

Welcome to BINF 5007

Introduction to Python

Learning Objectives

1. Understand the Course: Summarize the course description, objectives, and key policies.



2. Explore Bioinformatics: Review published research and the importance of programming in bioinformatics.



3. Differentiate Roles: Distinguish between users (analysts) and programmers (developers).



4. Learn Python Basics: Explain Python's relevance for biologists, apply basic rules, and use the interactive shell.



5. Set Up Tools: Install PyCharm and understand its use for Python programming.



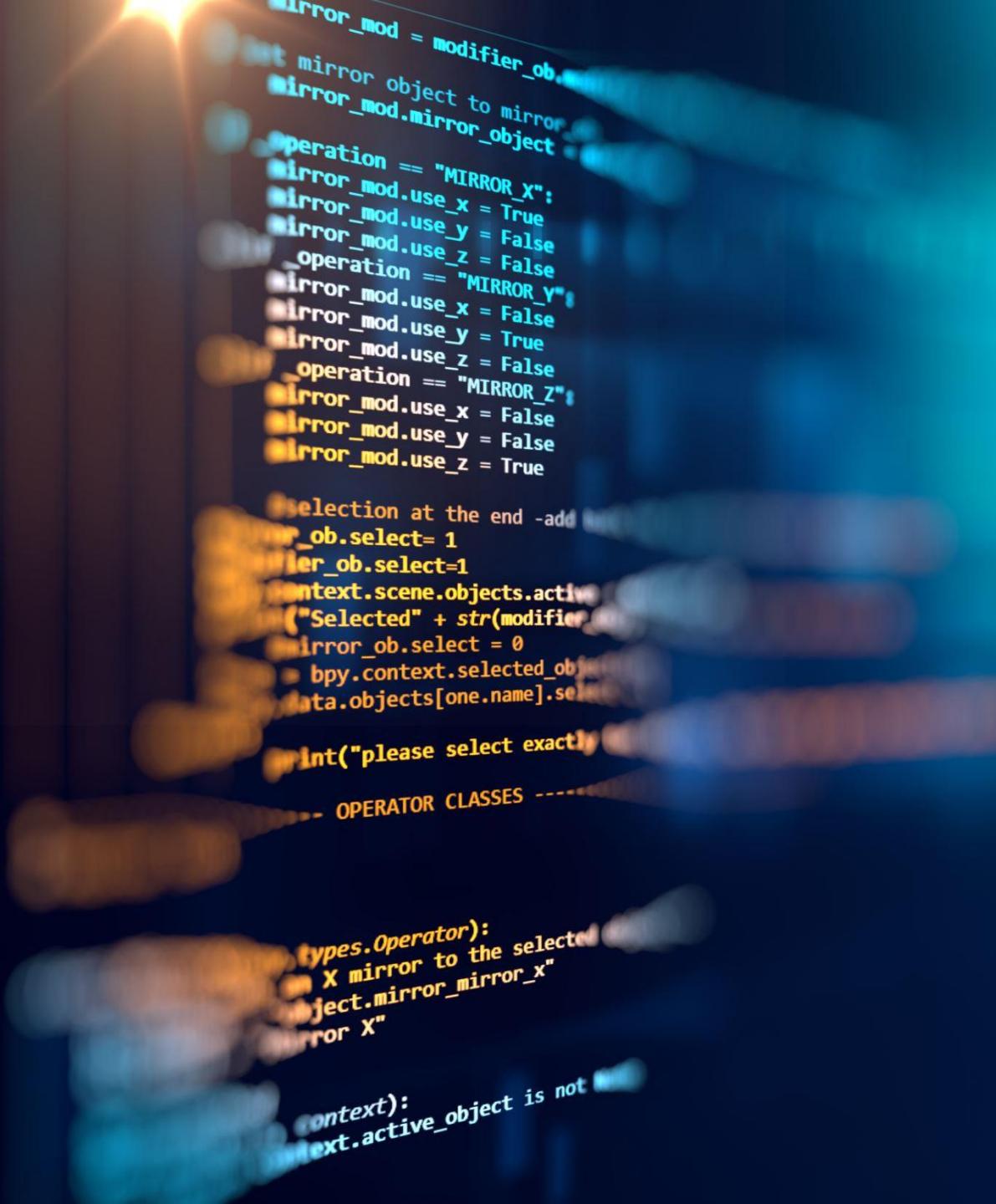
Course Description

Focuses on the fundamental programming skills required in the bioinformatics industry. Python is the main programming language used. Topics include string operations, file manipulation, regular expressions, Data structures. Includes substantial out-of-classroom assignments.

In this course, we'll concentrate on the essential coding abilities vital for a career in bioinformatics. Effective program design stands out as a crucial skill. Python, being an advanced language, offers numerous advantages, especially in bioinformatics. This allows students to immediately create effective programs tailored for their biological data and research needs.

Course Objectives

- Understand Python Syntax
- Write Python Functions
- Write Abstract Python Functions and How to Refactor Python Code
- Implement Type Hinting
- Effectively Document Code
- Properly Handle Exceptions
- Leverage Advanced Data Structures Pandas
- Implement Python Modules and Packages
- Parse Text Files and Using Regular Expressions
- Manipulate a Wide Range of Data Formats (TSV, CSV, FASTA, VCF, XML, JSON)

A photograph of a person's hand pointing towards a computer monitor. The monitor displays a block of Python code. The code appears to be a script for a 3D modeling application, likely Blender, based on the bpy context and object manipulation. The background is dark, and the screen is brightly lit, creating a glow around the text.

```
mirror_mod = modifier_obj
# Set mirror object to mirror
mirror_mod.mirror_object = selected_object
if operation == "MIRROR_X":
    mirror_mod.use_x = True
    mirror_mod.use_y = False
    mirror_mod.use_z = False
elif operation == "MIRROR_Y":
    mirror_mod.use_x = False
    mirror_mod.use_y = True
    mirror_mod.use_z = False
else:
    mirror_mod.use_x = False
    mirror_mod.use_y = False
    mirror_mod.use_z = True

# Selection at the end - add
modifier.select = 1
selected_object.select = 1
context.scene.objects.active = selected_object
("Selected" + str(modifier))
selected_object.select = 0
bpy.context.selected_objects.append(selected_object)
data.objects[one.name].select = 1
print("please select exactly one object")
print("operator classes")

# types.Operator:
#     X mirror to the selected object.mirror_mirror_x
#     mirror X"
#     context:
#         context.active_object is not None
```

Code Policies

We take the code policy very seriously

- Do not copy code!
 - We have ways to check for duplication
 - It is better to learn for yourself than risk getting a 0
 - We encourage working together but code should be *different*
- Programming style requirements:
 - PEP8 style Guide for Python Code
 - <https://www.python.org/dev/peps/pep-0008/>
- Each assignment will specify whether
 - You may use code from other sources
 - If you can use non-standard Python Modules, i.e. do not unless you have been allowed to

Policies

- Communication Policies: For reaching out, please use **email**.
- Policies Regarding Submitting Projects:
 - Projects submitted after the deadline will lose 5% each day for up to five days (maximum penalty of 25%).
 - After five days, the work will be given a zero.
 - Please read more : <https://healthsciences.humber.ca/current-students/resources/fhsw-student-handbook.html?page=Policies%20%26%20Processes&tab=fhswpolicyCategory1>
- Code policies – this is a good slide to introduce how you will be handling AI in the classroom

Why Python for Bioinformatics? What can it do?

