

Department of Molecular Life Sciences



BIO334

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courtesy of Danaila, Sumner, Lyon, Tackmann, Dmitrijeva & Gable

https://github.com/meringlab/Bio334.git

What is Python?

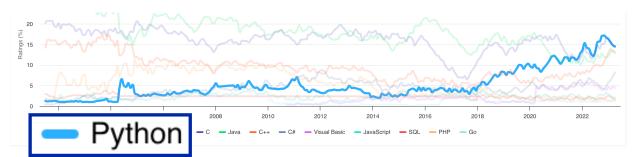
- Dynamic, interpreted, open source programming language
- Simple syntax with fast learning curve

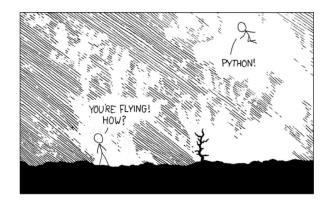
Octoverse by Github: Top languages used in 2022

Stack Overflow 2022: most popular languages

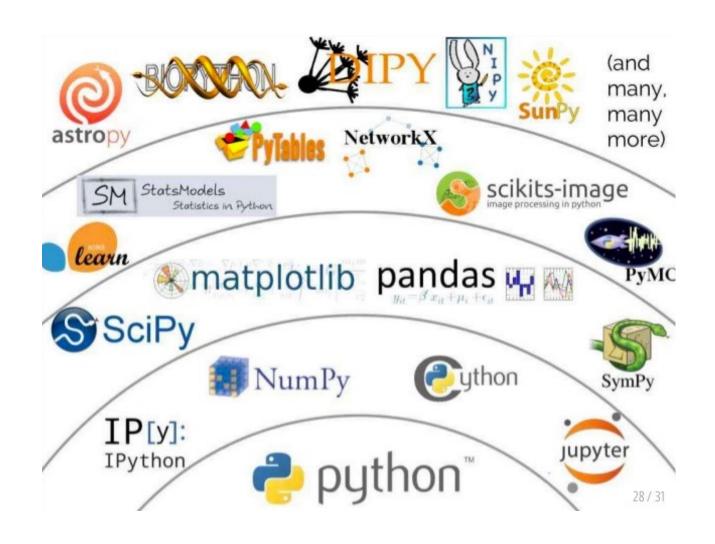


TIOBE Programming Community Index





Scientific Python stack: more than core Python



Today's program

- 1. Introduction to programming: refresh the basic concepts with small hands-on sessions in JupyterLab
- 2. Break
- 3. Writing Python code to solve Exercises 1 and 2 (and optional Exercises 3 and 4)
- 4. Optional: go through solutions of exercise 1 and 2 at 4pm

Part 1: Programming basics

Variables

Store a piece of data and give it a specific name with the "=" (equal) operator

```
a = 4
pi = 3.14159

my_string = "hello"
# single and double ticks are equal
# (just stick to one) 'hello' or "hello"

my_protein_sequence = 'MRHIAHTQRCLSRLTSLVALLLIVLP'
```

Basic **rules** for variable names:

- NO spaces
- don't start with numbers
- use descriptive names

Operators

Arithmetic operations

- addition + , subtraction -
- division /, multiplication *
- exponent **

Boolean operations

- return True or False
- comparison operators: == , > , <
- logical operators: and, or, not

Get ready for live sessions in JupyterLab: Download course files

- 1. open a **Terminal window** (e.g. Spotlight search: terminal):
- 2. change to your personal course folder (or where you would like the files to go)
 - > cd /Volumes/Kurs/UsersBio/mheiml #replace with your shortname
- 3. clone the git repository by entering:
 - > git clone https://github.com/meringlab/Bio334.git
- checkout our course files under 02_python_introduction
- > next up: installation and short demo on how to use JupyterLab

