# Polish Collegiate Programming Contest 2021 Editorial 

Jagiellonian University

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# Problem H Hidden password 

Author: Krzysztof Maziarz

We are given string $H_{1}$ and we know that there exists string $H_{2}: H_{2} \neq H_{1}$ and integer $d: 0<d<26$ such that Caesar cipher with shift $d$ transforms $H_{1}$ into $H_{2}$ and $H_{2}$ into $H_{1}$.

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We see that after double encryption $H_{1}$ transforms into itself, and because $H_{1} \neq H_{2}$ it follows that $d$ must be equal to 13 . This information is sufficient for obtaining $H_{2}$ as it is just $H_{1}$ encrypted with shift 13 Caesar ciper.

# Problem D Divided mechanism 

Author: Daniel Goc

## Task

We are given two rectilinear shapes on the plane and a sequence of instructions telling, that we should move one of them either up, down, left or right until it hits the other shape. We need to decide, if during the process we were able to move that part arbitrarily away from the other shape.

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To simplify the implementation, we can store the cells of the stationary part in the input array, and keep the cells of the moving part, for example, in a list.

# Problem L Lemurs 

Author: Daniel Goc

## Task

On a $n \times m$ map, there are some cells inhabited by lemurs. Each lair has a foraging area - every cell not further than $k$ cells in taxicab metric from it. Given cells, in which lemurs are foraging, we have to decide, if there exists specification of lairs location corresponding to that foraging area.


Let's note, that if lemurs are not foraging somewhere, then any lair cannot be located closer or with distance equal to $k$ from that cell. Let's cross out all such cells from the map, for example using BFS.

Can we place lemur lairs on the remaining cells, so that every marked cell is reachable from them? Anyhow we place them, the foraging area generated by them won't be "too large", it could only not span all the needed cells.

Therefore, we can wlog put lairs at every non-crossed out cells - such lair won't be ever inconsistent with our data. We place lairs everywhere, where it is possible and examine their total range - again, we can use BFS for that part, which gives linear solution to the whole problem.

# Problem K Kitten and Roomba 

Author: Krzysztof Maziarz

## Task

We are given a tree of size $n$ and a sequence $a_{1}, a_{2}, \ldots, a_{m}$ of vertices that Roomba will visit (in that order). Initially, a kitten is sleeping in vertex c. When Roomba enters vertex with a cat, it will wake up and escape to a randomly chosen, neighbouring vertex. Calculate the expected number of times the kitten will be woken up by the robot.

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Limits: $n \leq 1000000, m \leq 5000000$.

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When Roomba enters vertex $v$, we add $p[v]$ to the result, we update probabilities for all $d_{v}$ neighbours by adding $\frac{p[v]}{d_{v}}$, and finally, we set $p[v]=0$.

If we implement it naively, then the operation of adding to $v$ 's neighbours will take $O\left(d_{v}\right)$ time, which is too much (the whole algorithm may take $\mathcal{O}(n m)$ time $)$.

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To optimize it, for each vertex $v$ we will keep two values: $p[v]$ and lazy[ $v]$, where lazy $[v]$ is the value that we want to lazily add to $v$ 's children. To calculate the probability that the kitten is in the vertex $u$ we will check value of $p[u]+$ lazy $[$ parent $[u]]$.

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Complexity: $\mathcal{O}(n+m)$.

# Problem J Jungle Trail 

Author: Krzysztof Maziarz

## Task

We are given a $n \times m$ grid. Each square is either empty, blocked (impassable) or contains a den of snakes, either poisonous or benign (not poisonous).

We can tap a column/row. In that case all poisonous snakes in this column/row are turned to benign, and vice versa.

We have to tap some of the columns/rows, so that we can get from top-left corner, to the bottom-right one, moving only down or right, and visiting only empty squares or squares with benign snakes.

If there is no path from the top-left corner, to the bottom-right one that goes through non-blocked squares, then the answer is obviously "NO ". If there is such a path, we will show that the answer is always "YES".

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Key observation: in every step the path visits a new row or a new column.