Quantitative Modeling of COVID-19 Vaccines: A Bioinformatics Approach



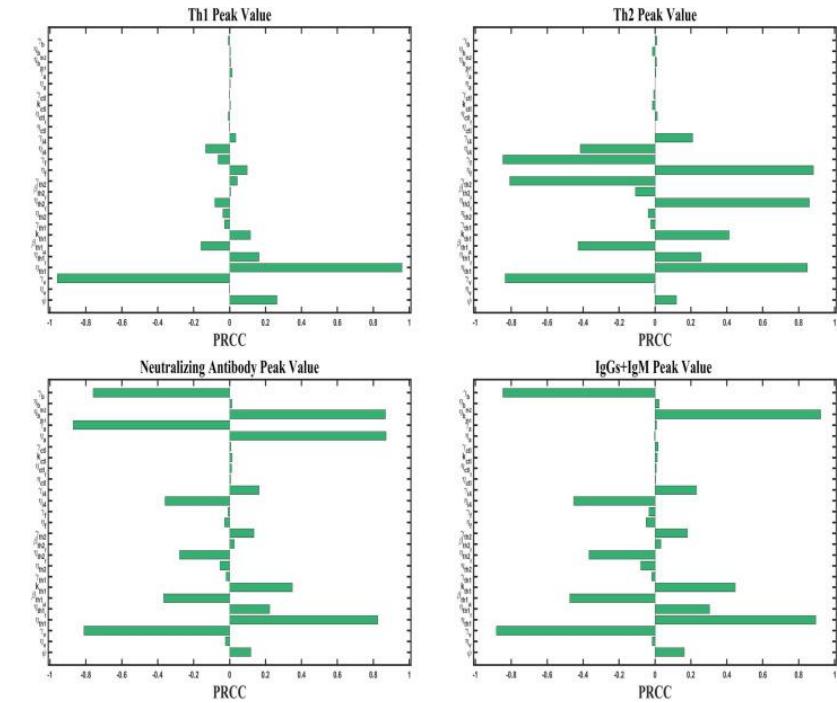
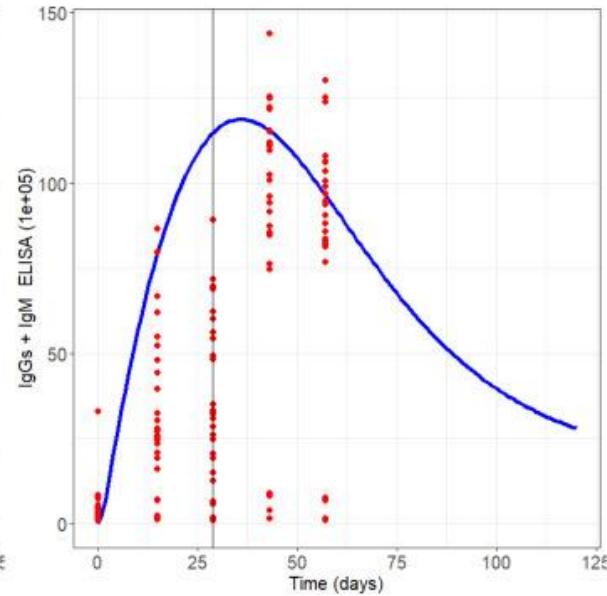
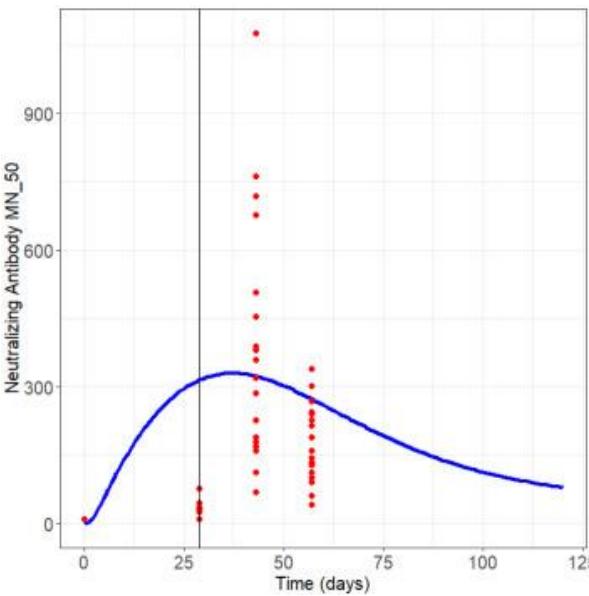
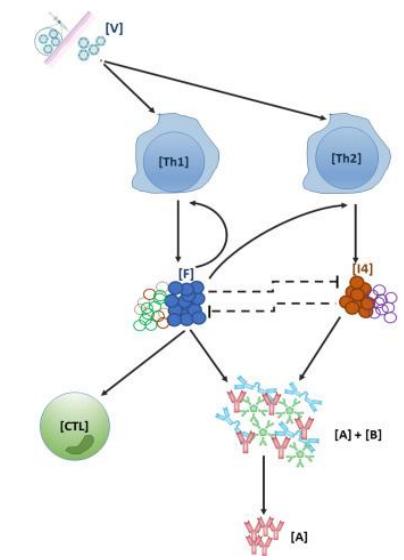
Mathematical Biosciences
Volume 358, April 2023, 108970



Original Research Article

A mathematical model of protein subunits COVID-19 vaccines

Samaneh Gholami ^a , Chapin S. Korosec ^a, Suzan Farhang-Sardroodi ^{b,a}, David W. Dick ^a, Morgan Craig ^c, Mohammad Sajjad Ghaemi ^d, Hsu Kiang Ooi ^d, Jane M. Heffernan ^a

