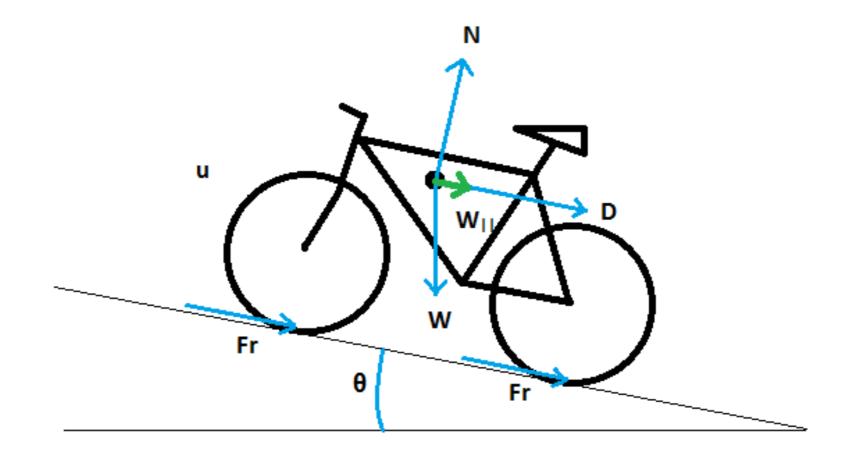
Programming with Python

Duke UPGG Scientific Computing Bootcamp August 12, 2019 Dan Leehr <u>dan.leehr@duke.edu</u>



What book should I read?

How many books about riding a bike did you read?



"You can be a scientist in the science of bike ride mechanics and it still won't help you one bit to do the actual thing."

http://twonontechies.com/bicycles-can-help-you-learn-programming/

Why Python?

- We have to use *something*
- It's free, well-documented, and runs everywhere
- Large community among scientists
- Relatively easy to pick up, but programming is hard!

Goals

- Write and run programs in Python
- Understand basic data types and functions
- Work with files and libraries
- Know where to look for more help

I know, I'll use **Python**!

Download

- Download the python-fasta.zip file from the course website - Syllabus.
- Unzip it and place on your Desktop:

```
python-fasta/
  ae.fa
  ls_orchid.fasta
```

1. Open **Anaconda Navigator** (installed with Anaconda)

2. Click to launch Jupyter Notebook

• • •	🔾 Ana	conda Navigator					
	NDA NAVIGATOR	() Upgrad	de Now Sign in to Anaconda Cloud				
A Home	Applications on base (root)	~ Channels	Refresh				
The Environments	\$	\$	^				
🗳 Learning	lab	jupyter		localhost:8888/tree	C	0	
	jupyterlab	notebook	0	Home		•	
Community	↗ 0.32.1	⊅ 5.5.0		Home		0.1] [
	An extensible environment for interactive and reproducible computing, based on the	Web-based, interactive computing notebook environment. Edit and run human-readable	💭 Jupyter			Quit	Logout
	Jupyter Notebook and Architecture.	docs while describing the data analysis.	Files Running Clusters				
Documentation							
	Launch	Launch	Select items to perform actions on them.			Upload	New - 2
Developer Blog					Name 🕹	Last Modified	File size
Feedback	¢	\$	anaconda3			a year ago)
	IP[y]:					a year ago)
			C Applications			3 hours ago	,
			D D bin			4 days ago)
			Box Sync			2 years ago	2
			capture			2 years ago	>
			Code			4 months ago	>
			C Cookied_conf			4 years ago	,
			Data			2 months ago	>
			C dc_workshop			3 days ago	>
						2 months ago	
			Desktop			2 minutes ago	
						2 years age	

Begin Jupyter Notebook

Data Types

- Numeric:
 - Integer: 1, 76, 400
 - Float: -1.2, 0.5, 3.1415926 (Use a decimal point)
 - Boolean: True, False
- Text:
 - Strings: 'ACTGACAG' (Wrap in quotes)

Strings

• Strings can be created with quotes or double quotes:

```
name = 'Daniel'
```

• Access individual letters as strings with [] (starting at 0)

name[0] # D name[1] # a

• Check if a letter exists in a string

```
'a' in name # True
'a' not in name # False
```

Variables

• Assign variables with equals

x = 3

Access variables by name

print x # 3

 Variables work like sticky notes, they're just a label on top of a value

What do we know?

- Our sequence is a string, in **seq10**
- Strings are sequences of characters, each at a numbered position (starting from 0)
- We can extract characters as strings with square brackets []
- We can combine strings together with +

Exercise: Reverse

- Write some code that **reverses** the sequence in seq.
- It should
 - 1. Create an empty string variable **rev**

rev = ''

- 2. Loop over the items in **seq**, adding these to rev in reversed order
- 3. Print the contents of **rev**

Loops

• Write a loop with **for** item **in** collection:

for letter in word: print letter

• Always put a colon at the end of the line, indented lines are run for every item in the collection

Complementing

- We can loop over all the bases in a sequence
- Each base has a complement that we should substitute:
- We can use a **Dictionary** to store this mapping.

А	→	Т
С	\rightarrow	G
Т	→	А
G	\rightarrow	С

Dictionaries and Lists

• Create dicts with {}, lists with []

nucs = {'A': 5, 'C': 4, 'T': 8}
counts = [5,4,8]

Both accessed with [] - dicts by key, lists by index

nucs['A'] # 5
counts[0] # 5

nucs['A'] = 3 # now 3counts[0] = 3 # now 3

GC-content percentage

- Calculated as (G + C) / (A + T + G + C)
- Create a GC count variable and an ATGC count variable
 - Loop over each base in the sequence
 - If G, add 1 to GC count
 - If C add 1 to GC count
 - For everything, add 1 to ATGC count

Conditionals

Test c1 for True or False
if c1:

print "c1 was True"
c1 was False, check c2
elif c2:
 print "c1 False but c2 True"
All checks False
else:

print "Both False"

Exercise: Functions

bases = 'adenine cytosine guanine thymine'

Write some code that:

- Makes a **list** of these bases from the string
- **Uppercases** the names (e.g. ['ADENINE', ...])
- **Reverse**s the order (e.g. ['THYMINE',...])

Hint: Use help(str) and help(list) to see what functions are available for strings and lists

Bonus: Write a for loop to print the first letter of each (e.g. A, C, ...)

Exercise

Strings can be reversed with this special slicing notation:
 [::-1]

```
s = 'abc'
r = s[::-1]
print(r)
```

cba

- Update reverse() function to use [::-1] instead of a loop.
- Do we need to do anything to complement()?
 What about reverse_complement()?

Functions

- Calling functions: length = len('abc')
- Defining functions:

def double(x):
 return x * 2

• Composing functions:

def reverse_complement(seq):
 return reverse(complement(seq))

• Avoid using global variables in functions

Exercise

- Write a function, **read_fasta(filename)** that:
 - Takes 1 argument: filename
 - Reads the file line-by-line
 - Strips/combines the lines into one long line
 - Skips the line if it contains a >
- Hint: if not 'i' in 'team':

Reading files

• Open a file with the **open()** function:

f = open('ae.fa')

• Loop over lines, and strip() each one

```
for line in f:
    print line.strip()
```

Close with f.close()

Scripts

- Put code in a file, give it the .py extension
- Read command line-arguments from sys.argv:

```
import sys
print sys.argv[0]
print sys.argv[1]
```

```
$ python script.py hello
script.py
hello
```

• Check the length of sys.argv to be helpful!