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Key observation: in every step the path visits a new row or a new column.
If we visit a square that contains poisonous snakes we can tap either the current column or row, so the current square turns into a den of benign snakes, and the squares that we have already visited remain untouched.

# Problem C Cake 

Author: Krzysztof Maziarz

## Task

We are given two arrays $2 \times n$ of integers. We want to transform the second one into the first one by preforming the operation: rotate a $2 \times 2$ square by 180 degrees. Compute the minimum number of operations required or decide that it is impossible.

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Limits: $n \leq 500000$.

Rotation by 180 degrees corresponds to swapping $t[0][i]$ with $t[1][i+1]$ and $t[1][i]$ with $t[0][i+1]$ for some $i$.

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Let's swap $t[0][i]$ and $t[1][i]$ for even $i$ (in both arrays). After that change a rotation by 180 degrees corresponds to just swapping two consecutive columns.

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The problem simplifies to: given a sequence of pairs, transform it into another sequence by swapping consecutive elements.

Let's enumerate pairs in the target sequence with the numbers $1,2, \ldots, n$ (we use all numbers even if there are duplicates in the sequence).

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If we enumerate pairs in the origin sequence with the numbers corresponding to them in the target sequence (be careful with duplicates), then the answer is number of inversions in a sequence (it is well known problem).

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If we enumerate pairs in the origin sequence with the numbers corresponding to them in the target sequence (be careful with duplicates), then the answer is number of inversions in a sequence (it is well known problem).

We can count the inversions using merge sort, segment tree or indexed set. Complexity $\mathcal{O}(n \log n)$.

# Problem F Fence 

Author: Krzysztof Maziarz

## Task

We are given a sequence $a_{1}, a_{2}, \ldots, a_{n}$. For a fixed integer $b$ we split all the $a_{i}$ 's into $\left\lfloor\frac{a_{i}-1}{b}\right\rfloor$ copies of a number $b$ and $\left(\left(a_{i}-1\right) \bmod b\right)+1$.

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Example: For $a=[4,5,6]$ and $b=2$ we get $[2,2,2,2,1,2,2,2]$.

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Example: For $a=[4,5,6]$ and $b=2$ we get $[2,2,2,2,1,2,2,2]$.
For every $b$ compute the sum of elements with odd indices in the resulting sequence.
Limits: $\sum a_{i} \leq 10^{6}$.

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[5, 4, 3, 2, 1]
Number of elements in the sequence $\rightarrow 5$.
Sum of elements with odd indices $\rightarrow 9$.
Sum of elements with even indices $\rightarrow 6$.

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Iterating over $b=1 \ldots$ we need to update only leaves with corresponding $a_{i} \geq b$. The total number of updates equals $\sum_{i=1}^{n} a_{i}$ which yields the $O\left(\left(\sum_{i=1}^{n} a_{i}\right) \cdot \log (n)\right)$ complexity.

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Bonus: solve it in $O\left(\sum_{i=1}^{n} a_{i}\right)$.

# Problem I Interesting numbers 

Author: Team work

## Statement

A sequence of integers ( $a_{1}, a_{2}, \ldots a_{n}$ ) and a number $k$ are given. Let's consider a graph in which vertices $i$ and $j$ are connected iff $a_{i} \oplus a_{j} \leq k$. We are asked to find the size of a maximum clique in this graph.

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How to solve the problem for a fixed $k$ and a sequence of integers $a_{i}$, where $a_{i} \in\left[0,2^{g}\right)$ ?

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Case $k<2^{g-1}$. If $a_{i}$ and $a_{j}$ differ on the $(g-1)$-th bit, then there is no edge between them, so we can independently solve the problem for the ranges: $\left[0,2^{g-1}\right)$ and $\left[2^{g-1}, 2^{g}\right)$.

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Case $k \geq 2^{g-1}$. If $a_{i}$ and $a_{j}$ don't differ on the $(g-1)$-th bit, then there exists an edge between them, so an edge may not exist only if $a_{i}<2^{g-1}$ and $a_{j} \geq 2^{g-1}$ (i.e. $a_{i}$ and $a_{j}$ differ on the $(g-1)$-th bit).

