing two numbers with their product.

 $n, q \leq 2 \times 10^5, \ a_i \leq 10^7.$ 

If every number is a power of the same prime number, then the cost is -1.

If every number is a power of the same prime number, then the cost is -1. Otherwise, we can multiply them all together so the cost is positive.

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We only have to consider prime powers.

In one operation, we can remove 1 or 2 prime powers. If two numbers are powers of different primes, we can remove them together with one operation. The strategy is to do operations on as many pairs of different primes as possible.

Let k be the number of prime powers in the range, and  $p = [p_1, p_2, \dots, p_k]$  be the primes.

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Claim: If no prime has majority in p, then we need  $\lceil \frac{k}{2} \rceil$  operations.

We can form  $\lfloor \frac{k}{2} \rfloor$  pairs of different primes by greedily picking the two primes with largest frequency in each operation.

Claim: If a prime has majority and it appears t times, then we need t operations.

Claim: If a prime has majority and it appears t times, then we need t operations. We can form at most k-t pairs of different primes. The remaining 2t-k majority elements have to be removed one by one.

The problem reduces to finding the frequency of the majority element in a range, which is a standard problem and can be done in  $O(\log n)$ ,  $O(\log^2 n)$  or  $O(\sqrt{n})$  time per query.

# J - Joining Arrays

Define the permutation scaling P(B) of  $[B_1,B_2,\ldots,B_m]$  as the permutation of  $[1,2,3,\ldots,m]$  such that  $P(B)_i < P(B)_j$  iff  $B_i \leq B_j$ . Given n sets  $S_1,S_2,\ldots,S_n$ , we make a sequence  $[A_1,A_2,\ldots,A_n]$  such that  $A_i \in S_i$ . Count the number of ways to choose A such that P(A) is lexicographically largest. P(A) = P(B) is P(A) = P(B).

# J - Joining Arrays

Let's find the lexicographically largest permutation scaling.

- 1. Choose the biggest element from the first set.
- 2. For each element we want to maximize the number of previous elements that are bigger.
- 3. If there are multiple values that maximize that, we take the biggest one.

# J - Joining Arrays

After knowing the maximal permutation scaling, we can reorder the sets in the order of the permutation.

Now the problem becomes counting the number of ways to choose A such that it is increasing. This can be done with DP using two pointers.

## I - Incomplete Information Queries

There is an unknown tournament graph with n vertices. The i-th vertex has  $c_i$  reachable vertices.

There are m bidirectional edges that you can add to the graph. Answer q queries: What is the minimum number of edge additions required to travel from a to b in the best/worst case?

 $n, m, q \le 5 \times 10^5.$ 

## I - Incomplete Information Queries

Contract all the SCCs and sort them in topological order. Notice that the result is uniquely determined by the sequence  $c_i$ .

Suppose we can add an edge between the u-th SCC and the v-th SCC. This operation makes every SCC between u and v strongly connected.

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Contract all the SCCs and sort them in topological order. Notice that the result is uniquely determined by the sequence  $c_i$ .

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The queries now become: Given a segment [l,r], find the minimum number of segments from a given set needed to cover [l,r].

The answer to each query can be found in  $O(\log n)$  time using binary lifting.

Consider a forest with n vertices. k special vertices are secretly chosen, and you are allowed to start a walk from vertex x with length m. Find the maximum probability that you visit a special vertex on the walk if you choose it optimally. You should process q queries online. Each query is either adding an edge to the forest or asking the answer for some x, k and m.

 $n \le 2 \times 10^5$ ,  $q \le 5 \times 10^5$ .

The optimal strategy is choosing a walk that visits as many different vertices as possible. Suppose a different vertices are visited. Then the answer is

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The queries become "How many different vertices can you visit with a walk of length m starting from x?".

Observation: The vertices visited must form a tree.

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Suppose we visit all vertices of a tree starting at vertex x and ending at y. The minimum number of edges required is equal to  $2 \times \#$ edges in the tree — the distance between x and y.

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Suppose we visit all vertices of a tree starting at vertex x and ending at y. The minimum number of edges required is equal to  $2 \times \#$ edges in the tree — the distance between x and y.

The problem reduces to finding the longest path starting from x.