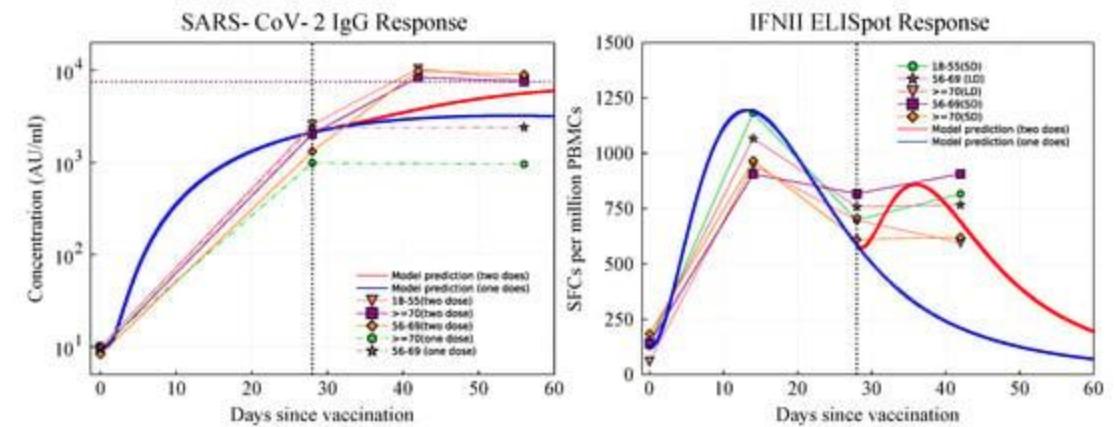
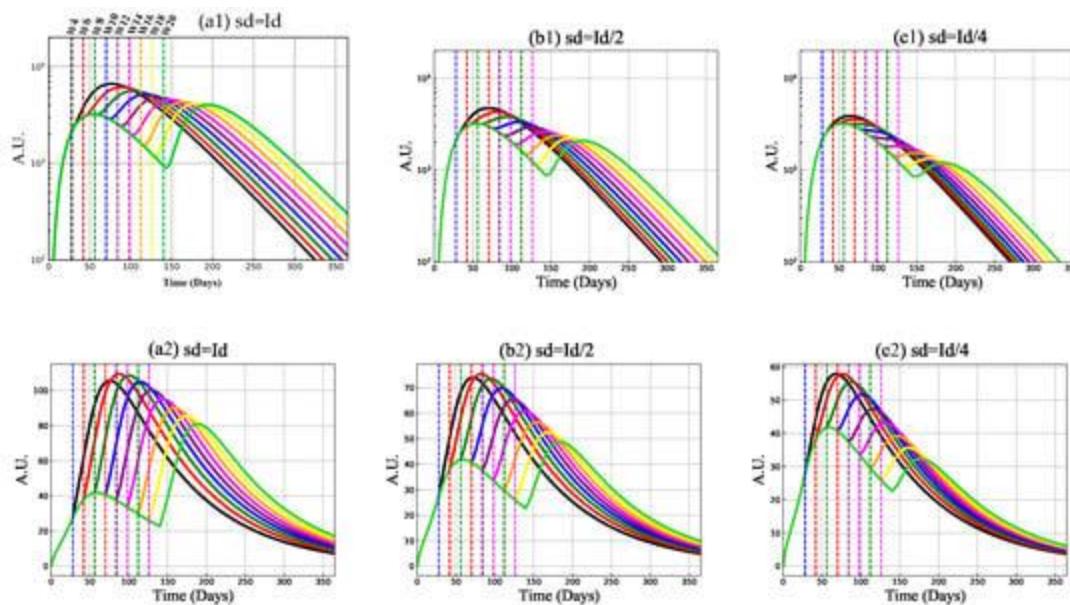


- <https://www.sciencedirect.com/science/article/pii/S0025556423000111>

A Bioinformatics Method for Modeling the AstraZeneca COVID-19 Vaccine

Analysis of Host Immunological Response of Adenovirus-Based COVID-19 Vaccines

by Suzan Farhang-Sardroodi ^{1,2,*}    , Chapin S. Korosec ^{1,2}    , Samaneh Gholami ^{1,2}   , Morgan Craig ³    , Iain R. Moyles ²    , Mohammad Sajjad Ghaemi ⁴    , Hsu Kiang Ooi ⁴    and Jane M. Heffernan ^{1,2,*}   

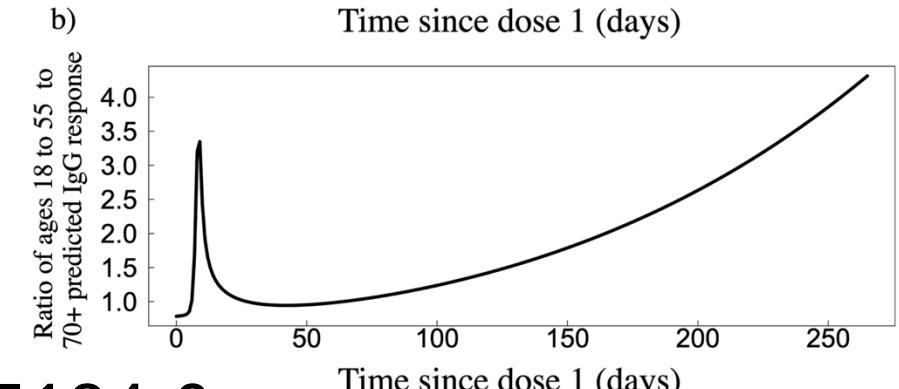
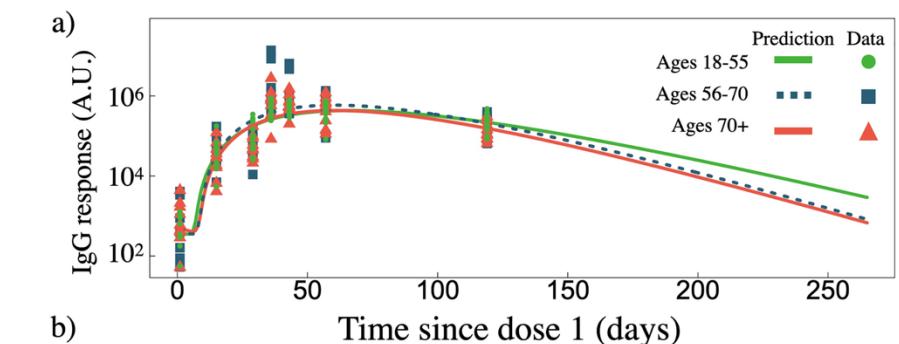
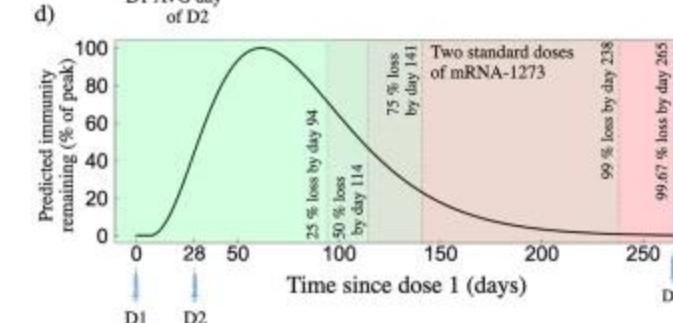
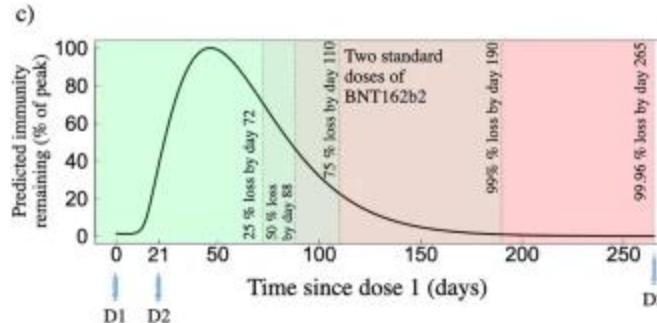
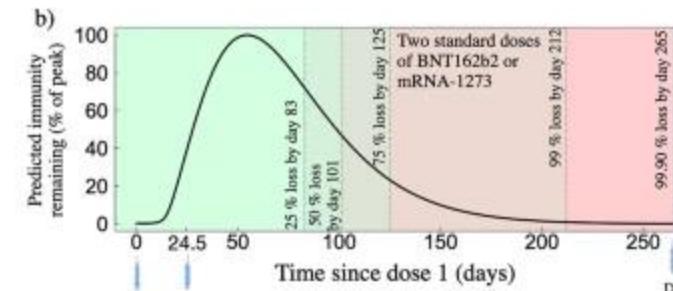
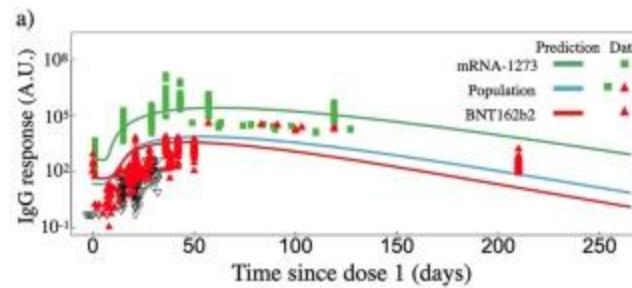


