



University of
Zurich^{UZH}

Department of Molecular Life Sciences



Introduction to Python

BIO334

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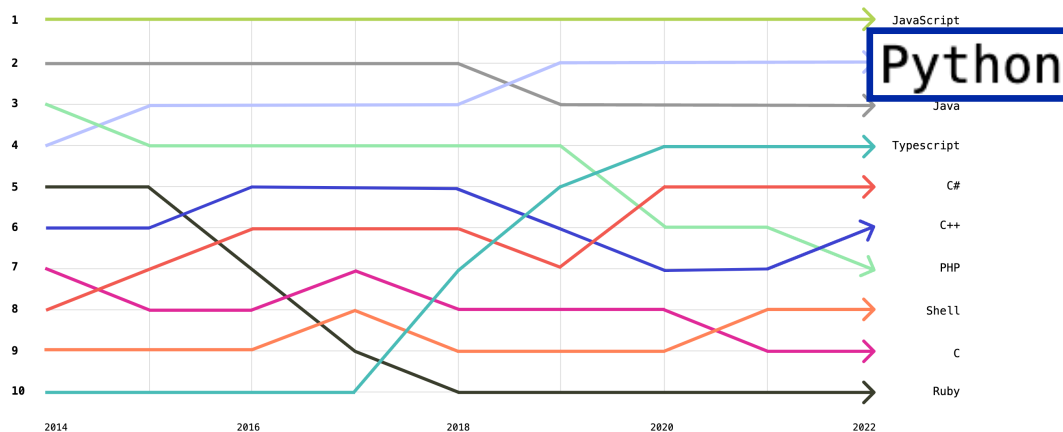
courtesy of Danaila, Sumner, Lyon, Tackmann, Dmitrijeva & Gable

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

What is Python?

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

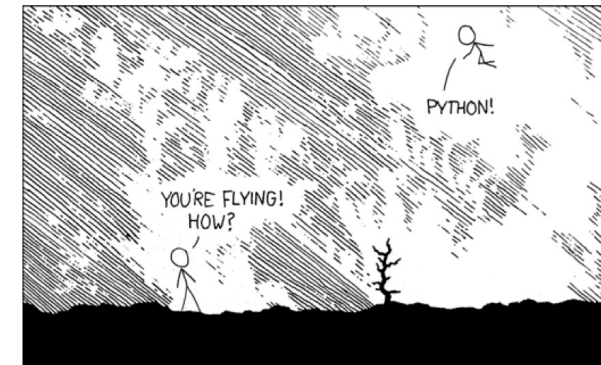
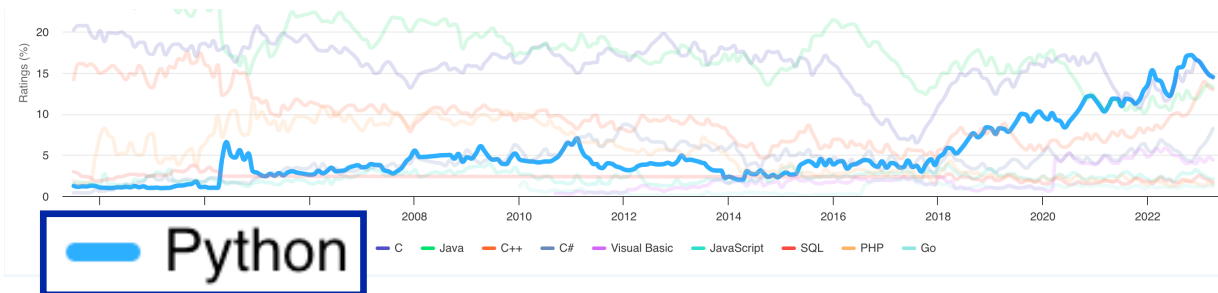
Octoverse by Github: Top languages used in 2022