Lemma: The farthest vertex from any vertex in a tree is the endpoint of a diameter.

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We want to create a data structure on a forest that supports:

- ▶ Adding an edge between two different trees.
- Maintaining a diameter of each tree.
- Querying the distance between two vertices.

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Instead of merging two trees, we support adding a leaf instead. Maintain the LCA binary lifting array and depth for each vertex.

#### Finding distance between two vertices:

- Find a' and b' such that a' is an ancestor of a, b' is an ancestor of b and  $\operatorname{depth}(a') = \operatorname{depth}(b') = \max(\operatorname{depth}(a), \operatorname{depth}(b))$
- ▶ Jump with both a' and b' at the same time to see if they reach the same ancestor at the same depth.
- ▶ Return depth(a) + depth(b) 2 depth(lca(a, b)).

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- ▶ Return depth(a) + depth(b) 2 depth(lca(a, b)).

Updating the depths and binary lifting array take O(1) and  $O(\log n)$  time.

Maintaining diameters:

Let s and t be current diameter endpoints and v be the added leaf. The diameter endpoints after the addition will be one of (s,t), (s,v) or (v,t). We can find the correct one by using the distance query.

We can use leaf addition to merge trees by adding vertices from the smaller tree to the larger tree one by one in DFS order.

The final time complexity will be  $O(n \log^2 n + q \log n)$ .

There are n numbers  $a_1, a_2, \ldots, a_n$ . Count the number of ways to choose a subsequence that has length divisible by k and has bitwise XOR 0.  $n < 10^6$ , k < 20,  $0 < a_i < 10^6$ .

Let's solve for k = 1.

The XOR convolution  $a \star b$  of sequences a and b is defined as

$$c_k = \sum_{i \oplus j = k} a_i \cdot b_j.$$

The Walsh-Hadamard transform (WHT) of a sequence a is defined as

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$$(a)_i = \sum_{j=0}^{n} (-1)^{\text{popent}(i \text{ AND } j)} \cdot a_j.$$

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Let  $A_i$  be a sequence  $A_{i,0},A_{i,1},\dots,A_{i,2^{20}-1}$  s.t.  $A_{i,0}=1$ ,  $A_{i,a_i}=1$  and  $A_{i,j}=0$  for every other j. The solution when k=1 is the 0-th element of

$$A_1 \star A_2 \star \cdots \star A_n$$
.

With WHT, the product is equal to

$$WHT^{-1}(WHT(A_1) \cdot WHT(A_2) \cdot \ldots \cdot WHT(A_n)).$$

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Since  $\operatorname{popent}(0\operatorname{AND} j) = 0$ , each element of  $\operatorname{WHT}(A_i)$  is either 0 or 2. Each element of  $\operatorname{WHT}(A_1) \cdot \operatorname{WHT}(A_2) \cdot \ldots \cdot \operatorname{WHT}(A_n)$  is either  $2^n$  or 0. We can check which one it is by doing SOS DP.

For a different k, we treat each  $A_{i,j}$  as a polynomial in x modulo  $x^k-1$ . Then  $A_{i,0}=1$  and  $A_{i,a_i}=x$ .

#### B - Beautiful XOR Problem

For a different k, we treat each  $A_{i,j}$  as a polynomial in x modulo  $x^k - 1$ . Then  $A_{i,0} = 1$  and  $A_{i,a_i} = x$ . Each element of  $\mathrm{WHT}(A_1) \cdot \mathrm{WHT}(A_2) \cdot \ldots \cdot \mathrm{WHT}(A_n)$  is equal to

$$(1+x)^m \cdot (1-x)^{n-m}$$
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We can precompute  $(1+x)^m$  and  $(1-x)^m$  for all m in O(nk) time.

For each value  $(1+x)^m \cdot (1-x)^{n-m}$  we only care about the coefficient of  $x^0$ , so we can multiply two polynomials in O(k)

so we can multiply two polynomials in O(k).

The total time complexity is O(nk).

Solve the following problem in an assembly-like language with only 4 32-bit registers:

Given n integers  $a_1, a_2, \ldots, a_n$ , find the product of the k largest primes in it modulo 2023.

