

Problem B. Bioengineering

Input file: *standard input*
Output file: *standard output*
Time limit: 33.2 seconds
Memory limit: 512 mebibytes

Viruses are usually bad for your health. How about fighting them with... other viruses? In this problem, you need to find out how to synthesize such good viruses.

We have prepared for you a set of strings of the letters 'A', 'G', 'T' and 'C'. They correspond to the DNA nucleotide sequences of viruses that we want to synthesize, using the following operations:

- Adding a nucleotide either to the beginning or the end of the existing sequence.
- Replicating the sequence, reversing the copied piece and gluing it with the original (so that e.g., "AGTC" becomes "AGTCCTGA").

We're concerned about efficiency, since we have very many such sequences, some of them very long. Find a way to synthesize them in a minimum number of operations.

Input

The first line of input contains the number of test cases T . The descriptions of the test cases follow:

Each test case consists of a single line containing a non-empty string. The string uses only the capital letters 'A', 'C', 'G' and 'T' and is not longer than 100 000 characters.

Output

For each test case, output a single line containing the minimum total number of operations necessary to construct the given sequence.

Example

standard input	standard output
4	3
AAAA	8
AGCTTGCA	6
AAGGGGAAGGGGAA	18
AAACAGTCCTGACAAAAAAAAAAAAAC	