Get ready for live sessions in JupyterLab: (Install and) launch JupyterLab

open a **Terminal window** and enter the following commands

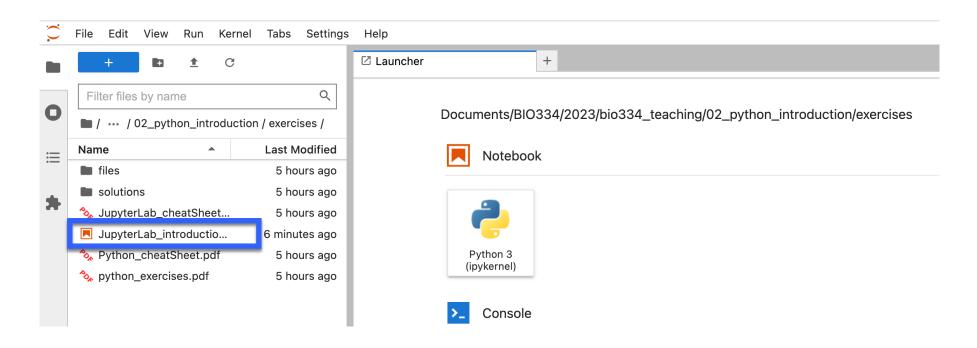
- > cd /Volumes/Kurs/UsersBio/yourShortname
- > . /opt/miniconda3/etc/profile.d/conda.sh #space after dot matters!
- > conda create --prefix /Volumes/Kurs/UsersBio/yourShortname/py38_envs
 python=3.8 #copy whole line starting from conda
- > conda activate /Volumes/Kurs/UsersBio/yourShortname/py38_envs
- > conda install jupyter #press y and Enter when asked
- > jupyter lab #to open jupyter lab

first time: run all the commands, blue and yellow

after this: run only the blue commands to open JupyterLab again

Get ready for live sessions in JupyterLab: start your first Jupyter Notebook

- in Jupyter: go to Bio334/02_python_introduction/exercises
- launch the Jupyter Notebook JupyterLab_introduction.ipynb for an interactive session



Session I: variables and basic operations

```
welcome message = 'hello world!' # hit [shift + enter] after every line
welcome message # more general alternative: print(welcome message)
# Arithmetics: use python as your advanced calculator
a = 4
b = a + 3
(a + b) * 4
a / 8
a**2
# Let's try with strings
welcome message + welcome message
a = '4'
a + 3 # anything strange?
# whats the difference?
a + str(3)
int(a) + 3
```

Discussion session I: Arithmetics

```
[1]: welcome_message = 'hello world!' # hit [shift + enter] after every line
[2]: welcome_message
[2]: 'hello world!'
[3]: # Arithmetics: use python as your advanced calculator
     a = 4
[4]: b = a + 3
[5]: (a + b) * 4
[5]: 44
[6]: a/8
[6]: 0.5
[7]: a**2
[7]: 16
```

Discussion session I: Concatenation

```
[8]: # Let's try with strings
 [9]: # use + to add two strings
     welcome_message + welcome_message
 [9]: 'hello world!hello world!'
[10]: a = '4'
[11]: # We need to have the same type to add elements
[12]: a + 3 # anything strange?
                                                Traceback (most recent call last)
     TypeError
      /var/folders/3s/6lmkv5_10g1g4vx768t1yh_0g4tfkb/T/ipykernel_50845/173162209.py in <module>
      ---> 1 a + 3 # anything strange?
     TypeError: can only concatenate str (not "int") to str
 [ ]: # whats the difference?
[13]: a + str(3)
[13]: '43'
[14]: int(a) + 3
[14]: 7
```

List data structure

 like a shopping list we have an object that can store multiple objects at once.

```
my list = ['butter','milk','oranges']
```

it can hold different objects as well

```
my_list = ['butter',1,1.5,'milk']
```

 every element has an index, starting from 0 which you can access with the square brackets []

```
>>> my_list[0]
'butter'
```

List data structure

get subset of a list ("slicing")

```
>>> my_list[0:2]
['butter','milk']
```

assign new values to list elements

```
my_list[0] = 'bananas'
```

- powerful list operations
 - e.g. append(), len(), sort(), reverse(), insert()