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Another approach is to notice that now we are looking for the maximum independent set in a bipartite graph.

Kőnig theorem: Size of a maximum independent set equals $|V|$ minus maximum matching in bipartite graphs.

In order to find the matching we can either modify matching algorithm to make it work efficiently for this specific bipartite graph...

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or use a segment tree to compress the graph and then find the maximum flow in the resulting graph using any reasonable flow algorithm (i.e. Dinic).

# Problem A AMPPZ in the times of disease 

Author: Krzysztof Maziarz

## Task

Partition $n$ points on the plane into $k$ (non-empty) groups, such that the longest distance between points inside the same group is less than the shortest distance between points from different groups.

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Limits: $n \leq 2000000, k \leq 20$.

## Solution 1:

Take any point $x_{1}$, wlog add it to the group 1 .

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In general: if we have assigned points $x_{1}, \ldots, x_{i}$ into groups (respectively) $1, \ldots, i$ so far, then $x_{i+1}$, which maximizes minimal distance to any of $x_{1}, \ldots, x_{i}$ has to go into a new group.

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At the end, we assign each of the remaining $n-k$ points into a group represented by the nearest from $x_{1}, \ldots, x_{k}$.

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Naive implementation achieves $\mathcal{O}\left(n k^{2}\right)$ complexity, but it can be easily sped up to $\mathcal{O}(n k)$.

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The complexity of a naive implementation is $\mathcal{O}\left(n k^{2}\right)$, using std::sets we can improve the theoretical complexity to $\mathcal{O}(n k \log k)$ (in practice, it's really slow).

# Problem G Gebyte's Grind 

Author: Krzysztof Maziarz

## Task

Consider a witcher with health points $H$ and three kinds of objects:

- beast, which decreases hp by $b_{i}$ (and kills if $H \leq b_{i}$ );
- inn, which kills if $H<k_{i}$, and sets $H$ to $k_{i}$ otherwise; and
- witch, that sets $H=\max \left(H, c_{i}\right)$.

The witcher's trail is a sequence of $n$ objects. We have to handle $q$ operations:

- changes: one object changes into another
- queries: we start at the position $l_{i}$ with hp $H_{0}$, and we wonder to which $r_{i} \geq l_{i}$ we are able to get, without dying
Limits: $n \leq 2000000, q \leq 4000000$.

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We can assemble functions for single objects easily, a sequence of many objects is also a function, however, more complicated...
...but can it be very complicated?

Turns out that an arbitrary sequence of objects after composing gives a function having following representation:

$$
f(x)= \begin{cases}0 & \text { for } x \in[0, a] \\ y & \text { for } x \in[a+1, b] \\ x-b+y & \text { for } x \in[b+1,+\infty]\end{cases}
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Proof: every base object is a special case of such function and a composition of such functions returns a function in that form too.

We maintain a segment tree, in every node keeping composition of objects from the corresponding base segment as a function in the above form (three numbers $a, b, c$ ).

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Change operation translates to leaf update and recalculation of $\mathcal{O}(\log n)$ nodes. To answer a query, we start in a leaf corresponding to $l_{i}$, firstly climbing up, and later traversing down (also $\mathcal{O}(\log n)$ time).

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Final complexity: $\mathcal{O}((n+q) \log n)$.

# Problem E Epidemic 

Author: Krzysztof Kleiner

## Task

You are given $n$ people, every person might be infected or not. Then $k$ consecutive events of the following form occur:
(1) Group of people have a meeting - if any of them were infected, then everybody from this group becomes infected (and remain infected until the end of their lives).
(3) Some person is tested and receives a negative result.

- Some person is tested, receives a positive results and is put under quarantine.
- You receive a query: Is it possible to prove that people put under quarantine are the only ones infected?

You need to be able to answer all the queries online. Limits: $n \leq 500000, k \leq 1000000$.

## We can represent current state of our knowledge as a directed acyclic graph. Initially, it consists of $n$ isolated vertices, on the $i$-th of then we put a pawn corresponding to the $i$-th person.