- <https://www.mdpi.com/2076-393X/9/8/861>

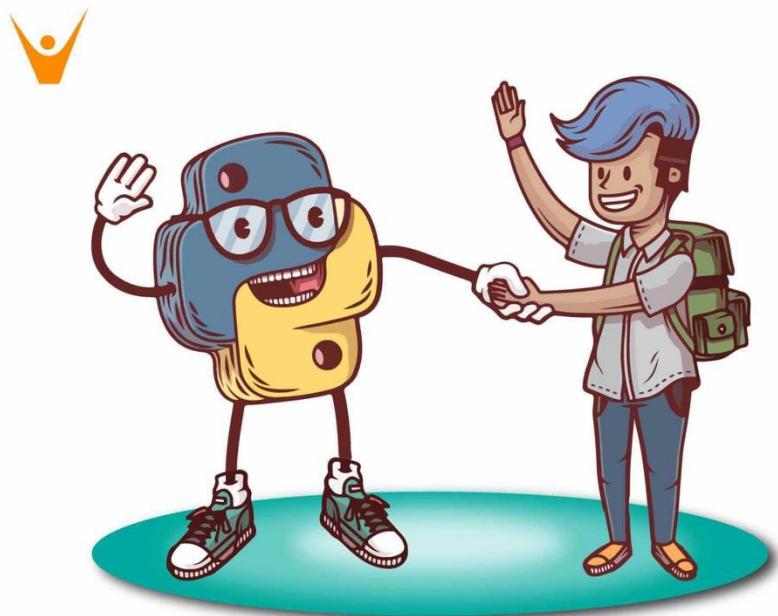
Article | [Open access](#) | Published: 08 December 2022

Long-term durability of immune responses to the BNT162b2 and mRNA-1273 vaccines based on dosage, age and sex

[Chapin S. Korosec](#)  , [Suzan Farhang-Sardroodi](#), [David W. Dick](#), [Sameneh Gholami](#), [Mohammad Sajjad Ghaemi](#), [Iain R. Moyles](#), [Morgan Craig](#), [Hsu Kiang Ooi](#) & [Jane M. Heffernan](#) 



Let's Get Started



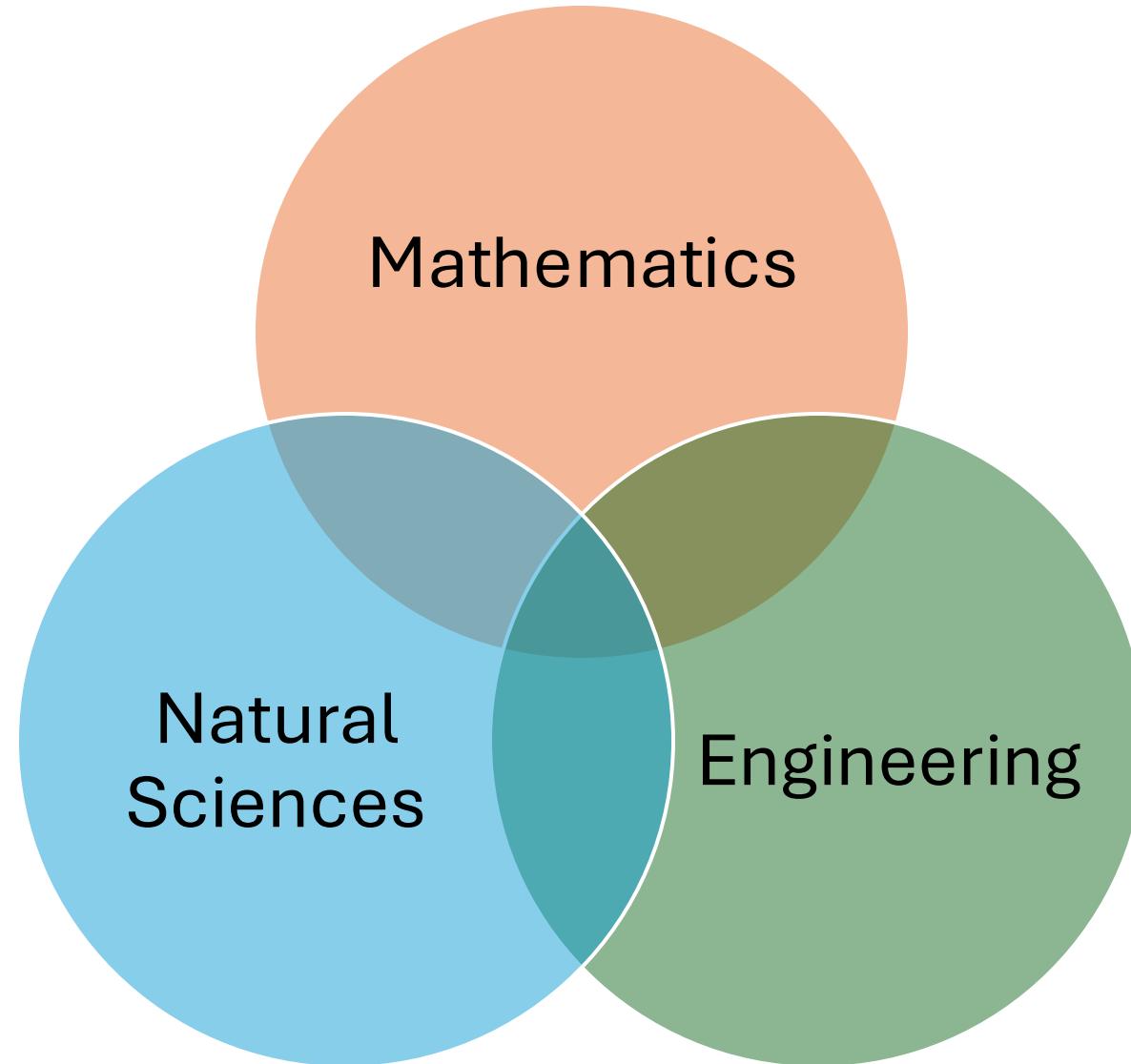
Summary

- A quick review of some aspects of Python
- We will revisit some of these concepts throughout the course. Especially the idea of Mutable / Immutable
- Remember to focus on the big picture
- Go through and implement the slides that have "Implement in PyCharm" on your own. Do this, so you can begin to practice