 $n \le 100$ ,  $a_i \le 500$ ,  $k \le 4$ 

We can store three numbers in one register in base 1000, i.e., we store (x,y,z) as  $x+y\times 10^3+z\times 10^6$ .

Let  $a \geq b \geq c \geq d$  be the four currently biggest prime numbers in the sequence. Initially each of them is equal to 1.

The initial registers setup will look like this:

- ightharpoonup A: (a,b,n)
- $\triangleright$  B: (c, d, 0)
- ightharpoonup C: (0,0,0)
- ightharpoonup D: (0,0,0)

#### Our algorithm looks like this:

- 1. Repeat n times:
- Read a number and check if it is prime.
- 3. If it is prime, update the list of biggest primes.
- 4. Find their product modulo 2023.

If A stores (a, b, n), we can make a loop that runs n times.

- ▶ DECREASE A BY 10<sup>6</sup>
- ▶ IF  $A \ge 10^6$  GOTO loop ELSE GOTO break

We can make a prime checking procedure.

- 1. Read input to C.
- 2. Set *D* to 2.
- 3. While D < C:
- 4. If D divides C, exit the procedure.
- 5. Increase D by 1.
- 6. Continue to the next step where  ${\cal C}$  is prime.

To check whether D divides C:

- 1. Set a new variable e to 0.
- 2. While C > D:
- 3. Decrease C by D.
- 4. Increase e by 1.
- 5. D divides C iff C = D.

We can store e in the most significant digit of B.

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After the procedure, we need to restore C:

- 1. While e > 0:
- 2. Increase C by D.
- 3. Decrease e by 1.

Suppose we found a prime number p and want to update the values of a,b,c,d.

- ▶ If  $p \ge d$ , swap p, d
- ▶ If  $d \ge c$ , swap d, c
- ▶ If  $c \ge b$ , swap c, b
- ▶ If  $b \ge a$ , swap b, a

We can move a value d from B to D:

- 1. DECREASE B BY  $10^3$
- 2. INCREASE D BY 1
- 3. IF  $B \geq 10^3$  GOTO 1 ELSE GOTO 4
- 4. ...

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- 2. INCREASE D BY 1
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- 4. ...

The result becomes

- ightharpoonup A: (a,b,n)
- ightharpoonup B: (c, 0, 0)
- ightharpoonup C: (0,0,0)
- ightharpoonup D: (d, 0, 0)

The most significant digits of other registers can be moved similarly — they behave like stacks.

We may use  ${\cal C}$  and  ${\cal D}$  as temporary values to swap variables.

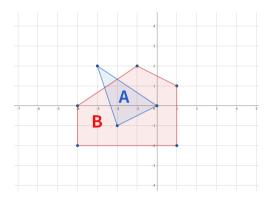
There is a convex polygon A and k convex polygons  $B_1, B_2, \ldots, B_k$ . Initially A intersects every of  $B_i$ .

Find the minimum value of |dx| + |dy| such that after you move A by (dx, dy), A intersects none of  $B_i$ .

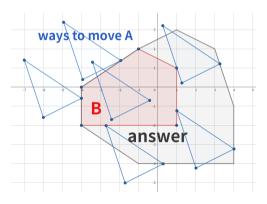
Total #vertices of  $B_i \leq 75000$ , #vertices of  $A \leq 75000$ ,  $k \leq 25000$ .

Consider k = 1.

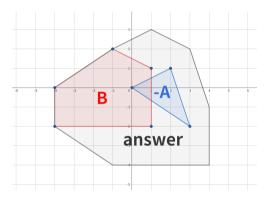
A and B are convex polygons. What is the set of vectors (dx,dy) such that A+(dx,dy) intersects  $B\mbox{\it ?}$ 



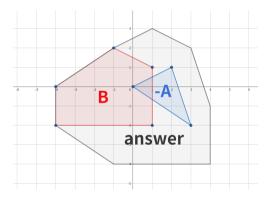
Intuitively, it should look like this:



Theorem: It is the Minkowski sum of -A and B.



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The sum is  $\{b-a \mid a \in A, b \in B\}$ . A vector d=(dx,dy) is in the set if and only if a+d=b for some points  $a \in A$  and  $b \in B$ .

