## Frequent

An astrobiologist studies life on the planet Alphabet. Life here is DNA-based and there are 26 nucleotides. Consequently, the DNA of a life form from Alphabet can be represented as a string of lowercase letters of the Latin alphabet. The astrobiologist has sequenced the DNA of  $\mathbf{K}$  life forms, not necessarily distinct, with a total length of  $\mathbf{N}$  nucleotides. Now she would like to find strands (substrings) of DNA that occur frequently among these life forms. Let  $\mathbf{L}(\mathbf{i})$  be the length of the longest strand of consecutive DNA nucleotides common to at least  $\mathbf{i}$  life forms, for  $2 \le \mathbf{i} \le \mathbf{K}$ . Note that  $\mathbf{L}(\mathbf{i})$  can be 0.

#### Task

Help the astrobiologist compute the array **L**.

#### Input data

The file **frequent.in** contains an integer number on the first line, **K**, representing the number of life forms. Each of the following **K** lines contains a non-empty string of lowercase letters, terminated by a newline character.

#### **Output data**

The file **frequent.out** must contain K - 1 lines with the values L(2), L(3), ..., L(K), each on its own line.

#### **Limits and constraints**

- $2 \le N \le 200,000$
- $2 \le \mathbf{K} \le \mathbf{N}$
- Time limit: 0.5 seconds
- Memory limit: 128 MB

#### **Subtasks**

Test cases will be scored **individually**.

Subtask	Percentage of test cases	Additional input constraints
1	30%	$N \le 10,000$
2	40%	$N \le 100,000$
3	30%	none



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### Example

frequent.in	frequent.out	Explanation
6	5	atter appears in two of the strings
matter	3	mat appears in three of the strings
animate	2	<i>ma</i> (or <i>at</i> or <i>te</i> ) appear in four of the strings
pattern	2	<i>ma</i> appears in five of the strings
thermal	1	<i>a</i> appears in all the strings
domain		
teammate		