Primal Brackets

Input file:	`standard input`
Output file:	`standard output`
Time limit:	1 second
Memory limit:	256 megabytes

A sequence of integers is balanced if it satisfies the following rules:

- An empty sequence is balanced.
- If A and B are balanced sequences, then their concatenation AB is balanced.
- If A is a balanced sequence, then for any positive integer x, the sequence obtained by simultaneously adding $+x$ at the beginning of A and $-x$ at the end of A is balanced.

For example, the sequence $[1, 2, -2, -1]$ is balanced, but $[-1, 1, 2, -2]$ and $[1, 1]$ are not.

A balanced sequence of length $2k$ is called primal if the first k integers are positive and the last k integers are negative. For example, the sequence $[3, 2, 1, -1, -2, -3]$ is primal, but $[1, -1, 2, -2]$ is not.

For a balanced sequence b, let $f(b)$ be the minimum number of non-empty subsequences* into which b can be partitioned† into such that each subsequence is a primal sequence.

For example, for the balanced sequence $b = [1, 2, -2, 3, -3, -1]$, one possible partition into a minimal number of primal subsequences is: $[1, 2, -2, -1]$ and $[3, -3]$. Therefore, $f(b) = 2$.

You are given a balanced sequence $a_1, a_2, \ldots, a_{2n}$ where all elements are distinct, and q queries. In each query, you will given two integers l and r, and you need to determine $f(a[l \ldots r])$ where $a[l \ldots r]$ is the subarray‡ of a starting at index l and ending at index r. Additionally, it is guaranteed that this subarray is a balanced sequence.

*A sequence c is a subsequence of sequence b if c can be obtained from b by deleting some (possibly zero or all) elements.

†A partition of a sequence b is a way to divide it into one or more subsequences such that every element of b belongs to exactly one subsequence. In other words, each element must be used exactly once across all subsequences in the partition.

‡A sequence a is a subarray of a sequence b if a can be obtained from b by deletion of several (possibly, zero or all) elements from the beginning and several (possibly, zero or all) elements from the end.

Input

The first line contains an integer t ($1 \le t \le 10^4$) — the number of test cases.

The first line of each test case contains two integers n and q ($1 \le n,\ q \le 10^5$) — the length of the sequence a and the number of queries.

The second line contains $2n$ integers $a_1, a_2, \ldots, a_{2n}$ ($1 \le |a_i| \le n, a_i \neq 0$). **It is guaranteed that a is a balanced sequence and all elements of a are distinct.**

Each of the next q lines contains two integers l and r ($1 \le l < r \le 2n$) representing the indices of the subarray $a[l \ldots r]$. **It is guaranteed that $a[l \ldots r]$ is also a balanced sequence.**

It is guaranteed that the sum of n over all test cases does not exceed $5 \cdot 10^5$, and the sum of q over all test cases does not exceed $5 \cdot 10^5$.

Output

For each query, print a single integer — the minimum number of subsequences into which the specified subarray can be partitioned into such that each subsequence is a primal sequence.

Example

standard input	standard output
2 3 3 1 2 -2 -1 3 -3 1 6 5 6 1 4 1 1 1 -1 1 2	2 1 1 1