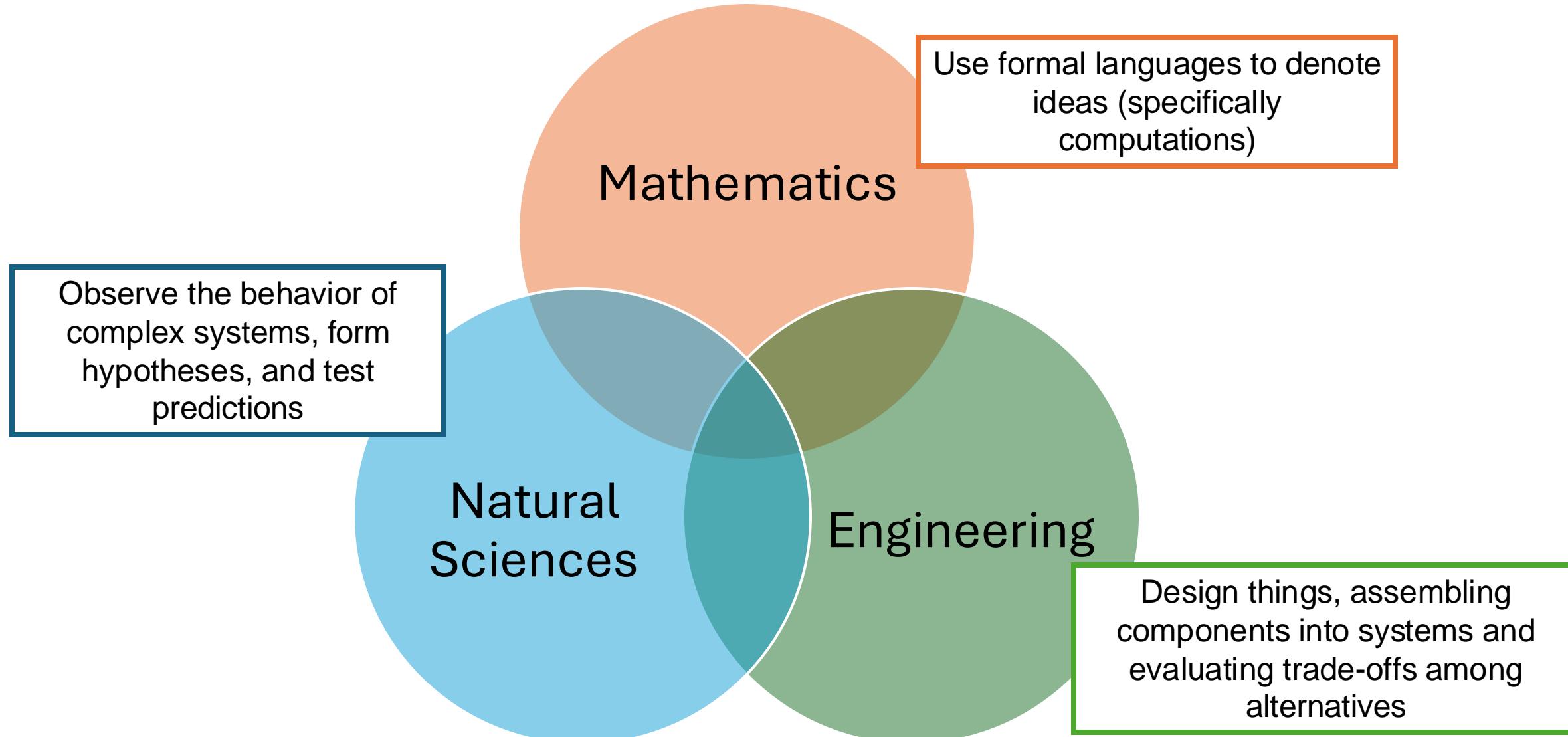
Why Learn to Program?

- I want to use programs to analyze the data I run in my lab experiments in my research
- I want to compile the data from the programs I run my lab
- I want to automate tedious tasks that I do over and over
- I found out I like the challenge of solving problems with programming
- I need to be able to communicate with other people about programming

Thinking like a Computer Scientist



Thinking like a Computer Scientist



Computers Want to be Helpful...

- Computers are built for one purpose - to do things for us
- But we need to speak their language to describe what we want done
- Users have it easy - someone already put many different programs (instructions) into the computer and users just pick the ones they want to use



What
Next?

What
Next?

What
Next?

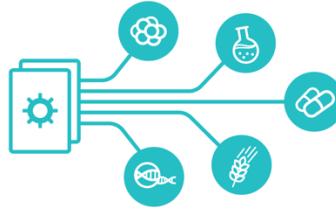
What
Next?

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Next?

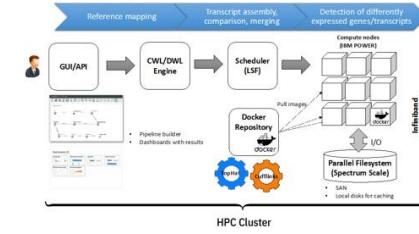
What
Next?

Users vs. Programmers

- Users see computers as a set of tools - word processor, spreadsheet, map, to-do list, etc.
- Programmers learn the computer "ways" and the computer language
- Programmers have some tools that allow them to build new tools
- Programmers sometimes write tools for lots of users and sometimes programmers write little "helpers" for themselves to automate a task



User



Computer
Hardware + Software

Programmers
(us)

Data

Information

....

Networks

From a software creator's point of view, we build the software. The end users (stakeholders/actors) tell us what they would like the software to do. We the programmers, develop software and get paid in exchange for good work. However, the data, information, and networks are our problem to solve on their behalf. The hardware and software are our friends and allies in this quest.

So what is code?

To some it looks like this:



```
for i in people.data.users:
    response = client.api.statuses.user_timeline.get(screen_name=i.screen_name)
    print 'Got', len(response.data), 'tweets from', i.screen_name
    if len(response.data) != 0:
        ldate = response.data[0]['created_at']
        ldate2 = datetime.strptime(ldate, '%a %b %d %H:%M:%S +0000 %Y')
        today = datetime.now()
        howlong = (today-ldate2).days
        if howlong < daywindow:
            print i.screen_name, howlong
            totaltweets += 1
            for j in response.data:
                if j.entities['urls']:
                    for l in j.entities['urls']:
                        if l['expanded_url'].startswith('http://t.co'):
                            print l['expanded_url']
                else:
                    print i.screen_name, j['text']
```

A photograph of a person with light brown hair, wearing a white shirt, sleeping with their head resting on a keyboard. A laptop screen is visible in the foreground, showing a dark interface with some text or code. The background is slightly blurred, suggesting an indoor office or study environment.

