

Introduction to Python

Day Two Exercises

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1 For-loops

First, define the following variables:

- `numbers = [0, 1, 1, 2, 3, 5, 8, 13]`
- `animals = ["gorilla", "canary", "frog", "moth", "nematode"]`

1. Loop over the list "numbers". At each iteration, print the value in "numbers" plus 2. Your code should print out the following:

```
2
3
3
4
5
7
10
15
```

2. Loop over the list "animals". At each iteration, print out the value in "animals" followed by its length. Your code should print out the following:

```
gorilla 7
canary 6
frog 4
moth 4
nematode 8
```

3. Modify the previous for-loop to print out the capitalized version of each animal (do not redefine anything, just print!). Your code should print out the following:

```
GORILLA
CANARY
FROG
MOTH
NEMATODE
```

4. Write a new for-loop to create a new list called "cap_animals" which should contain the capitalized versions of all entries in "animals." For this task, you will need to define the list "cap_animals" before the for-loop, and use `.append()` to build up this list as you go. Print the final list after the for-loop.
5. Write a for-loop to create a new list called "negative_numbers" which should contain the negative values of the entries in "numbers", following a similar procedure to the previous task. Once this list is complete, write an `if/else` statement to check if the sum of "negative_numbers" equals -1 times the number of "numbers". Use the function `sum()` in the if statement. Print informative messages in the `if/else` construct.

6. Write a for-loop, using the `range()` function, to print the powers of 2 from 2^0 to 2^{15} . (Note that in Python, the exponent symbol is `**`, as in `3**2 = 9`).

2 Working with dictionaries

1. Define this dictionary: `molecules = {"DNA": "nucleotides", "protein": "amino acids", "hair": "keratin"}`, and perform the following tasks:
 - (a) Create two lists called `molecules_keys` and `molecules_values`, comprised of the keys and values in `molecules`, respectively. Use dictionary methods for this task.
 - (b) Add the key:value pair `"ribosomes": "RNA"` to the `molecules` dictionary. Print the dictionary to confirm.
 - (c) Add yet another key:value pair, `"ribosomes": "rRNA"`, to the `molecules` dictionary, and print out the new dictionary. Understand why the result contains the key:value pairs shown.
2. Congratulations, you've been hired as a zoo-keeper! Now you have to feed the animals. You received these handy Python dictionaries which tells you (a) to which category each animal belongs, and (b) what to feed each animal category:

```
category = {"lion": "carnivore", "gazelle": "herbivore", "anteater":  
"insectivore", "alligator": "homovore", "hedgehog": "insectivore", "cow":  
"herbivore", "tiger": "carnivore", "orangutan": "frugivore"}
```

```
feed = {"carnivore": "meat", "herbivore": "grass", "frugivore": "mangos",  
"homovore": "visitors", "insectivore": "termites"}
```

- (a) Copy and paste these dictionaries into a Python script. Use indexing to determine what you should feed the orangutan and print the result.
- (b) Write a for-loop to loop over "feed" and print out what food each animal type eats. Your code should ultimately print the following (in any order!):
The carnivore eats meat
The herbivore eats grass
The frugivore eats mangos
...etc
Hint: You might find it helpful to first loop over the "feed" dictionary and simply print the loop variable. Extend the code from there to print the full sentence.
- (c) Write a for-loop to print out what each animal eats. Your code should ultimately print the following (in any order!):
The lion eats meat
The gazelle eats grass
The anteater eats termites
The alligator eats visitors
... etc
For this task, you should loop over the dictionary "category" and use indexing to obtain the relevant information from the "food" dictionary to create your sentence.
Finally, modify the previous for-loop so that it creates a new dictionary called "animals_eat" while looping over "category". This dictionary should contain the exact animal:food pairs, e.g. "lion": "meat" will be one key:value pair. Print out the resulting dictionary.