Session II: Work with lists

```
# create your first list
# try accessing it with indices
my list = [1, 2, 3, 4, 5]
my list[1]
my list[5]
my list[0:3]
my list[-1]
len(my list)
# mixed lists:
# put different data types in your list
my mixed list = ['UZH', 'founded in', 1934]
my mixed list[2] = 1834
del my mixed list[1]
my mixed list.append('A.D.')
my mixed list
```

```
# apply arithmetic operators on lists
[1,2,3] + [3,4,6] # list concatenation
['Hello'] * 4
```

Discussion session II: Accessing list elements

```
[22]: # create your first list and try accessing it with indices
      my_list = [1,2,3,4,5]
[18]: my_list[1]
[18]: 2
[19]: my list[5]
                                                Traceback (most recent call last)
      IndexError
      /var/folders/3s/6lmkv5_10g1g4vx768t1yh_0g4tfkb/T/ipykernel_50845/303719451.py in <module>
      ----> 1 my_list[5]
      IndexError: list index out of range
[42]: my_list[0:3] # slice your list with index ranges
[42]: [1, 2, 3]
[44]: my_list[-1] # get the last element with the index -1
[44]: 5
[45]: len(my_list) # check how many elements your list contains
[45]: 5
```

Discussion session II: List modifications and arithmetic operations on lists

```
[55]: # mixed lists, put different variable types in your list
      my_mixed_list = ['UZH','founded in',1934]
[56]: my_mixed_list[2] = 1834
[57]: del my_mixed_list[1] # delete an element at a specific index
[58]: my_mixed_list.append('A.D.') # add a new item to the end of the list
[59]: my_mixed_list
[59]: ['UZH', 1834, 'A.D.']
[60]: # apply the arithmetic operators on lists
      [1,2,3] + [3,4,6]
[60]: [1, 2, 3, 3, 4, 6]
[61]: ['Hello'] * 4
[61]: ['Hello', 'Hello', 'Hello', 'Hello']
```

Strings & Lists

- Strings can also be considered lists of characters and can be accessed by index
- To convert them to an actual list (and make them mutable) use list()

```
>>> my_sequence = 'MRHIAHTQRCLSR'
>>> list(my_sequence)
['M', 'R', 'H', 'I', 'A', 'H', 'T', 'Q', 'R', 'C', 'L', 'S', 'R']
```

- While this opens many possibilities, check the cheat-sheet for convenient built-in string operations
 - e.g. split(), join(), replace()

Dictionaries

Similar to lists, but elements are accessed through a user defined key

```
my_proteins_seqs = {}
my_proteins_seqs = dict() # same as above
my_proteins_seqs['DROME_HH_Q02936'] = 'MRHIAHTQRCLSRLTSLVA'
my_proteins_seqs['DROME_PATC_P18502'] = 'MDRDSLPRVPDTHGDVVD'
```

retrieve their value by using the key

```
>>> my_proteins_seqs['DROME_HH_Q02936']
'MRHIAHTQRCLSRLTSLVA'
```

Session III: Strings and dictionaries

```
# Try accessing a string like a list # create your first dictionary
                                        dna to rna = {}
my string = 'hello world!'
                                        dna to rna['A'] = 'A'
my string[6]
                                        dna to rna['T'] = 'U'
                                        dna to rna['C'] = 'C'
my string[-1] = '?'
                                        dna to rna['G'] = 'G'
my list = list(my string)
                                        dna to rna
my list[-1] = '?'
my modified string = ''.join(my list)
                                      #or in one line
                                        dna to rna =
                                        {'A':'A','T':'U','C':'C','G':'G'}
my modified string
                                        #dictionary operations
                                        dna to rna['U']
                                        'U' in dna to rna
                                        dna to rna.keys()
                                        dna to rna + dna to rna
```

Discussion session III: String lists

```
[65]: # Try accessing a string like a list
     my string = 'hello world!'
[72]: my_string[6] # accessing a character by index
[72]: 'w'
[73]: my string[-1] = '?' # changes are not allowed - strings are immutable!
                                                Traceback (most recent call last)
      TypeError
      /var/folders/3s/6lmkv5_10g1g4vx768t1yh_0g4tfkb/T/ipykernel_50845/280325205.py in <module>
      ----> 1 my_string[-1] = '?' # changes are not allowed - strings are immutable!
     TypeError: 'str' object does not support item assignment
[75]: my list = list(my string) # Convert your string to a list to change it
[76]: my list[-1] = '?'
[77]: my modified string = ''.join(my list) # combine your list into a string again
     my modified string
[77]: 'hello world?'
```