What is Code?

```
name = input('Enter file: ')
handle = open(name, 'r')
```

```
counts = dict()
for line in handle:
    words = line.split()
    for word in words:
        counts[word] = counts.get(word, 0) + 1
```

```
bigcount = None
bigword = None
for word, count in counts.items():
    if bigcount is None or count > bigcount:
        bigword = word
        bigcount = count
print(bigword, bigcount)
```

```
python3 words.py
Enter file: words.txt
to 16
```

```
python3 words.py
Enter file: clown.txt
the 7
```

A lot going on here! Much to discuss!

<https://www.py4e.com/code3/words.txt>

<https://www.py4e.com/code3/clown.txt>

What is Clinical Bioinformatics?

- Clinical bioinformatics is a subfield of bioinformatics that involves analyzing and interpreting 'omics data to solve biological problems with implications in health and disease.
- We can assess risk, improve diagnostic tools, uncover biomarkers for disease, as well as inform treatment approaches through targeted drug therapies.



In Clinical Bioinformatics, we can use Python to...



Handle large clinical datasets with data from multiple samples

Pandas and NumPy - data mining, organizing and cleaning



Write scripts to filter and annotate genetic variants in order to flag variations as potentially pathogenic and linked to disease

PyVCF - Library for parsing variant call files



Develop machine learning models to predict disease risk and clinical outcomes

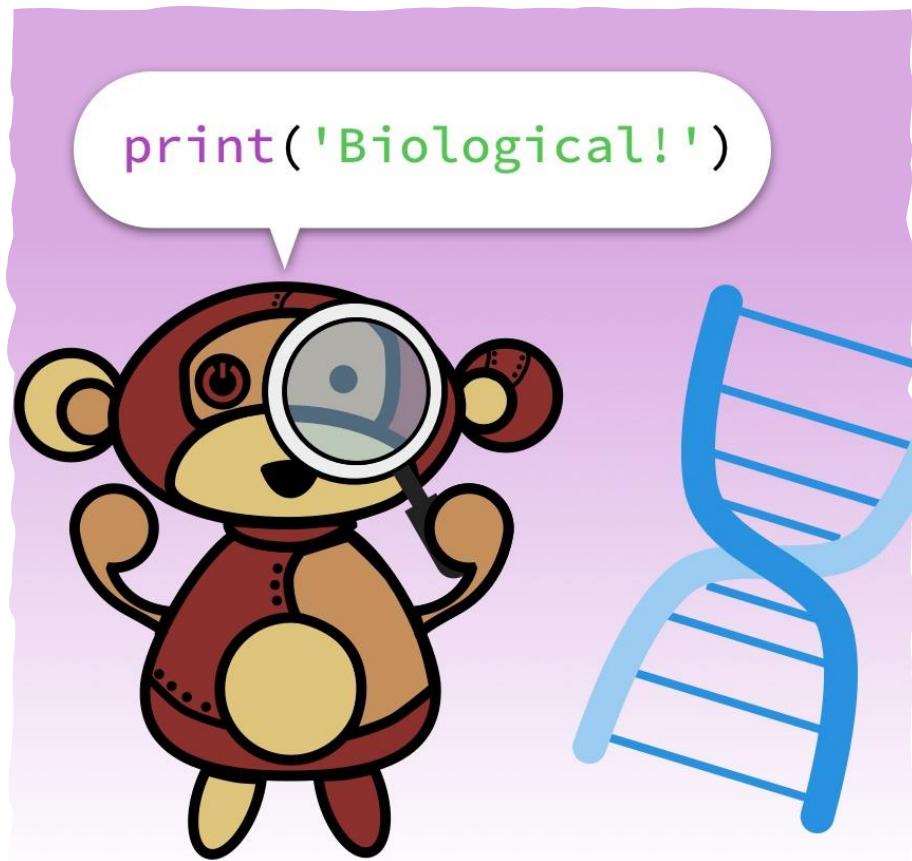
SciKit Learn - ML tasks



Visualize patterns of gene expression to identify differences between conditions (i.e. healthy and diseased cells) for identifying potential drug targets

Matplotlib
Seaborn

Why Python for a Biologist?



- Versatile language that can make our work easier with tools to explore our complex (and messy!) datasets as well as automate repetitive processes
- Allows us to create a collection of small and functional scripts we can reuse for reoccurring tasks
- Useful features for easy text manipulation, pattern recognition and data integration that are helpful for working with sequencing data
- Can integrate with different software tools, process their outputs, and move data smoothly to create multi-stage analysis workflows for complex tasks

Guido van Rossum, Creator Python

- Born in the Netherlands
- Began to develop Python in 1989
- Has worked for Google, DARPA, Dropbox at present
- Named Python after *Monty Python's Flying Circus*
- Released Python 1.0 in 1994



Python Philosophy

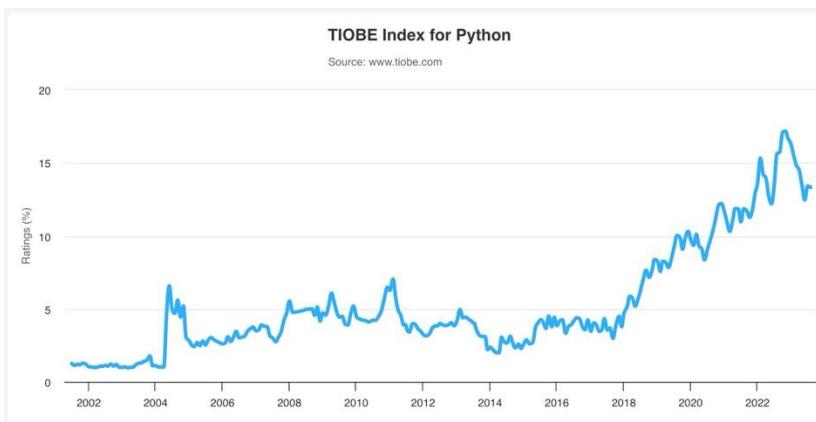
- The Zen of Python is a collection of 19 "guiding principles" for writing computer programs
- The language's core philosophy is summarized in those documents
- Includes set of principles, like:
 - Beautiful is better than ugly
 - Simple is better than complex
 - Complex is better than complicated
 - Readability counts
 - **Explicit is better than implicit**
- Rather than having all its functionality built into its core, Python was designed to be highly extensible
- This compact modularity has made it particularly popular as a means of adding programmable interfaces to existing applications

Python is meant to be an easily readable language...