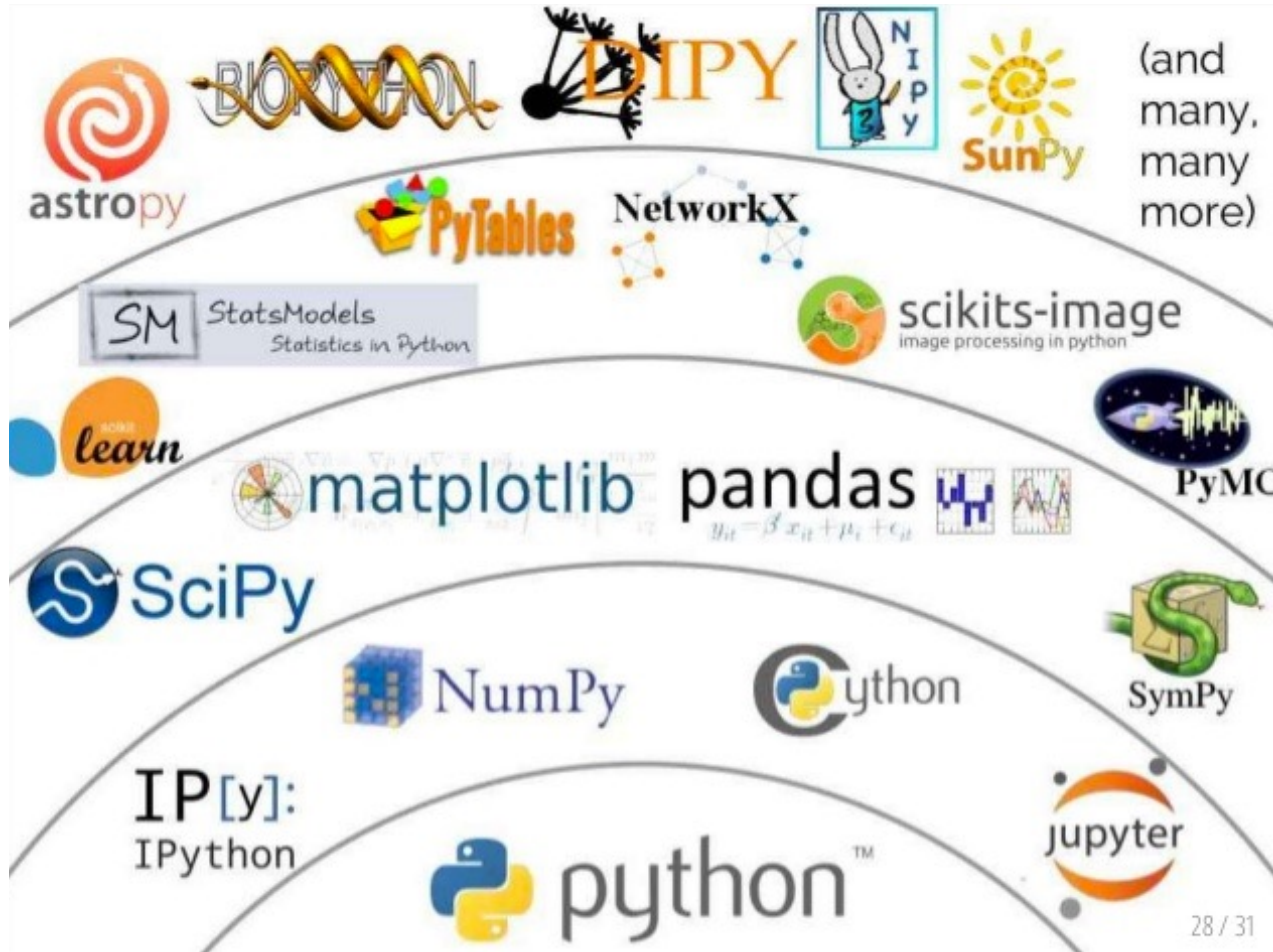
Stack Overflow 2022: most popular languages



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 **

```
>>> 2 + 3
5
>>> 23 - 3
20
>>> 22.0 / 12
1.8333333333333333
>>> (1 + 2) * 3
```

Boolean operations

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

```
>>> 1 < 2
True
>>> 23 == 45
False
>>> 2<3 and 4!=3
True
```

Get ready for live sessions in JupyterLab:

Download course files

1. open a **Terminal window** (e.g. Cmd-f: terminal):
2. change to the desired folder, where you would like the files to go
 - to save it to your personal course folder, enter:
`cd /Volumes/Kurs/UsersBio/mheiml` replace with your shortname
 - to save it to this computer (might not be permanently stored), enter:
`cd Documents`

