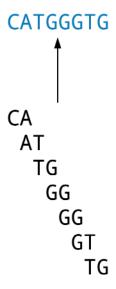
3B Reconstruct a String from its Genome Path

String Spelled by a Genome Path Problem

Find the string spelled by a genome path.

Input: A sequence of k-mers $Pattern_1$, ..., $Pattern_n$ such that the last k-1 symbols of $Pattern_i$ are equal to the first k-1 symbols of $Pattern_i+1$ for i from 1 to n-1.

Output: A string *Text* of length k + n - 1 where the *i*-th *k*-mer in *Text* is equal to *Pattern*_i for all *i* from 1 to *n*.



Formatting

Input: A space-separated list of strings *Patterns*.

Output: A string *Text*.

Constraints

- The number of patterns in the string-set *Pattern* will be between 1 and 10⁴.
- The length of any one pattern in *Pattern* will be between 1 and 10^2 .

Test Cases 🖸

Case 1

Description: The sample dataset is not actually run on your code.

Input:

ACCGA CCGAA CGAAG GAAGC AAGCT

Output:

ACCGAAGCT

Case 2

Description: The sample dataset is not actually run on your code.

Input:

CTT TTT TTG

Output:

CTTTG

Case 3

Description: The sample dataset is not actually run on your code.

Input:

TT TG GT TT

Output:

TTGTT

Case 4

Description: The sample dataset is not actually run on your code.

Input:

AGCAGATC GCAGATCA CAGATCAT AGATCATC GATCATCG ATCATCGG

Output:

AGCAGATCATCGG

Case 5

Description: A larger dataset of the same size as that provided by the randomized autograder. Check input/output folders for this dataset.