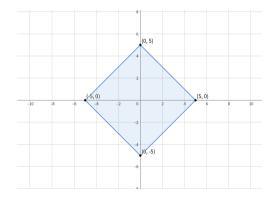
#### Rewrite the problem:

There is a convex polygon -A and k convex polygons  $B_1, B_2, \ldots, B_k$ . Find the minimum value of |x| + |y| of a point (x, y) that is not inside the Minkowski sum of A and any  $B_i$ . The point (0, 0) is contained in all these sums.

Try binary search: For a value c, we want to check whether the region  $|x|+|y|\leq c$  contains a point outside all the sums.

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Fact: The region  $|x| + |y| \le c$  is a rotated square centered at (0,0).



Claim: We only have to check the boundary of the square.

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Reason: If a point is inside one of the sums, every point between it and (0,0) is also inside that sum, since the sum is a convex set and (0,0) belongs to it.

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Reason: If a point is inside one of the sums, every point between it and (0,0) is also inside that sum, since the sum is a convex set and (0,0) belongs to it. Therefore, if the boundary doesn't contain a point outside the sums, then

neither does its interior.

Let's make life easier:

- ▶ Check one edge at a time (e.g., assuming  $dx, dy \ge 0$ ).
- ▶ Rotate the problem by 45 degrees, so the squares become axis-aligned.

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- ▶ Check one edge at a time (e.g., assuming  $dx, dy \ge 0$ ).
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Now we're interested about an edge (-c,-c)-(c,-c) of the square.

Suppose we want to check whether the segment (-c,-c)-(c,-c) satisfies the constraint.

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Each Minkowski sum is a convex polygon, which covers a contiguous interval on the segment. It is sufficient to find the intersections of each sum with the line y=-c and check whether the entire segment is covered.

Recall how the Minkowski sum of two convex polygons is calculated:

- ► Find the lowest point of each polygon. Their sum is the lowest point of the result.
- ▶ Merge the lists of edge vectors of both polygons sorted by angle. These are the edges of the result.

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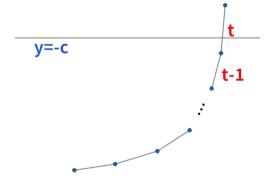
Doing this naively for every  $B_i$  takes

 $O(n+m_1)+O(n+m_2)+\cdots+O(n+m_k)=O(kn+\sum m_i)$  time which is too slow.

Observation: To find an intersection, we don't need the entire Minkowski sum. We only need the edge that intersects y=-c.

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Sort the edges by angle. We want the minimum t such that the sum of the first t edges have  $y \geq -c$ .



The edges of the i-th Minkowski sum contain the edges of A and  $B_i$ . We have two sorted list of edges, and we want to find a target prefix sum of their union.

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We could use binary search again to find the slope of the target edge, but doing this directly uses two additional  $\log$  factors.

To optimize, let's use a data structure that supports:

- ▶ Maintaining a list of vectors sorted by angle.
- Adding and deleting vectors.
- ightharpoonup Querying the smallest prefix of vectors with total  $y \ge$  some value.

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A segment tree or Fenwick tree can do each operation in  $O(\log n)$  time.

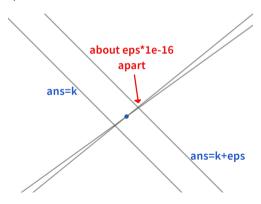
Alternative solution: For each i, binary search on the number of edges in A below y=-c. Then, we can find the exact number of edges in  $B_i$  below it with another binary search. It's easy to check whether we took too many or too less edges from A.

This requires trickier implementation details and a lot of edge case handling but also uses only one  $\log$  factor.

Caveat: Using floating points probably doesn't pass.

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Reason: Coordinates can be up to  $10^8$ , and the angle between two lines can be about  $10^{-16}$ . The answer must have a relative error below  $10^{-8}$ , so about  $10^{-24}$  precision is required.



The total time complexity:

$$O(N \log N \log C)$$

where  $N = n + \sum m_i$  and C is the coordinate range.

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Fun fact: The original (easier) version is the validator for this problem, and its intended time complexity is  $O(N \log N)$ .