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- When group of people meet we create a new "meeting vertex". We add edges to this vertex from vertices in which pawns of meeting's participants currently are. Now, we move these pawns to the new vertex. If any of the people in the meeting were in possibly_infected, then we need to put all of the participants in this set.

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- If some person receives positive test result, then we erase this person from possibly_infected and remove the corresponding pawn from the graph.
- If some person receives negative test result, then we need to update the state of our knowledge.

When person $p$ receives negative test result, then we traverse our graph with DFS, starting from a vertex $v$ in which $p$ 's pawn currently stands.

## Step of DFS(v) algorithm:

- We know that there is a healthy person who took part in the meeting $v$. Therefore, all participants of this meeting were healthy when they met. For every pawn that is still in $v$ we can deduce that corresponding person is healthy and erase them from possibly_infected ${ }^{1}$.
- We erase vertex $v$ from the graph.
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- We erase vertex $v$ from the graph.
- For every edge $u \rightarrow v$, we call $\operatorname{DFS}(u)$. We can do it because all participants of $v$ were healthy, so all people that they met must have been healthy too.
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## Step of DFS(v) algorithm:

- We know that there is a healthy person who took part in the meeting $v$. Therefore, all participants of this meeting were healthy when they met. For every pawn that is still in $v$ we can deduce that corresponding person is healthy and erase them from possibly_infected ${ }^{1}$.
- We erase vertex $v$ from the graph.
- For every edge $u \rightarrow v$, we call $\operatorname{DFS}(u)$. We can do it because all participants of $v$ were healthy, so all people that they met must have been healthy too.
- For every edge $v \rightarrow w$, we erase such edge from the graph and check if it was the last going into $w$ from a meeting with potentially infected people. If so, then we call DFS( $w$ ).
${ }^{1}$ The rest of the participants must have attended other meetings since $v$ took place, so we cannot be sure that they didn't get infected

When we receive a query, then we can search for an answer in possibly_infected.

If this set is empty, then nobody (but people under the quarantine) is infected and the epidemic has been contained.

## Complexity analysis

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- All DFS calls in all iterations take $\mathcal{O}(n+s+c)$ operations because every visited vertex (or edge) is permanently erased after processing it.
- Number of operations on possibly_infected set is linear with respect to the input size. Every person can be put to this set at most as many times as there were meetings that this person attended (plus one time during initialization). The same goes for the number of times one person can be erased from possibly_infected. Every operation on the set takes $\mathcal{O}(\log n)$ time.


# Problem B <br> Babushka and her pierogi 

Author: Daniel Goc

## Task

You are given a sequence of $n$ integers $a_{i}$ and an integer sequence $p_{i}$ of the same length. In both of these sequences elements are pairwise distinct and consist of the same numbers ( $p$ is permutation of $a$ ). You are also given number $C$.
In one move you can choose two indices $i, j$ and swap $a_{i}, a_{j}$ paying $\left|a_{i}-a_{j}\right|+C$.
Your task is to transform a into $p$ at the lowest possible cost.

We can get a lower bound on the cost:
$\frac{1}{2} \sum_{i=1}^{n}\left|a_{i}-p_{i}\right|+\left(n-n u m b e r \_o f\right.$ _permutation_cycles $) \cdot C$
The second part of the sum comes from the fact that after all operations we have $a=p$, so there are $n$ cycles and a single swap can add at most one cycle, therefore we need at least $n$ - number_of_permutation_cycles operations.

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It turns out that there exist an algorithm that solves this problem at exactly that cost.

We can solve the problem for each permutation cycle separately, so we will focus on a single cycle.

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Our goal is to find two elements in the cycle, such that after swapping them we have split the cycle into two cycles, and we are still able to achieve the optimal cost. First condition is true for every two distinct elements of the cycle. The second is true in a following situation: let $i, j$ be indices of cycle elements in the sequence, then:

$$
a_{j} \in\left[\min \left(a_{i}, p_{i}\right), \max \left(a_{i}, p_{i}\right)\right] \wedge a_{i} \in\left[\min \left(a_{j}, p_{j}\right), \max \left(a_{j}, p_{j}\right)\right]
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We choose element of the cycle such that corresponding element in $p$ is the largest possible. Then there always exists a different element of the cycle such that paired with our chosen element it fulfils the aforementioned condition (it is easy too see that otherwise it would be a cycle with a single element and we wouldn't need to split it).