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

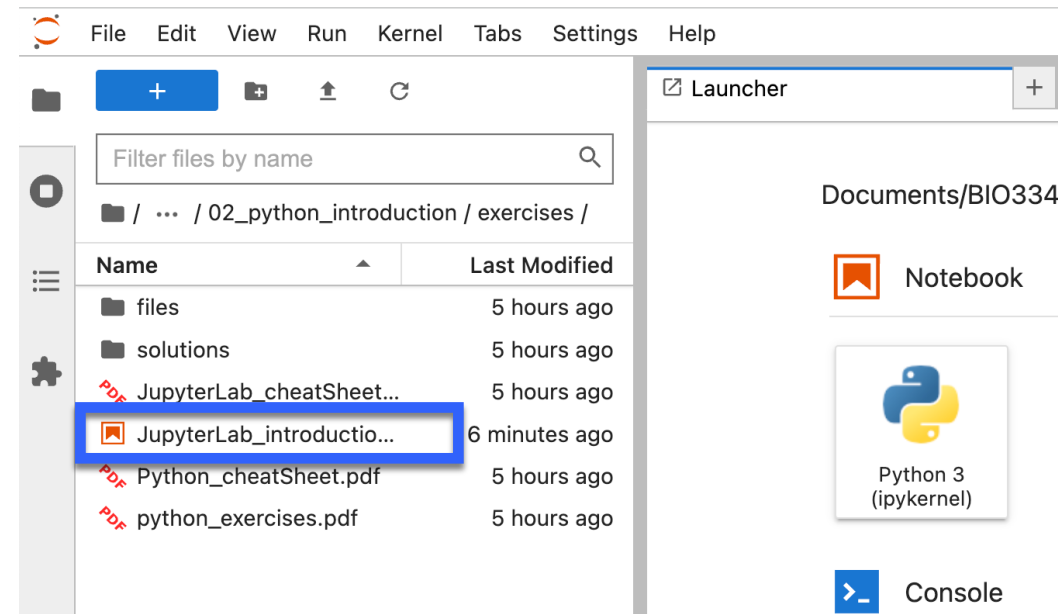
```
. /opt/miniconda3/etc/profile.d/conda.sh
```

```
conda create --prefix ~/Documents/py38_envs python=3.8
```

```
conda activate ~/Documents/py38_envs
```

```
conda install jupyter
```

- 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?
```

```
-----  
TypeError                                Traceback (most recent call last)  
/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]
```

```
-----  
IndexError                                Traceback (most recent call last)  
/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

```
my_string = 'hello world!'
```

```
my_string[6]
```

```
my_string[-1] = '?'
```

```
my_list = list(my_string)
```

```
my_list[-1] = '?'
```

```
my_modified_string = ''.join(my_list)
```

```
my_modified_string
```

create your first dictionary

```
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
```

#or in one line

```
dna_to_rna =
```

```
{ 'A': 'A', 'T': 'U', 'C': 'C', 'G': 'G' }
```

#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!
```

```
-----  
TypeError                                Traceback (most recent call last)  
/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 first 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
```

```
-----  
KeyError                                Traceback (most recent call last)  
/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  
      4     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

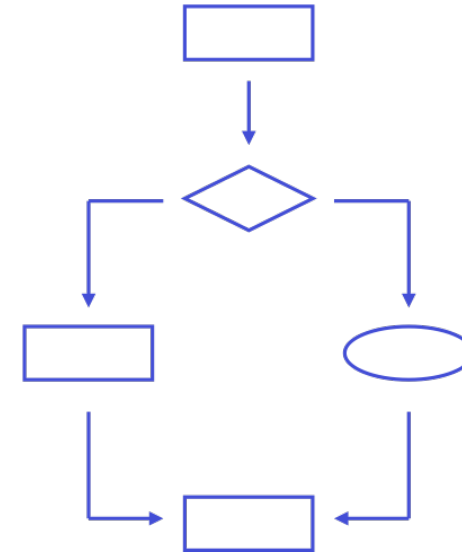
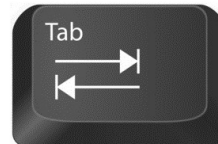
```
>>> print("Hello", "Python!")  
Hello Python!
```

