Hand-in 4, due Tuesday, September 30

Background

You have been provided with a file in Absalon, which contains amino acid sequences for all known proteins in the Ecoli organism. It is in the so-called Fasta format, which means that for every protein, we first have a line starting with > containing the name:

>YBGC_ECOLI

and then some lines containing the sequence of amino acids:

MNTTLFRWPVRVYYEDTDAGGVVYHASYVAFYERARTEMLRHHHFSQQALMAERVAFVVRKMTVEYYAPA RLDDMLEIQTEITSMRGTSLVFTQRIVNAENTLLNEAEVLVVCVDPLKMKPRALPKSIVAEFKQ

Exercises

- 1. Create a module called handin4.py, and inside that module create a function called read_fasta that reads a fasta file and creates a dictionary, where the keys are the names of the proteins and the sequences are the values.
 - Create a new file, called handin4_test.py where you import the handin4 module and write some code that tests whether the function works on the Ecoli data file mentioned above.
- 2. In the same module, create another function called find_prot that takes a dictionary as first argument, and as second argument takes a protein name. The function should return the sequence corresponding to the provided name and provide an error message if the name is not present.
 - Again, write some test code in handin4_test.py that repeatedly (e.g. 3 times) calls this function on the dictionary from before include one case where the name is not present.
- 3. Finally, in the same module, create a function called find_prot2 that takes a dictionary and regular expression (as a string), and returns all of the keys in the dictionary that the pattern matches.
 - In the test program, use this function to count the number of the protein names in Ecoli that only consist of three letters before _ECOLI (e.g. VSR_ECOLI).

Please include both the Python code and the output produced from running the code. The code should be in a .py file, while the output should be a simple text file. You can easily save your output by copy&pasting from the PyCharm console window.

If your code produces external output files, please include those as well.

Please remember to comment your code, use doc-strings in function definitions and to use meaningful variable and function names!