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Warsaw U Contest Editorial

November 14th, 2016

by Mikhail Tikhomirov (MIPT)

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A. Ar	rkanoid	1				

There are $k \ 1 \times 1$ square obstacles on a rectangular $n \times m$ field. A ball starts moving diagonally to axes from a certain point. The ball reflects from the walls. When the ball collides with an obstacle, the latter is destroyed and the ball reflects naturally. Determine the time when the last obstacle is destroyed.

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Outline: modelling with effective finding of the next obstacle to break.

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The hardest part is to find for a certain ball position and a set of obstacles (some of the initial ones could get destroyed) which obstacle is the next to be destroyed.

A. Arkanoid

Let us number all possible pairs of position and direction of the ball starting from a certain point.



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If one has the number k of the situation (position + direction), one can transform it to actual position and velocity coordinates. Indeed, consider time moment k. If the ball started from (0.5, 0) moving to the upper right, it must have made $\lfloor (0.5k + 0.5)/m \rfloor$ bounces from vertical sides and $\lfloor 0.5k/n \rfloor$ from horizontal sides. This information is enough to find the coordinates and velocity.

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To answer a converse question, that is, determine the number k from coordinates and velocity, first notice that the pair (x-coordinate, x-velocity) periodically repeats with period 4m; the same holds for y and period 4n. Thus, one has to solve a system of modular equations of sort:

$$k \equiv k_x \pmod{4m}, k \equiv k_y \pmod{4n},$$

where numbers k_x and k_y correspond to the position and direction.

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One way of solving a system of similar form is to use the Chinese remainder theorem.

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Suppose that we are currently in a situation with index k. The current direction can correspond to increasing or decreasing k over time. Now, finding the next collision can be done with certain lower/upper bound query to the data structure.

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The solution has $O(k \log k)$ complexity.

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B. Va	ari-dire	ctio	nal St	reets			

A vertex v of a directed graph is *good* if for every vertex u either u is reachable from v or v is reachable from u. Find the set of good vertices of a given digraph.

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Outline: condense the strongly connected components of the digraph, obtain a simple criterion for a DAG using topsort properties.

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Choose a topological ordering of the DAG arbitrarily. For simplicity we will identify vertices of the DAG with their indices in topsort.

A vertex v is good iff all vertices u > v are reachable from v, and v is reachable from any u < v. We will check the first condition for all vertices; the second one can be checked in a completely symmetrical way.



For a vertex u > v let $deg_+(v|u)$ denote the number of edges (w, u) with $w \ge v$. Let us call a vertex u a v-source if $deg_+(v|u) = 0$ (that is, u is a source in the part of the graph to the right of v including v).

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Let us iterate over all possible v from left to right, and maintain $deg_+(v|u)$ for all $u \ge v$ along with the number of vertices with $deg_+(v|u) = 0$.

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To move from v to v + 1 simply decrease in-degree of all vertices directly reachable from v.

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To check the symmetrical condition consider the reversed graph. A vertex v is good if doesn't have a v-source in both cases.

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Build all SCC's and the condensation of the digraph (compressed graph with vertices in SCC's and edges between different SCC's). Apply the DAG solution to the condensation, output all vertices in good SCC's.

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Both condensation construction and DAG case are solvable in O(n + m) time.

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Find a hamiltonian cycle in an cube graph with 2^n vertices that contains a given perfect matching.

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Outline: constructive solution that reduces to smaller problems.

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Outline: constructive solution that reduces to smaller problems.

For details read the enclosed solution.

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We are given an array of *n* integers. All subsequences of the array are ordered by sum of the elements; subsequences with equal sum are ordered lexicographically as sorted tuples of indices. Find *k*-th subsequence in this ordering. $n, k \leq 10^6$.

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Outline: a Dijkstra-like approach with enough optimizations.

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We have to take care not to add the same entry to the structure twice. One way of doing it is to only append the elements that go later in the sequence than the previous last element, hence there is a unique order of adding elements for each subsequence.

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Optimization 1: if the size of the structure is greater than k, we can remove the largest entry.

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Optimization 1: if the size of the structure is greater than k, we can remove the largest entry.