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

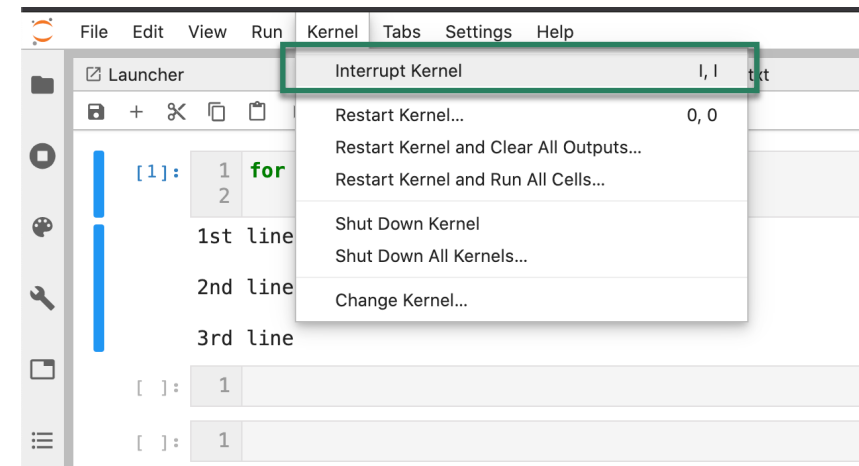
EOF

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

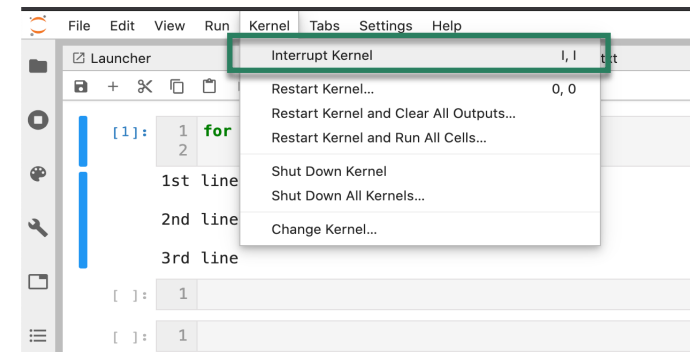
```
# 1. try out this for loop
for my_number in range(6):
    print(my_number)
```

```
# 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
```

```
0
1
2
4
5
after loop: 5
```

```
[142]: # 3. A new loop key-word: break
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
```

```
0
1
2
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>
      5     print("hurray! my number is increasing:", my_number)
      6     # 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

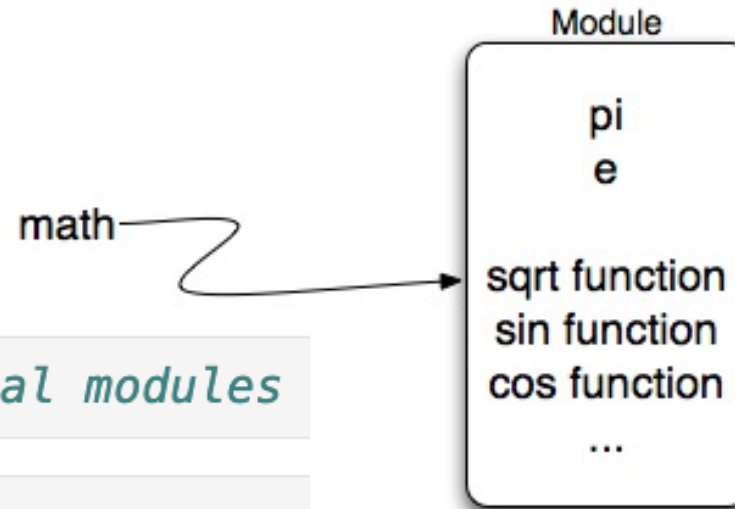
- For example the **math** module:

```
[6]: # import statement to load additional modules
```

```
[7]: import math  
print('Pi is equal to', math.pi)  
print('or in degrees', math.degrees(math.pi))
```

