## Exercise of Programming Language, Quiz 1

Write 4 Python programs to solve the following questions. Please name your program files as Q1.py, Q2.py, and so on, i.e., according to the serial number of questions. All data files you need can be obtained from the e3 system. Submit your programs to e3 before the end of the quiz.

- 1. Print out the AT content of the DNA sequence recorded in the file dna.txt.
- 2. The file *protein.txt* contains a protein sequence. Compute and print out the number of 20 standard types of amino acids that the protein comprises. Your output may look like this:

```
A: 12
C: 8
D: 1
... etc ...
(Hint: A, C, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, Y)
```

- 3. There are several blood pressure values saved in the file *blood\_pressures.txt*. Write a program to read the values and print to the screen their average. ( Hint: to make a string variable become a numeric variable with an integer value, you can use the function int() )
- 4. FASTA file format is a commonly-used DNA and protein sequence file format. Sequence(s) in FASTA format look like this:

```
>sequence_one
ATCGATCGATCGAT
>sequence_two
ACTAGCTAGCTAGCATCG
>sequence_three
ACTGCATCGATCGATCGTACCT
```

where the text after ">" is a header that describes the sequence (the ">" symbol indicates the start of the header line). Write a program that will create a FASTA file containing the following sequences.

Sequence header	DNA sequence
ABC123	ATCGACGATCGATCGCAGACGTATCG
DEF456	ACTGATGACGATGATCGACACGACT
HIJ789	ACTGACACTGTATGTACATGTG