Optimization 2: sort the given numbers (don't forget to store their original indices), stop trying to append a number when the sum becomes too great to fit in the structure.



This approach might still take too long since we have a lot of options to append a number on each step.

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This approach might still take too long since we have a lot of options to append a number on each step.

Optimization 3: leave only O(1) transitions from each state by introducting new information. Instead of (sum, subset of indices, [possibly lower bound for index]), we will now have (lower bound for sum after adding the number *i*, subset of indices, current position *i*).

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While processing the next state we have to try to add number i and add the subset to the list of generated sequences. However, in the sequel we may opt to skip the number. The two transitions are to continue with the number a_i included or not included in the subset; in both cases the current position is increased by 1. Note that the lower bound for the sum does not decrease in any case.

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It is easy to see that the target sequence will not have more than $\log_2 k$ elements since all the subsets of a subsequence precede it in the order. Thus the described solution has complexity $O(n \log n + k \log^2 k)$, since we compare two states in $O(\log_2 k)$ time.

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We are given a connected graph without multiple edges. Determine if all simple closed paths in the graph have the same length. If that is the case, count these cycles modulo $10^9 + 7$.

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Outline: if the condition holds, the biconnected blocks have very special form.

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If the component is a cycle of length I, then we have 2I closed paths for every choice of starting point and direction.

Otherwise, suppose that all simple cycles contain I edges each. Consider a pair C_1 , C_2 of intersecting cycles, and take their symmetrical difference S (that is, the set of edges that are present in exactly one of the cycles).

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The set S has less than 2/ edges, and can be decomposed into simple cycles. By assumption, S must itself be a cycle of length *I*. If that is the case, the intersection of C_1 and C_2 must be a path of length I/2. Thus $C_1 \cup C_2$ is a graph that consists of three edge-disjoint paths of length I/2 between a pair of vertices v and u.

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Note that no additional edges can be added between vertices of $C_1 \cup C_2$ so that each cycle has length *I*.

If $C_1 \cup C_2$ is not the whole component yet, there must be a cycle that has common edges with C_1 or C_2 , but doesn't lie completely inside $C_1 \cup C_2$. By a similar argument, the new cycle must consist of two paths of length I/2 between v and u: one old and one new.

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Finally, we can decompose the graph into biconnected components in O(n + m) time. We should also check that the cycle lengths for different components are equal.

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Count the number of ways to remove kd Nim heaps out of n so that the second player wins. $d \leq 10$, total size of the heaps $\leq 10^7$.

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Outline: standard DP with optimization.

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Fact

The second player wins iff XOR of all heap sizes is zero.



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Note that XOR of heap sizes a_i is less than $2 \max a_i$.

An $O(nd \max a_i)$ solution: $dp_{i,m,x} =$ number of subsets among first *i* heaps such that the number of omitted heaps is *m* modulo *d*, and XOR of all taken heaps' sizes if *x*.

This DP has $O(nd \max a_i)$ states and transitions.

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This DP has $O(nd \max a_i)$ states and transitions.

Optimization: let us process a_i by increasing. By the time we process a_i , we can't get XOR of some smaller numbers greater than $2a_i$, so we won't store such values. Now appending a single number a_i is done in $O(a_i)$, for the total complexity $O(d \sum a_i + n \log n)$.

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In a given tree find a simple path such that the number of edges with exactly one endpoint inside the path is maximized.

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Outline: standard subtree DP.

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First we solve a "vertical" version of the problem. Make the tree rooted, and let down(v) be the answer if the path goes from v into its subtree, and we only consider the edges in the subtree (no edge from v to the parent). down(v) will allow for a single-vertex path (unlike the original problem).

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If ch_v is the number of children of v, then

$$down(v) = \max\left(ch(v), ch(v) - 1 + \max_{u \text{ is a child of } v} down(u)\right).$$



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If v is the root of the tree, then the maximal length of such path is either

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$$\mathit{ch}(v) - 1 + \max_{u ext{ is a child of } v} \mathit{down}(u)$$

if v is an endpoint,

or

$$ch(v) - 2 + \max_{u_1, u_2 - \text{ different children of } v} (down(u_1) + down(u_2))$$

if v is a proper LCA of endpoints.