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If so, then we can search for such an element in our cycle, split it, update sequence $a$ and proceed recursively with two smaller cycles.

Naive search through the whole cycle is too slow, but we can alternate between searching from the beginning and the end of the cycle, which assures that we will check number of elements that is proportional to the size of the smaller of two cycles obtained from the split. This gives $\mathcal{O}(s \log s)$ operations where $s$ is the size of the initial cycle.

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We need to keep the cycle in a structure that will allow us to quickly find the largest element and to add/erase elements. A standard set is a good option which gives us overall complexity of $\mathcal{O}\left(n \log ^{2} n\right)$. It is possible to achieve $\mathcal{O}(n \log n)$ using other structures, but it was not required.

# Problem M Median 

Author: Krzysztof Maziarz

## Task

We have an incorrect algorithm for computing the median of a sequence. It works in a following way: if the sequence has at most 2 elements then return the correct answer, otherwise split the sequence into 3 parts with equal length, recursively compute the answer for them and then return median of these 3 values.

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Given sequence of $n$ integers in range $[0, m-1]$, with $q$ unknown numbers, compute (modulo $10^{9}+7$ ) in how many ways can we choose the unknown values (from range $[0, m-1]$ ) in such a way that the algorithm returns the correct median of a sequence.

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Limits: $n \leq 3^{8}=6561, q \leq 30, m \leq 10^{9}$.

Let's fix $t \in[0, m-1]$. In how many ways can we choose the unknown values such that both the correct median and the one returned by the algorithm equals $t$ ?

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Let's replace the numbers smaller than $t$ with -1 , the equal ones with 0 and the bigger ones with 1 . Let's denote the number of -1 s and $1 s$ that replaced the unknown numbers by $x$ and $y$ respectively $(x, y \in[0, q])$. It is easy to verify if the correct median is 0 knowing $x, y$.

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We will compute in how many ways can we choose the unknown values (from set $\{-1,0,1\}$ ) in such a way that the algorithm returns 0 .

It can be done for all $(x, y)$ at once using dynamic programming on a tree of the recurrence. In every node we keep the result for every tuple: $(x, y, a)$ where $a \in\{-1,0,1\}$ is an answer for the node.

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Let $X=\left\{x_{1}<x_{2}<\ldots<x_{l}\right\}$ be the set of known values in the sequence .
Let $S=\left\{\left[0, x_{1}-1\right],\left[x_{1}, x_{1}\right],\left[x_{1}+1, x_{2}-1\right],\left[x_{2}, x_{2}\right], \ldots,\left[x_{l}+1, m-1\right]\right\}$, $|S| \in \mathcal{O}(n)$. If $p \in S$ the for each $t \in p$ the dynamic solution returns the exact same results.

One can prove that only $\mathcal{O}(q)$ of the ranges contributes to the final answer. Informal argument: the median of all values won't diverge much from the median of all known values.

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The only problem is that we have to multiply the results from dp by $\sum_{t=L}^{R} t^{x}(m-1-t)^{y}$ for many different $x, y, L, R$.

$$
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& \text { Let } P_{x, y}(t)=t^{x}(m-1-t)^{y} . \\
& \text { Let } Q_{x, y}(t)=\sum_{i=0}^{t} P_{x, y}(i) .
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Total complexity $\mathcal{O}\left(n q+q^{5}\right)$.

## Jury

# Lech Duraj <br> Krzysztof Maziarz <br> Krzysztof Kleiner <br> Daniel Goc Mateusz Radecki 

## Beta testers

Rafał Burczyński Marcin Briański Kamil Rajtar Witold Jarnicki Jan Tułowiecki Adam Szady Mateusz Radecki Kamil Dębowski Marek Sommer

Thank you!