- Strives for a simpler, less-cluttered syntax and grammar
- Still gives developers a choice in their coding methodology
- In contrast to Perl's "there is more than one way to do it" motto
 - Python embraces a "***there should be one—and preferably only one—obvious way to do it***" design philosophy
- Alex Martelli, a Fellow at the Python Software Foundation and Python book author, writes that "To describe something as 'clever' is not considered a compliment in the Python culture."
- "**Pythonic: ways..." can sometime lead to hard to read code**

Python and Programming Paradigms

- Python supports several programming paradigms
- **It fully supports structured programming and object oriented**
- It also supports various concepts in **functional** programming
- Python features a **dynamic type** system and automatic memory management
- The programming paradigms and language features help you to develop large and complex software applications with Python
- **This has led to widespread adoption**
- **See the TIOBE index**



Rank	Language	Icon	Name	Growth	Change
1	1	Python	Python	13.33%	-2.30%
2	2	C	C	11.41%	-3.35%
3	4	C++	C++	10.63%	+0.49%
4	3	Java	Java	10.33%	-2.14%
5	5	C#	C#	7.04%	+1.64%
6	8	JavaScript	JavaScript	3.29%	+0.89%
7	6	Visual Basic	Visual Basic	2.63%	-2.26%
8	9	SQL	SQL	1.53%	-0.14%
9	7	Assembly language	Assembly language	1.34%	-1.41%
10	10	PHP	PHP	1.27%	-0.09%
11	21	Scratch	Scratch	1.22%	+0.63%
12	15	Go	Go	1.16%	+0.20%
13	17	MATLAB	MATLAB	1.05%	+0.17%
14	18	Fortran	Fortran	1.03%	+0.24%
15	31	COBOL	COBOL	0.96%	+0.59%
16	16	R	R	0.92%	+0.01%
17	19	Ruby	Ruby	0.91%	+0.18%
18	11	Swift	Swift	0.90%	-0.35%
19	22	Rust	Rust	0.89%	+0.32%
20	28	Julia	Julia	0.85%	+0.41%

The TIOBE Programming Community index is an indicator of the popularity of programming languages
Language of the Year: 2007, 2010, 2018, 2020, 2021
<https://www.tiobe.com/tiobe-index/>

Examples of When is Whitespace Ignored:

```
# whitespace.py

long_winded_computation = (1 + 2 + 3 + 4 + 5 + 6 + 7 + 8 +
                            9 + 10 + 11 + 12 + 13 + 14 +
                            15 + 16 + 17 + 18 + 19 + 20)
# Alternatively:
long_winded_computation = 1 + 2 + 3 + 4 + 5 + 6 + 7 + 8 + \
                           9 + 10 + 11 + 12 + 13 + 14 + \
                           15 + 16 + 17 + 18 + 19 + 20

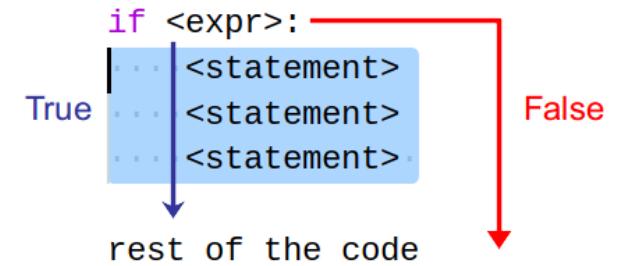
# lists
list_of_lists = [[1, 2, 3], [4, 5, 6], [7, 8, 9]]

easier_to_read_list_of_lists = [[1, 2, 3],
                                 [4, 5, 6],
                                 [7, 8, 9]]


print(type(long_winded_computation))          <class 'int'>
print(type(list_of_lists))                     <class 'list'> What's being printed
print(type(easier_to_read_list_of_lists))       <class 'list'>
```

Indentation

- Increase indent after an `if` statement or `for` statement (after `:`)
- Maintain indent to indicate the block (which lines are affected by the `if/for`)
- Reduce indent back to the level of the `if` statement or `for` statement to indicate the end of the block
- Blank lines are ignored - they do not affect indentation
- Comments on a line by themselves are ignored with regard to indentation



```
# Python program to explain continue statement
string1 = "Steckies"

# Continue with for loop
for value in string1:
    if value == 'e':
        # If Letter is equal to 'e' next statement will skip
        # Skip statement after continue
        print('This is continue block: ',value)
    # Print the next iteration
    print("Value: ",value)

# Statement outside the for loop
print("Outside for Loop")
```

Warning: Turn Off Tabs!!

- Some editors automatically uses spaces for files with ".py" extension (nice!)
- Most text editors can turn tabs into spaces - make sure to enable this feature
- Python cares a *lot* about how far a line is indented. If you mix tabs and spaces, you may get "indentation errors" even if everything looks fine, i.e. you won't see them
- PyCharm convert tabs to spaces automatically

<https://www.youtube.com/watch?v=wYQxx2ewrbA>

Early Learner: Syntax Errors

- We need to learn the Python language to communicate our instructions. We will make lots of mistakes in the beginning and speak gibberish like *someone learning a new language*
- When you make a mistake, the computer says "syntax error" – It will seem like Python is cruel
- Must remember that you are intelligent one in the relationship. The computer is simple and very fast, but cannot *learn*
- You don't need to understand every detail of Python to be productive
- The Principles we cover throughout this semester are vital to your growth as a programmer
- **But remember: Mastery comes with practice of reading and writing code**

A Simple Task to Handle with Python

- Question:
 - What 6-mer containing only A's and T's occurs mostly commonly in the *P. falciparum* genome
- Answer:
 - Scan through the 2.2 million nucleotides of the 14 chromosomes using a 6-nt window and find that ATATAT occurs most frequently

```
-----  
A: 40.31 %  
T: 40.30 %  
C: 9.70 %  
G: 9.69 %  
total(ATCG only ) = 22853497  
nucleotides  
total(everything) = 22853764  
nucleotides  
-----
```

1	ATATAT	570577
2	TATATA	529826
3	AAAAAA	420232
4	TTTTTT	417439
5	TATTAT	147506
6	ATAATA	146948
7	TATATT	146711
8	AATATA	146695

-- snip --

57	ATTTAA	50617
58	ATTAAT	49856
59	TTAAAT	49731
60	TAATTA	45781
61	AATTAA	45278
62	TTAATT	45212
63	AATTTA	45101
64	TAAATT	44553

The Process of Programming

- In scripted languages, you can write it with ANY text editor
 - **Sublime is good, but PyCharm is superior**
- As you can imagine, programming is a **process**:
 - Write code
 - Try to run the program
 - Correct the errors (debugging)
 - Look at the output
 - **Compare previous results (use your `diff` or `sdiff`)**
 - Edit the program and try again
- **If you want to program as a profession, learn to love the process. You will be required over and over and over**
- **To write better code, we'll learn more about this process over the semester**

Getting python3

- If you need to get Python 3, it's straight forward. Download and follow instructions