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All the above can be computed in O(n) time.

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In a directed graph, count the number of paths of length l from v to u that don't pass through v and u other than at start and finish. Many queries of v, u, l.

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In a directed graph, count the number of paths of length l from v to u that don't pass through v and u other than at start and finish. Many queries of v, u, l.

Outline: count the standard DP $paths_{v,u,l}$ for the number of all paths without any constraints, then carefully subtract all excess paths.

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We will have to count several DP's. First is the standard $paths_{v,u,l}$ for the total number of paths of length *l* from *v* to *u*:

$$paths_{v,v,0} = 1$$
 $paths_{v,u,l} = \sum_{w:(v,w) - \text{ an edge}} paths_{w,u,l-1}$

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 $paths'_{v,u,l}$ — the number of paths of length *l* from *v* to *u* that contain *v* only as the start. We want to subtract all non-suitable paths from total. Let *l'* be the last moment a bad path passes through *v*. Hence the formula:

$$paths'_{v,u,l} = paths_{v,u,l} - \sum_{l'=1}^{l-1} paths_{v,v,l'} paths'_{v,u,l-l'}$$

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 $cycle'_{v,u,l}$ — the number of paths of length *l* from *v* to *v* avoiding *u*. Let *l'* be the last moment a bad path passes through *u*. Then

$$cycle'_{v,u,l} = dp_{v,v,l} - \sum_{l'=1}^{l-1} dp_{v,u,l'} paths'_{u,v,l-l'}$$

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Finally, $paths''_{v,u,l}$ is the answer to the original problem. In a similar way we have

$$paths_{v,u,l}'' = paths_{v,u,l}' - \sum_{l'=1}^{l-1} paths_{v,u,l'}' cycle_{u,v,l-l'}'$$

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All the above values can be computed in $O(nml + n^2l^2)$ time, and each query can be answered in O(1) time.

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We are given a permutation. We repeatedly go from the current permutation to lexicographically previous one using minimal number of element swaps (not necessarily adjacent!). How many swaps we will make in total before we arrive to the $(1, \ldots, n)$ permutation?

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We are given a permutation. We repeatedly go from the current permutation to lexicographically previous one using minimal number of element swaps (not necessarily adjacent!). How many swaps we will make in total before we arrive to the (1, ..., n) permutation?

Outline: combinatorial argument, then "number-by-permutation"-like algorithm with RSQ.

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Let f(n) be the total number of swaps to proceed from $(1, \ldots, n)$ to $(n, \ldots, 1)$.

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Let f(n) be the total number of swaps to proceed from $(1, \ldots, n)$ to $(n, \ldots, 1)$.

To do that, first we have to get from $(1, \ldots, n)$ to $(1, n, \ldots, 2)$. Next we have to change $(1, n, \ldots, 2)$ into $(2, 1, 3, \ldots, n)$, then to $(2, n, \ldots, 3, 1)$, and so on.

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In general, we have n steps of "swap the suffix" sort. Each of them take f(n-1) steps.

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In general, we have *n* steps of "swap the suffix" sort. Each of them take f(n-1) steps.

Between these steps we have to change $(k, n, \ldots, k+1, k-1, \ldots, 1)$ to $(k+1, 1, \ldots, k, k+2, \ldots, n)$, where $k = 1, \ldots, n-1$.

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Fact

The minimal number of swaps to change a permutation p into permutation q is equal to n - (number of cycles in $p^{-1}q$).

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One can check with some case analysis (or with brute-force for small numbers) that the number of swaps needed to change $(k, n, \ldots, k+1, k-1, \ldots, 1)$ into $(k+1, 1, \ldots, k, k+2, \ldots, n)$ doesn't depend on k and is equal to $\lceil n/2 \rceil$.

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Thus, we have $f(n) = nf(n-1) + (n-1)\lceil n/2\rceil$. Values of this recurrence can readily be found in O(n) time.

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Let x_l be the index of p_l among the remaining elements if we order them by increasing. To place p_l in *l*-th position we have to do $x_l - 1$ repetitions of "swap the suffix" and "apply next permutation so that *l*-th element increases".