```
Pi is equal to 3.141592653589793  
or in degrees 180.0
```

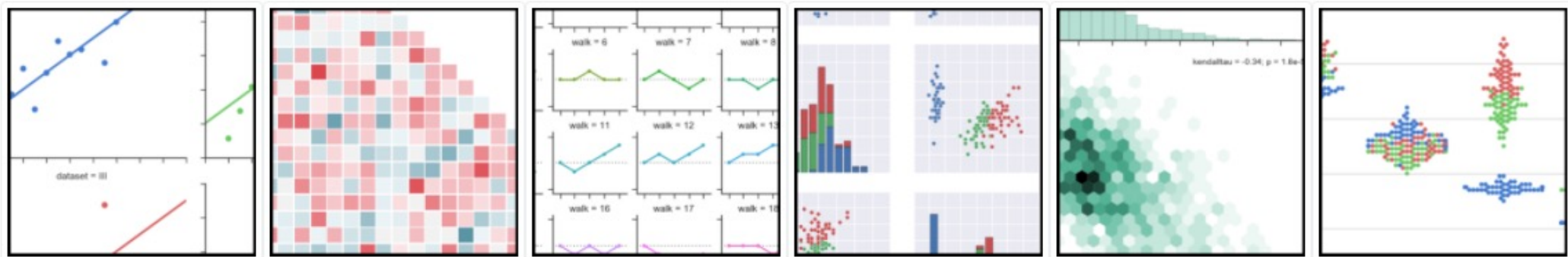
more details at: <https://docs.python.org/3/library/math.html>



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

IP[y]:
IPython

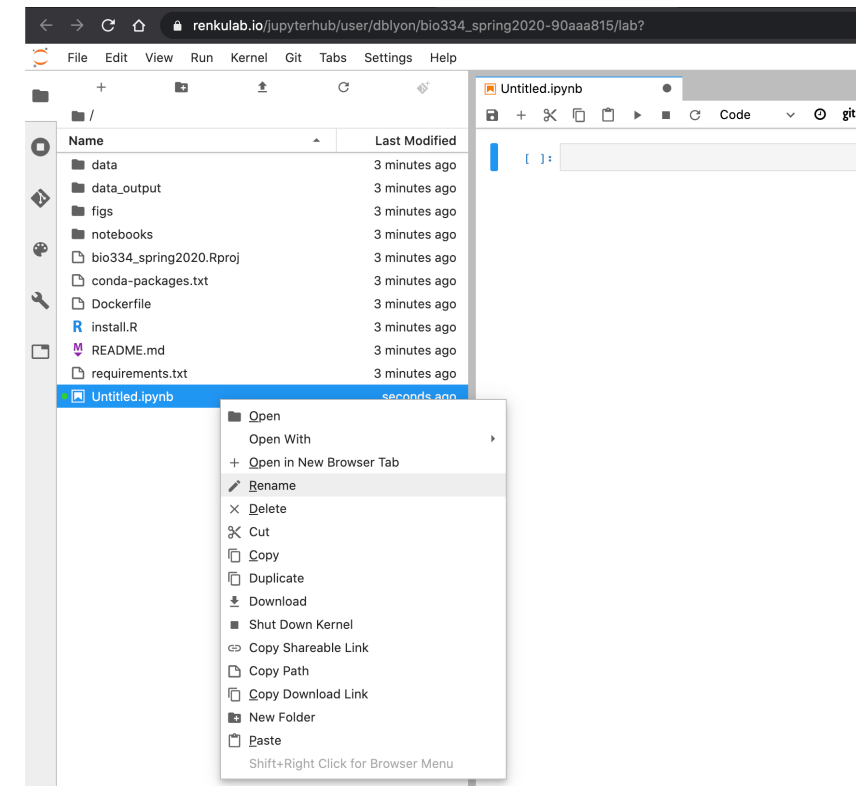


Further reading

- <https://snakify.org/> (interactive tutorial)
- <http://www.diveintopython.net/> (comprehensive, general purpose)
- <http://swcarpentry.github.io/python-novice-inflammation/> (scientific python by example)
- <http://www.scipy-lectures.org/intro/intro.html> (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

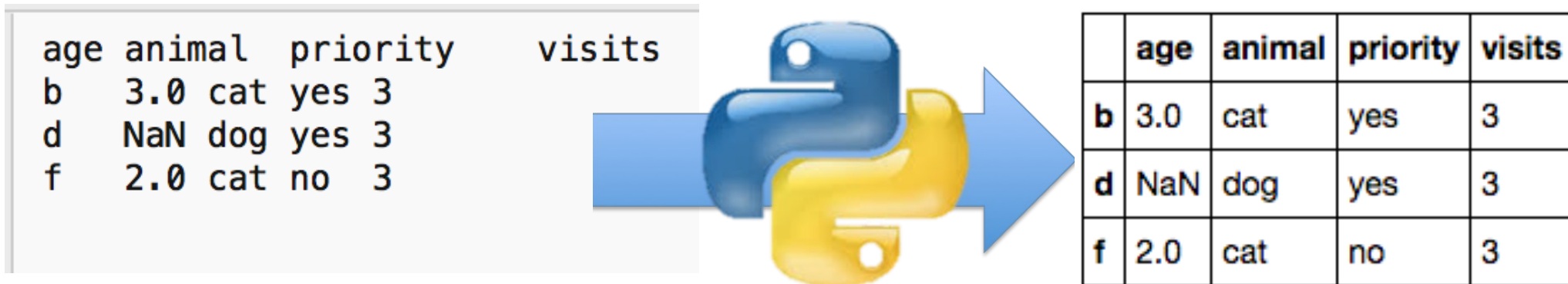
