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**Words 2****X86108\_en**

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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 recursive 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**

```
1
G T A C
```

**Sample output 1**

```
A
C
G
T
```

**Sample input 2**

```
2
G T A C
```

**Sample output 2**

```
AA
AC
AG
AT
CA
CC
CG
CT
GA
GC
GG
GT
TA
TC
TG
TT
```

**Sample input 3**

```
3
G T A C
```

**Sample output 3**

```
AAA
AAC
AAG
AAT
```

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

## Problem information

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