Discussion session III: Dictionary definition

```
[78]: # create your fist dictionary

[82]: dna_to_rna = {}
    dna_to_rna['A'] = 'A'
    dna_to_rna['T'] = 'U'
    dna_to_rna['C'] = 'C'
    dna_to_rna['G'] = 'G'
    dna_to_rna

[82]: {'A': 'A', 'T': 'U', 'C': 'C', 'G': 'G'}

[80]: #or in one line
    dna_to_rna = {'A': 'A', 'T': 'U', 'C': 'C', 'G': 'G'}
```

Discussion session III: Dictionary usage

```
[117]: # dictionary operations
[118]: dna_to_rna['U'] # access an element with its key
                                                 Traceback (most recent call last)
       KevError
       /var/folders/3s/6lmkv5_10g1g4vx768t1yh_0g4tfkb/T/ipykernel_50845/3658080417.py in <module>
       ----> 1 dna_to_rna['U'] # access an element with its key
      KeyError: 'U'
[119]: 'U' in dna_to_rna # Test if a key is in the dictionary
[119]: False
[120]: dna to rna.keys() # see all keys in the dictionary
[120]: dict_keys(['A', 'T', 'C', 'G'])
[121]: dna to rna + dna to rna # Dictionaries don't support concatenation
       TypeError
                                                 Traceback (most recent call last)
       /var/folders/3s/6lmkv5 10g1g4vx768t1yh 0g4tfkb/T/ipykernel 50845/1693159149.py in <module>
       ----> 1 dna to rna + dna to rna # Dictionaries don't support concatenation
      TypeError: unsupported operand type(s) for +: 'dict' and 'dict'
```

Functions

- function: stores instructions (not values)
- basic use: function_name(arguments)
- many functions loaded by default in python, e.g.
 - str() convert an object into a string
 - int() convert an object into an integer
 - float() convert a object into a floating point number
 - type() tells you the type of an object
- see many others in the cheat-sheet and find them on https://docs.python.org/3/library/functions.html

Function example

```
[122]: # Functions
[123]: # let's create a function using def (for define)
      def add_things(a, b):
          result = a+b
          return result
[124]: add_things(3,5)
[124]: 8
[125]: add_things('Hello ', 'there')
[125]: 'Hello there'
[127]: add things('3',5) # reminder: we need to have the same type to add elements
      TypeError
                                                Traceback (most recent call last)
      /var/folders/3s/6lmkv5_10g1g4vx768t1yh_0g4tfkb/T/ipykernel_50845/3573978534.py in <module>
       ----> 1 add_things('3',5) # reminder: we need to have the same type to add elements
       /var/folders/3s/6lmkv5_10g1g4vx768t1yh_0g4tfkb/T/ipykernel_50845/2105753661.py in add_things(a,
            1 # let's create a function using def (for define)
            2 def add_things(a, b):
        ---> 3 result = a+b
               return result
      TypeError: can only concatenate str (not "int") to str
```

Methods

- functions of a specific object class
- access by <variable_name>.<method>(arguments)

```
>>> s = 'The quick brown fox jumps over the lazy dog'
>>> s.split()
['The', 'quick', 'brown', 'fox', 'jumps', 'over', 'the', 'lazy', 'dog']
>>> s.split('fox')
['The quick brown ', 'jumps over the lazy dog']
```

- in iPython: write . after a variable name and press <tab> to get an overview about available methods
- alternatively: dir(object)
- use ? / ?? (in ipython) or help() to get information about a method and its arguments e.g. ?my_str.split or help(my_str.split)
- alternatively: google or ask chatGTP

Break

Part 2: Writing Python code

Why write a script?