- Create a new Python3 Jupyter Notebook, save it to the folder */Bio334/02_python_introduction/exercises* and start exploring
- Change the name of your newly created file by right clicking on *Untitled.ipynb*
- To execute a cell press Shift + Enter



optional exercise slides follow

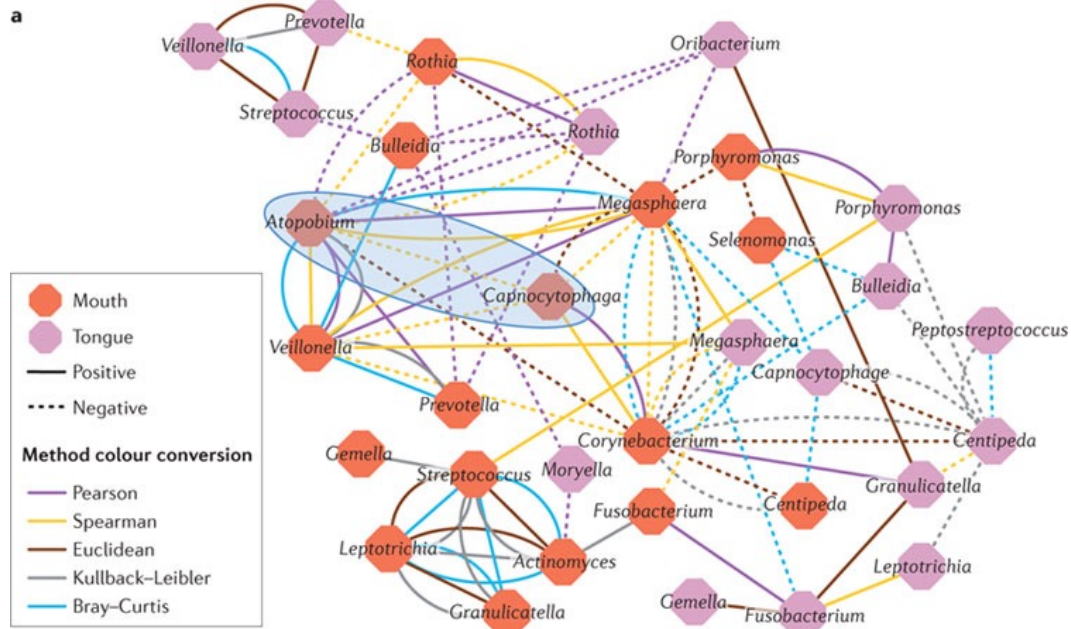
Optional Exercise #3

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

```
1 import os
2
3 dir_ = "Bio334"
4 if not os.path.exists(dir_):
5     print("It seems you haven't cloned the git repository yet. \
6         Please enter the following command in a terminal window.")
7     print("git clone https://github.com/meringlab/Bio334.git")
8 else:
9     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._
```