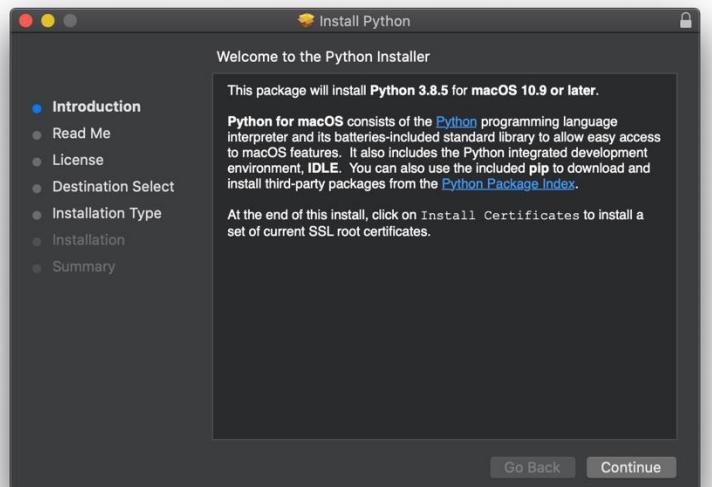
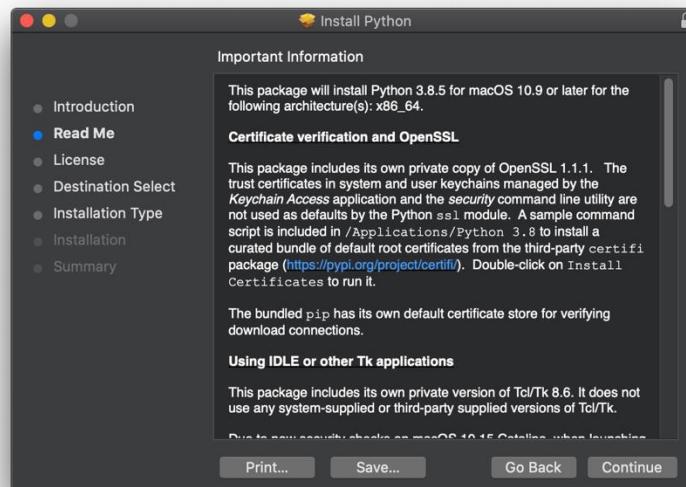
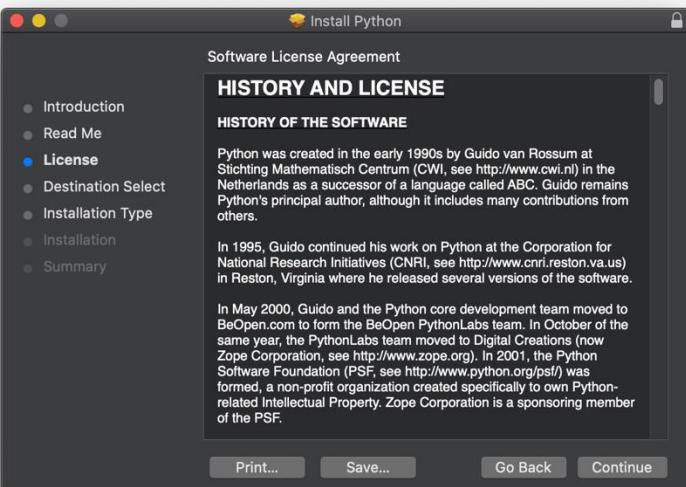
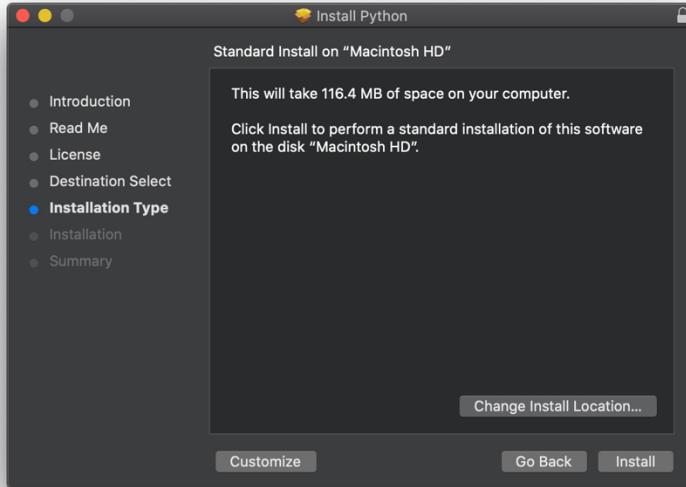
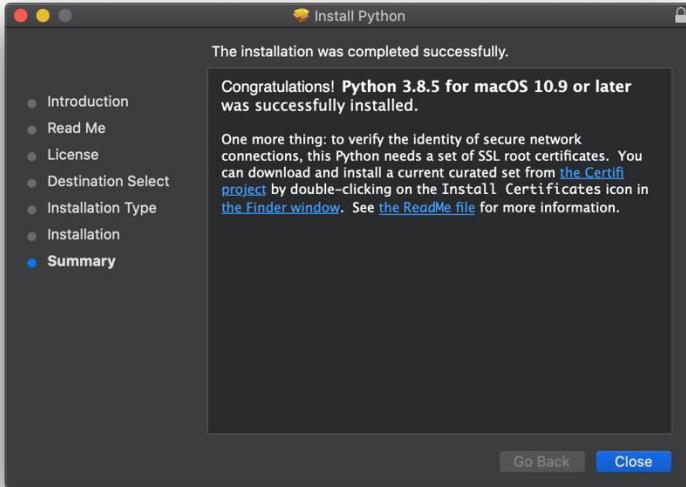
<https://www.python.org/ftp/python/3.11.5/python-3.11.5-macos11.pkg>



Mac Users link above

<https://www.python.org/downloads/>

Getting python3



Where Do Site Packages Go?

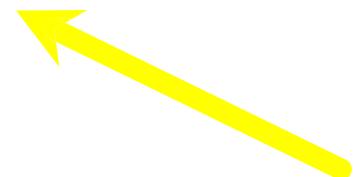
```
$ pip show numpy
Name: numpy
Version: 1.23.5
Summary: NumPy is the fundamental package for array computing with Python.
Home-page: https://www.numpy.org
Author: Travis E. Oliphant et al.
Author-email:
License: BSD
Location: /opt/homebrew/anaconda3/lib/python3.10/site-packages
Requires:
Required-by: astropy, bokeh, Bottleneck, contourpy, datashader, datashape, gensim, h5py,
holoviews, hvplot, ifCNV, imagecodecs, imageio, imbalanced-learn, matplotlib, numba,
numexpr, pandas, patsy, pyerfa, PyWavelets, scikit-image, scikit-learn, scipy, seaborn,
statsmodels, tables, tifffile, transformers, xarray
```

Note: I have Python 3.9 installed, and have installed `numpy`

Python Interactive Shell

- One of the best features of python is the REPL: is an **interactive** way to use Python
- The easiest way to open a Python interactive **shell** is to type `python3` on CLI
- To quit, `exit()`, `quit()` or press Control+D
- The REPL is useful for:
 - Experimenting with short expressions or statements to learn or test new features of the language
 - See how assignments work (not assignments you'll do in class)
 - Consulting the documentation
 - Etc.

```
[~]# python3  
Python 3.9.10 (main, Jan 15 2022, 11:48:00)  
[Clang 13.0.0 (clang-1300.0.29.3)] on darwin  
Type "help", "copyright", "credits" or "license" for more information.  
>>>
```



What next?

Note: if you install Python 3.9.10 yours will look like this

```
[~]# python3
Python 3.9.10 (main, Jan 15 2022, 11:48:00)
[Clang 13.0.0 (clang-1300.0.29.3)] on darwin
Type "help", "copyright", "credits" or "license" for more information.
>>> x = 1
>>> print(x)
1
>>> x = x + 1
>>> print(x)
2
>>> exit()
```

This is a good test to make sure that you have Python correctly installed. Note that `quit()` also works to end the interactive session.