- Organize your commands in a text file and build more complicated workflows that can be executed at once.
- save typing, make your work reproducible

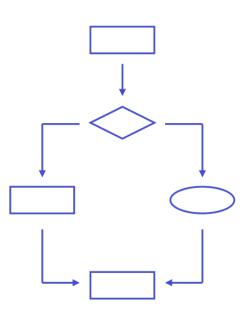
- Run it at any point by executing your script
- Use comments (#) to describe your code
- use the print() function to print the output or an intermediate result

```
>>> print("Hello", "Python!")
Hello Python!
```

Conditional statements

Use if/else clauses to make decisions

```
if boolean_expression:
    print("The statement is True")
else:
    print("The statement is False")
```



- Remember the indentation!
 - While other programming languages use brackets or end statements, Python uses whitespace to structure code.
 - > Simply use tabulator to indent.



Loops

- Loops are essential for repeating an action several times
- The for loop executes the nested statements as many times as there are elements in the input list.

```
input_list = [1,2,3,4,5]
for my_number in input_list:
    print(my_number)

1
2
3
4
5
```

for loop for reading text files

```
f = open('my_file.txt','r')
for line in f:
    print(line)
f.close()

1st line

2nd line

3rd line
```

File: my_file.txt

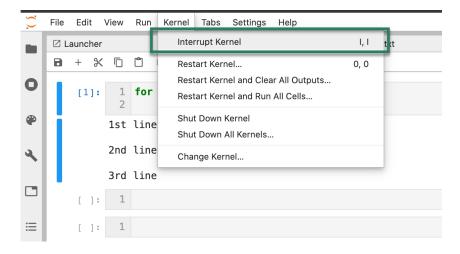
1st line
2nd line
3rd line
EOF

Conditional loop

Use the while loop to continue an action until a condition is not satisfied anymore

```
>>> number_apples = 3
>>> while number_apples > 0:
...     number_apples -= 1
...     print("Ate an
apple,",number_apples,'left')
...
Ate an apple, 2 left
Ate an apple, 1 left
Ate an apple, 0 left
```

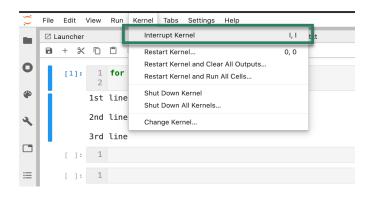
in case you need to terminate your loop...



Session IV: Loops and friends

```
# 1. try out this for loop
for my number in range(6):
    print(my number)
# 2. A new loop key-word: continue
for my number in range(6):
    if my number == 3:
        continue
    print(my number)
print('after loop: ', my number)
# 3. try using break instead of continue
# 4. when will this while loop finish?
my number = 1
from time import sleep
while my number < 5:
    print("hurray! my number is increasing:", my number)
    sleep(1)
```

in case you need to terminate your loop...



Discussion session IV: Loops and friends

```
[129]: # 1. try out this for loop
for my_number in range(6):
        print(my_number)

0
1
2
3
4
5
```

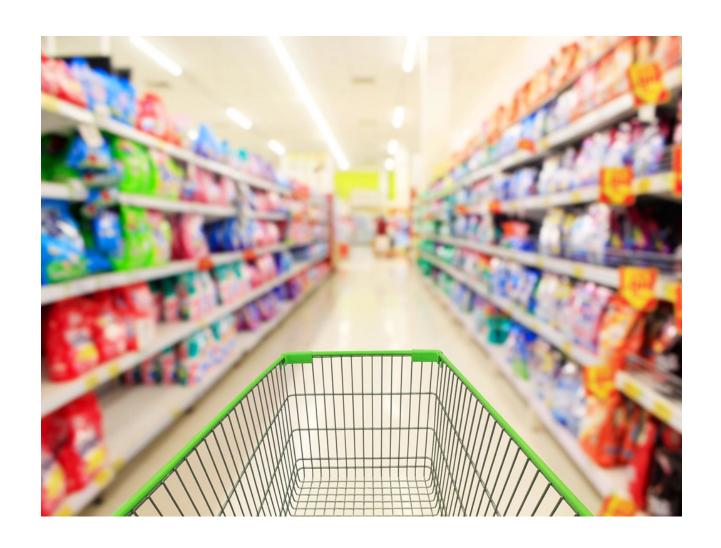
Discussion session IV: Loops and friends

```
[141]: # 2. A new loop key-word: continue
       for my_number in range(6):
           if my_number == 3:
               continue # skips the current iteration
           print(my number)
       print('after loop: ', my_number) # loop continued until the end
       after loop: 5
[142]: # 3. A new loop key-word: continue
       for my_number in range(6):
           if my_number == 3:
               break # terminates the loop immediately
           print(my number)
       print('after loop: ',my_number) # loop has terminated prematurely
       after loop: 3
```