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By previous arguments, we have to perform $(x_l - 1)(f(n - l) + \lceil (n - l)/2 \rceil)$ swaps. After that, p_l is in its place, and all the later elements are sorted, thus we reduce to a smaller problem.

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Values of x_l can be found using any kind of RSQ data structure in $O(n \log n)$.

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Two players are playing a game with n pairs of heaps of stones. Initially both heaps in *i*-th pair contain a_i stones. The first player can remove any number of stones from any heap. The second player must move several stones between heaps in some pair. The first player wants to remove all stones in minimal number of moves, while the second player wants to play as long as possible. Find out the number of moves in the game if both play optimally.

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Outline: evil problem with hard-to-identify cases.

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Couple of simple observations:

• The first player always empties one of the heaps.

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Couple of simple observations:

- The first player always empties one of the heaps.
- After the first player emptied a heap, the second player should try to even out the heaps in this pair when this is possible.

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We will say that the first player can *snatch* a move if he forces the second player to move in a situation when each pair of heaps is either empty or has $(2^k, 2^k)$ stones for certain integer k (probably different for different pairs).

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These are the only situations that help the first player to get under the upper bound F. Indeed, if the second player could skip moves, then F would be the exact number of moves. Thus, the only way to do better than F is to force the second player to do harmful moves.

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Note that if some pair contains unequal heaps, than the second player can effectively skip a move, thus such situations are not appealing to the first player.

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Suppose that the first player will move in a pair that currently contains (x, x) stones. Forcing a second player's move here will result in (x - 1, 0) instead of (x, 0) (if the second player could skip). The move will be harmful if $\lfloor \log_2(x - 1) \rfloor < \lfloor \log_2 x \rfloor$, or, equivalently, $x = 2^k$.

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We want to make sizes of all pairs as close to (1, 1) as possible without making unequal pairs. By making enough moves, a pair (x, x) can be forced to a pair $(2^k - 1, 2^k - 1)$, where k is the number of leading ones in binary representation of x. Moving further in this pair will result in unequal heaps. We suppose that at all times the heaps are reduced to this form.

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Observation

Both the first and the second player will choose a pair with maximal k.

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Similarly, in the end the second player wants to have as few (1,1) pairs as possible, so he'll avoid making them from (3,3). Similarly, to avoid making (3,3) he'll avoid moving in (7,7), and so on.

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Finally, we can model the game as follows. After reducing all heaps to $(2^k - 1, 2^k - 1)$ form we'll store the number of pairs with $k = 0, 1, \dots, \log_2 A$ (where A is the maximal value of a heap size).

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If you have trouble understanding this solution, try to work out why the answer for 3, 3, 3 input is 15 instead of 17.

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Find longest common subsequence of two sequences *a* and *b* that consists of pairs of equal numbers. You can perform O(nm) operations, but can't store $\Omega(nm)$ memory.

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Outline: optimize the memory with additional bookkeeping.

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The standard DP solution for LCS can be modified as follows. Let $dp_{i,j}$ be equal to the maximal length of LCS of first *i* and *j* elements of *a* and *b* respectively if we are forced to take both elements of each pair simultaneously.

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Let prev(a, i, c) be the last occurrence of c in a before position i. Then $dp_{i,j} = \max(dp_{i-1,j}, dp_{i,j-1})$ if $a_i \neq b_j$, and $\max(dp_{i-1,j}, dp_{i,j-1}, 2 + dp_{prev(a,i,c)-1, prev(b,j,c)-1})$ if $a_i = b_j = c$ (all prev's have to be defined, of course).

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Let us compute $dp_{i,j}$ row by row. We can't store the whole matrix, so we'll just have two last rows.

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To account for $dp_{prev(...),prev(...)}$ let us store

$$dp_j^{equal} = \max_{i|a_i=b_j} dp_{i-1,j-1},$$

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where *i* ranges over all processed rows.

Note that we can use $dp_{prev(b,j,c)}^{equal}$ in place of $dp_{prev(a,i,c)-1,prev(b,j,c)-1}$. This eliminates our need for $\Omega(nm)$ memory and requires only O(n+m) memory.