

Test Name:

Summary      Timeline

Tasks summary

| Task                        | Time spent | Score |
|-----------------------------|------------|-------|
| GenomicRangeQuery<br>Python | 6 min      | 100%  |

Total score



Tasks Details

|        |   |            |             |             |
|--------|---|------------|-------------|-------------|
| Medium | 1. <b>GenomicRangeQuery</b>                               | Task Score | Correctness | Performance |
|        | Find the minimal nucleotide from a range of sequence DNA. | 100%       | 100%        | 100%        |

Task description

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:

```
def 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:

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

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  $P$  and  $Q$  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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Solution

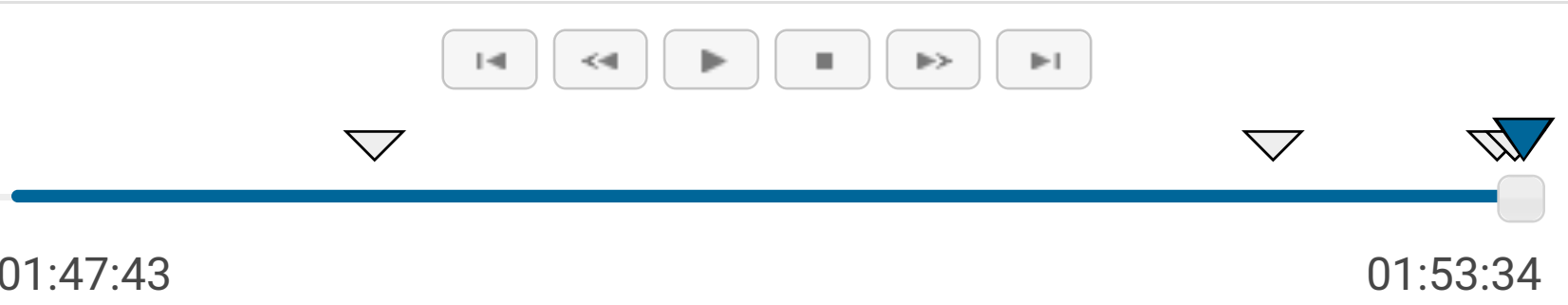
Programming language used: Python

Total time used: 6 minutes

Effective time used: 6 minutes

Notes: *not defined yet*

Task timeline



Code: 01:53:33 UTC, py, final, score: [show code in pop-up](#)  
100

```
1  # you can write to stdout for debugging purposes, e.g.
2  # print("this is a debug message")
3
4  def solution(S, P, Q):
5      # write your code in Python 3.6
6      M = len(P)
7      if M != len(Q):
8          return -1
9      N = len(S)
10     dict_ = {}
11     'A' : 1,
12     'C' : 2,
13     'G' : 3,
14     'T' : 4
15
16
17     arr=[]
18     for el in S:
19         arr.append(dict_[el])
20     global_minimum = min(arr)
21
22     if N == 1:
23         return ([arr[0]] * M)
24     else:
25         dict1 = {}
26         i = 0
27         for el in P:
28             if el not in dict1.keys():
29                 dict1[el] = []
30             dict1[el].append(i)
31             i += 1
32         dict2 = {}
33         i = 0
34         for el in Q:
35             if el not in dict2.keys():
36                 dict2[el] = []
37             dict2[el].append(i)
38             i += 1
39
40         min_ = max(arr)+1
41         min_before_reset = min_
42         dict_ = {}
43         queries = {}
44         for i in range(N):
45             min_ = min(min_, arr[i])
46
47             if i in dict1.keys() and i in dict2.keys():
48                 list_ = []
49                 for el1 in dict1[i]:
50                     for el2 in dict2[i]:
51                         if el1 == el2:
52                             queries[el1] = arr[i]
53                             list_.append(el1)
54
55                 list1 = []
56                 for el1 in dict1[i]:
57                     if el1 not in list_:
58                         list1.append(el1)
59                 dict1[i] = list1.copy()
60                 list2 = []
61                 for el2 in dict2[i]:
62                     if el2 not in list_:
63                         list2.append(el2)
64                 dict2[i] = list2.copy()
65
66             if i in dict2.keys():
67                 list_ = []
68                 for el in dict2[i]:
69                     if el in dict_.keys():
70                         dict_[el] = min(min_, dict_[el])
71                         queries[el] = dict_[el]
72                         list_.append(el)
73                 for el in list_:
74                     dict_.pop(el)
75
76             if i in dict1.keys():
77                 if min_before_reset != min_:
78                     for key in dict_.keys():
79                         dict_[key] = min(min_, dict_[key])
80                 for el in dict1[i]:
81                     if arr[i] == global_minimum:
82                         queries[el] = global_minimum
83                     else:
84                         dict_[el] = arr[i]
85                         min_before_reset = min_
86                         min_ = arr[i]
87
88             if min_ == global_minimum:
89                 for key in dict_.keys():
90                     queries[key] = global_minimum
91                 dict_ = {}.copy()
92
93     result = [ queries[i] for i in range(M)]
94     return result
```

Analysis summary

The solution obtained perfect score.

Analysis

|   |                   |
|---|-------------------|
| Detected time complexity: <b>O(N + M)</b> |                   |
| expand all                                | Example tests     |
| ▶ example                                 | ✓ OK              |
| example test                              |                   |
| expand all                                | Correctness tests |
| ▶ extreme_sinlge                          | ✓ OK              |
| single character string                   |                   |
| ▶ extreme_double                          | ✓ OK              |
| double character string                   |                   |
| ▶ simple                                  | ✓ OK              |
| simple tests                              |                   |
| ▶ small_length_string                     | ✓ OK              |
| small length simple string                |                   |
| ▶ small_random                            | ✓ OK              |
| small random string, length = ~300        |                   |
| expand all                                | Performance tests |
| ▶ almost_all_same_letters                 | ✓ OK              |
| GGGGGG.??..GGGGGG.??..GGGGGG              |                   |
| ▶ large_random                            | ✓ OK              |
| large random string, length               |                   |
| ▶ extreme_large                           | ✓ OK              |
| all max ranges                            |                   |