Discussion session IV: Loops and friends

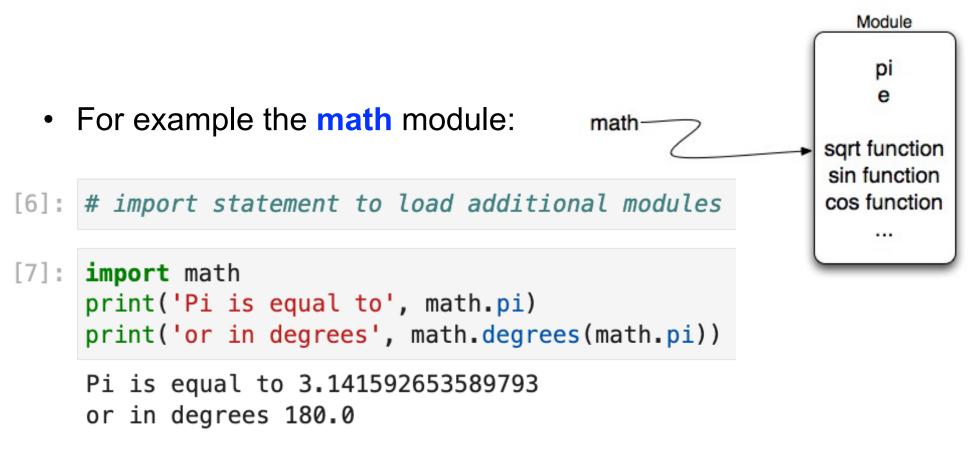
```
[144]: # 4. when will this while loop finish?
      my number = 1
      from time import sleep
      while my number < 5:
           print("hurray! my number is increasing:", my number)
          # my number += 1 # don't forget to update your counter!!
           sleep(1)
      hurray! my number is increasing: 1
       hurray! my number is increasing: 1
       hurray! my number is increasing: 1
      hurray! my number is increasing: 1
       hurray! my number is increasing: 1
       hurray! my number is increasing: 1
       hurray! my number is increasing: 1
       hurray! my number is increasing: 1
       KeyboardInterrupt
                                                Traceback (most recent call last)
       /var/folders/3s/6lmkv5_10g1g4vx768t1yh_0g4tfkb/T/ipykernel_50845/3886309862.py in <module>
                   print("hurray! my number is increasing:", my number)
                 # my number += 1 # don't forget to update your counter!!
       ---> 7 sleep(1)
       KeyboardInterrupt:
```

python modules



Import statement

Use import to load additional modules for more functionalities



more details at: https://docs.python.org/3/library/math.html

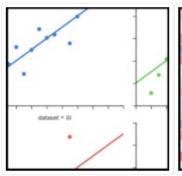
Much more to discover

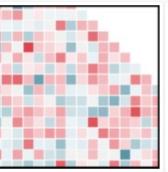
- ipython magics make your life a lot easier
 - time your code, debug it, call other languages
- make python lightning fast using numpy, scipy, and cython/pypy/numba
- visualize your data with seaborn, bokeh and matplotlib