3 For-loop and if/else

1. A professor has decided to curve grades in a very special way: grades above 95 are reduced by 10%, grades between 75-95 (inclusive) remain the same, and grades below 75 are raised by 10%. You have been tasked with crunching the numbers.

- (a) Create a list of new grades that reflects these rules from the following grade list:
`grades = [45, 94, 25, 68, 88, 95, 72, 79, 91, 82, 53, 66, 58]`
- (b) The professor has changed his mind: he now wants to use a scaling factor of 0.078325 (instead of 0.1), because why not! Recompute the grades from part 1 using this new scaling.
- (c) Finally, modify your code (if you need to) so that the scaling value (either 0.1 or 0.078325) used in your for-loop is a *pre-defined variable* to avoid hard-coding the scaling value.
- (d) (Note: this question should be skipped if you are pressed for time!) The *nested* list below contains three sets of grades for silly professor's three class sections:
`all_grades = [[45, 94, 25, 68, 88, 95, 72, 79, 91, 82, 53, 66, 58], [23, 46, 17, 67, 55, 42, 31, 73], [91, 83, 79, 76, 82, 91, 95, 77, 82, 77]]`
 Create a new nested list with the curved grades for each of these groups, using a scaling factor of 0.06.

2. This dictionary provides the molecular weight for all amino acids:

```
amino_weights = {"A":89.09, "R":174.20, "N":132.12, "D":133.10, "C":121.15,
                 "Q":146.15, "E":147.13, "G":75.07, "H":155.16, "I":131.17, "L":131.17,
                 "K":146.19, "M":149.21, "F":165.19, "P":115.13, "S":105.09, "T":119.12,
                 "W":204.23, "Y":181.19, "V":117.15}.
```

Perform the following tasks with this dictionary (for ease, copy/paste it into a python script):

- Determine the molecular weight for this protein sequence:
`"GAHYADPLVKMPWRTHC"`.
- This protein sequence, `"KLSJXXFOWXNN CPR"` contains some ambiguous amino acids, coded by "X" and "J". Calculate the molecular weight for this protein sequence. To compute a weight for an ambiguous amino acid "X" and "J", use the *average* amino acid weight.
 Hint: the `len()` and `sum()` functions and the `.values()` dictionary method will be useful! The `len()` and `sum()` functions can be used together to compute a mean value of a list.

4 File Input and Output

Be sure to always close the file once you are done with it! If you use `with` control flow, Python will close the file automatically. The following exercises make use of files distributed in today's course materials. Be sure that you are in the correct directory when interacting with the files!

1. Open the file `"file1.txt"` in read-mode, and print its contents to screen. For this, you should use the `.read()` method, which saves the contents of the file to a single string. Perform this task twice: once using `open` and `close`, and once using `with` control-flow.
2. Open the file `"file1.txt"` in read-mode, and save all lines in this file to a list using the `.readlines()` method. Write a new file called `"upper_file1.txt"` which contains the same contents of `"file1.txt"` but in upper-case. Try to do this task using a single for-loop, and don't forget that in order to write newlines (the "enter" key) to a file, you must include `"\n"` in the string you are writing to file!
3. Open the file `"upper_file1.txt"` in append-mode, and *append* the sentence: `"I just created this upper-case file!"` to the bottom of the file. Close the file and open it (separately) to examine the contents.
4. Again, open the file `"upper_file1.txt"`, this time in read-mode. Loop over the file lines *without* using `.read()` or `.readlines()`. Print out lines as you loop.
5. Modify the previous for-loop to only print out lines in `"upper_file1.txt"` which contain at least (i.e. `>=`) 5 letter "E"s. This will require an `if` statement as well as the method `.count()`.
6. You should notice 20 files named `file1.txt`, `file2.txt`, ...`file20.txt`. Write a for-loop to open each of these files (Hint: use the `range()` function to loop over file names). For each file, write a for-loop over the file lines (this does not use `.read()` or `.readlines()`). Print each line that contains more than 25 characters (determine this with `len()`).