

Candidate Report: Anonymous

Test Name:

[Summary](#) [Timeline](#)

Test Score

62 out of 100 points

62%

Tasks in Test

GenomicRangeQuery
Submitted in: JavaScript

Time Spent ⓘ
1 min

Task Score
62%

TASKS DETAILS

| | | | | |
|--------|---|------------|-------------|-------------|
| MEDIUM | 1. GenomicRangeQuery | Task Score | Correctness | Performance |
| | Find the minimal nucleotide from a range of sequence DNA. | 62% | 100% | 0% |

Task description

Solution

A DNA sequence can be represented as a string consisting of the letters A, C, G and T, which correspond to the types of successive nucleotides in the sequence. Each nucleotide has an *impact factor*, which is an integer. Nucleotides of types A, C, G and T have impact factors of 1, 2, 3 and 4, respectively. You are going to answer several queries of the form: What is the minimal impact factor of nucleotides contained in a particular part of the given DNA sequence?

The DNA sequence is given as a non-empty string $S = S[0]S[1] \dots S[N-1]$ consisting of N characters. There are M queries, which are given in non-empty arrays P and Q , each consisting of M integers. The K -th query ($0 \leq K < M$) requires you to find the minimal impact factor of nucleotides contained in the DNA sequence between positions $P[K]$ and $Q[K]$ (inclusive).

For example, consider string $S = \text{CAGCCTA}$ and arrays P, Q such that:

P[0] = 2 Q[0] = 4

P[1] = 5 Q[1] = 5

P[2] = 0 Q[2] = 6

The answers to these $M = 3$ queries are as follows:

- The part of the DNA between positions 2 and 4 contains nucleotides G and C (twice), whose impact factors are 3 and 2 respectively, so the answer is 2.
- The part between positions 5 and 5 contains a single nucleotide T, whose impact factor is 4, so the answer is 4.
- The part between positions 0 and 6 (the whole string) contains all nucleotides, in particular nucleotide A whose impact factor is 1, so the answer is 1.

Write a function:

function solution(S, P, Q);

that, given a non-empty string S consisting of N characters and two non-empty arrays P and Q consisting of M integers, returns an array consisting of M integers specifying the consecutive answers to all queries.

Result array should be returned as an array of integers.

For example, given the string $S = \text{CAGCCTA}$ and arrays P, Q such that:

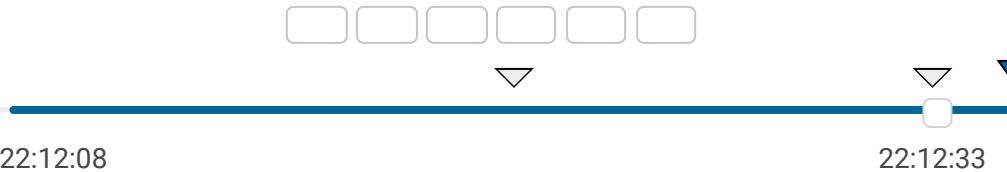
Programming language used: JavaScript

Total time used: 1 minutes

Effective time used: 1 minutes

Notes: *not defined yet*

Task timeline



Code: 22:12:32 UTC, js, final, score: 62 [show code in pop-up](#)

```
1  const ensureValid = (S, P, Q) => {
2    if (!S || !P || !Q || P.length !== Q.length) {
3      throw error("Data is invalid");
4    }
5  }
6
7  const getImpactFactor = (s) => {
8    if (s === 'A') {
9      return 1;
10   } else if (s === 'C') {
11     return 2;
12   } else if (s === 'G') {
13     return 3;
14   } else if (s === "T") {
15     return 4
16   } else {
17     return null;
18   }
```

```
P[0] = 2    Q[0] = 4
P[1] = 5    Q[1] = 5
P[2] = 0    Q[2] = 6
```

the function should return the values [2, 4, 1], as explained above.

Write an **efficient** algorithm for the following assumptions:

- N is an integer within the range [1..100,000];
- M is an integer within the range [1..50,000];
- each element of arrays P, Q is an integer within the range [0..N - 1];
- $P[K] \leq Q[K]$, where $0 \leq K < M$;
- string S consists only of upper-case English letters A, C, G, T.

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```
19 }
20
21 var solution = function solution(S, P, Q) {
22
23     const result = [];
24     const minImpactFactor = 1;
25
26     ensureValid(S, P, Q);
27
28     for (var i = 0; i < P.length; i++) {
29         var substr = S.substr(P[i], Q[i] + 1);
30
31         let impactFactor = null;
32
33         for (let j = 0; j < substr.length; j++) {
34             const element = substr.substr(j, j+1);
35
36             const currentImpactFactor = getImpactFactor(element);
37             if (currentImpactFactor === null) {
38                 throw error("Data is invalid");
39             }
40
41             if (impactFactor == null || impactFactor > currentImpactF
42                 impactFactor = currentImpactFactor;
43             }
44
45             if (impactFactor === minImpactFactor) {
46                 break;
47             }
48         }
49
50         result.push(impactFactor);
51     }
52
53     return result;
54 }
55 }
```

Analysis summary

The following issues have been detected: timeout errors.

Analysis ?

Detected time complexity: $O(N * M)$

| | | |
|---------------------------|------------------------------------|---|
| expand all | Example tests | |
| ▶ example | example test | ✓ OK |
| expand all | Correctness tests | |
| ▶ extreme_sinlge | single character string | ✓ OK |
| ▶ extreme_double | double character string | ✓ OK |
| ▶ simple | simple tests | ✓ OK |
| ▶ small_length_string | small length simple string | ✓ OK |
| ▶ small_random | small random string, length = ~300 | ✓ OK |
| collapse all | Performance tests | |
| ▼ almost_all_same_letters | GGGGGG..??..GGGGGG..??..GGGGGG | ✗ TIMEOUT ERROR Killed. Hard limit reached: 6.000 sec. |
| ----- | | |
| 1. | 6.000 s | TIMEOUT ERROR, Killed. Hard limit reached: 6.000 sec. |
| 2. | 0.084 s | OK |
| ▼ large_random | large random string, length | ✗ TIMEOUT ERROR Killed. Hard limit reached: 6.000 sec. |

| | | |
|----|---------------------------------|---|
| 1. | 6.000 s | TIMEOUT ERROR, Killed. Hard limit reached: 6.000 sec. |
| ▼ | extreme_large all max ranges | ✗ TIMEOUT ERROR Killed. Hard limit reached: 6.000 sec. |
| 1. | 6.000 s | TIMEOUT ERROR, Killed. Hard limit reached: 6.000 sec. |

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