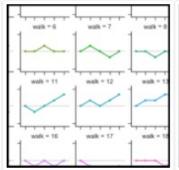


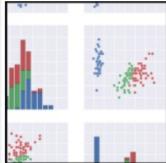


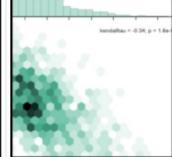


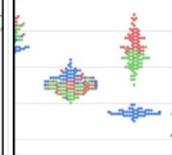












Further reading

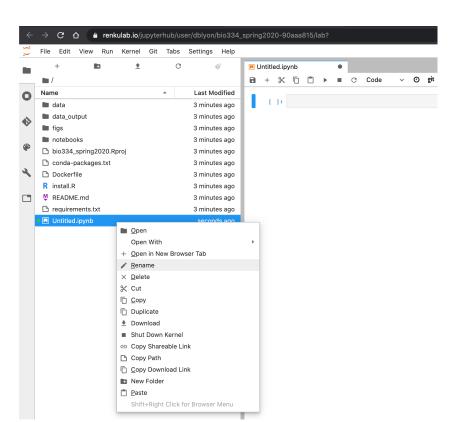
- <u>https://snakify.org/</u> (interactive tutorial)
- <u>http://www.diveintopython.net/</u>(comprehensive, general purpose)
- http://swcarpentry.github.io/python-novice-inflammation/ (scientific python by example)
- <u>http://www.scipy-lectures.org/intro/intro.html</u> (intro by the scientific python consortium)
- https://github.com/dblyon/pandasintro (extensive introduction to pandas)
- https://cs50.harvard.edu/college/2021/spring/weeks/0/

Exercise session

 Create a new Python3 Jupyter Notebook, save it to the folder /Bio334/02_python_introduction/exercises and start exploring



- Change the name of your newly created file by right clicking on *Untitled.ipynb*
- To execute a cell press Shift + Enter



optional exercise slides follow

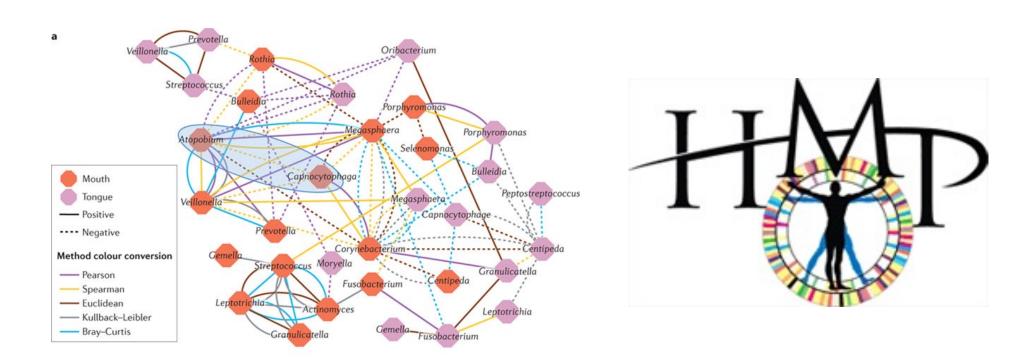
Optional Exercise #3

Learn the basics of **Pandas**, a powerful python module for data analysis, including reading, writing, filtering, merging, arithmetic operations on and sorting of tabular data



Optional Exercise #4

Create a pipeline to infer simple ecological relationships in the Human Microbiome



Optional exercise session

- Create a new python file in JupyterLab
- To execute, launch a terminal tab and enter
 - ipython <script_name>.py

Good luck!

sys module

- A built-in module that contains system-specific parameters and functions
- For example we can use it to read arguments from the command line:

```
Script: say_hello.py
import sys
print('Hello there', sys.argv[1])
# Command line arguments are in sys.argv[1], sys.argv[2]...
# sys.argv[0] is the script name itself
```

davide\$ python say_hello.py bio334_Students
Hello there bio334_Students

os module

- Another built-in module that contains miscellaneous operating system interfaces
- For example we can use it to obtain the name of the user currently logged in:

Script: check_directory.py

```
import os

dir_ = "Bio334"

f not os.path.exists(dir_):
    print("It seems you haven't cloned the git repository yet. \
        Please enter the following command in a terminal window.")
    print("git clone https://github.com/meringlab/Bio334.git")

else:
    print("Great! The 'Bio334' directory exists. You are good to go.")
```

Run the script in a terminal window

```
base > work > bio334_spring2020_dbl > master > 3? > $ > python check_directory.py
Great! The 'Bio334' directory exists. You are good to go._
```