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The Virtual Learning Environment for Computer Programming

### Words 1

Nucleic acid sequences are labeled over the alphabet  $\{A, C, G, T\}$ , and there are  $4^n$  possible genomic sequences of length *n*. Amino acid sequences, on the other hand, are labeled over the alphabet  $\{A, C, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, Y\}$ , and there are  $20^n$  possible proteomic sequences of length *n*. An interesting problem is the generation of all the genomic sequences with *n* nucleotides or all the proteomic sequences with *n* amino acids, that is, the generation of all the words of length *n* over an alphabet  $\Sigma$ .

Write code for the words problem. The program must implement and use the WORDS function in the pseudocode discussed in class, which is iterative and is not allowed to perform input/output operations. Make one submission with Python code and another submission with C++ code.

#### Input

The input is an integer n and an alphabet  $\Sigma$ .

#### Output

The output is a sorted list of all the words of length *n* over the alphabet  $\Sigma$ .

Sample input 1	Sample output 1
1	А
G T A C	С
	G
	Т
Sample input 2	Sample output 2
2	AA
G T A C	AC
	AG
	AT
	CA
	CC
	CG
	CT
	GA
	GC
	GG
	GT
	TC
	IG
	11
Sample input 3	Sample output 3
3	AAA
G T A C	AAC
	AAG
	AAT

	1
	GAG
ACA	GAT
ACC	GCA
ACG	GCC
ACT	GCG
AGA	GCT
AGC	GGA
AGG	GGC
AGT	GGG
ATA	GGT
ATC	GTA
ATG	GTC
ATT	GTG
CAA	GTT
CAC	TAA
CAG	TAC
CAT	TAG
CCA	TAT
ССС	TCA
CCG	TCC
ССТ	TCG
CGA	TCT
CGC	TGA
CGG	TGC
CGT	TGG
CTA	TGT
CTC	TTA
CTG	TTC
CTT	TTG
GAA	TTT
GAC	

## **Problem information**

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