

# Genome Sciences 373

# Genome Informatics

Quiz Section #1

March 28, 2017

# About me

**Email:** cnoecker@uw.edu

**Office hours:** Mondays, 4:30-5:30 PM

Foege S-040

Or by appointment!

My research focuses on modeling to integrate different kinds of data from microbial communities

# Quiz section goals

- solidify in-class material
- develop understanding of programming concepts
- learn basic Python to write bioinformatics programs

*attendance is not required, but the material covered in section is required*

# **Homework policy**

No late homework accepted without **prior arrangements**

Group work, Internet searching: You can (and should) use them, but don't copy exactly! We can tell!

The point is **to learn**.

Grading is equally about your **effort** and your **execution**

# Homeworks continued

- First assignment will be assigned tomorrow via Catalyst
- Due Wednesday 4/5 before class (1:30PM)  
Start early!

## Questions:

Catalyst discussion board:

<https://catalyst.uw.edu/gopost/board/cnoecker/43929/>

Email, office hours

Questions about  
course logistics?

# Today's goals

- Quick review on alignments
- Algorithms and programs: what and why
- Getting started programming in Python

# What is an alignment?

Arrangement of nucleotide (or amino acid) sequences, to identify **regions of similarity** that may be a consequence of functional, structural, or evolutionary relationships between the sequences.

G - A A T T C A G T T A
G G - A - T C - G - - A

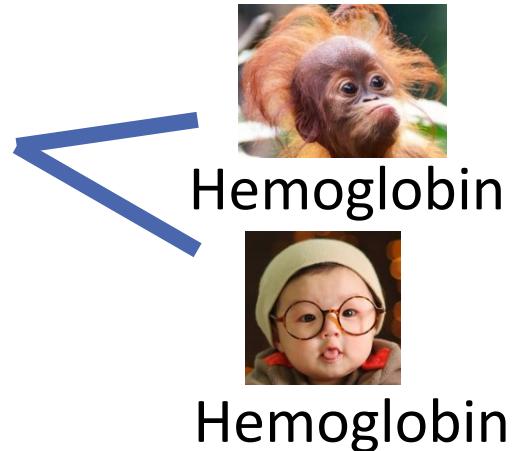
**What are some reasons to align sequences?**

# One big reason: compare *homologous* sequences

Sequences with shared ancestry

Orthologs

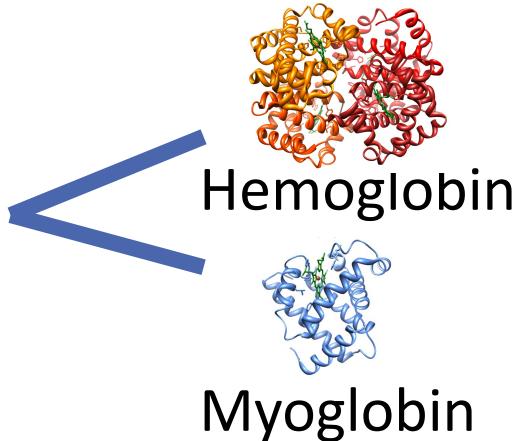
Common  
Ancestor



G - A A T T C A G T T A
G G - A - T C - G - - A

Paralogs

Common  
Ancestor



G - A A T T C A G T T A
G G - A - T C - G - - A

# Alignment example

Write down 2 possible alignments of the following two sequences:

ACCTTGT

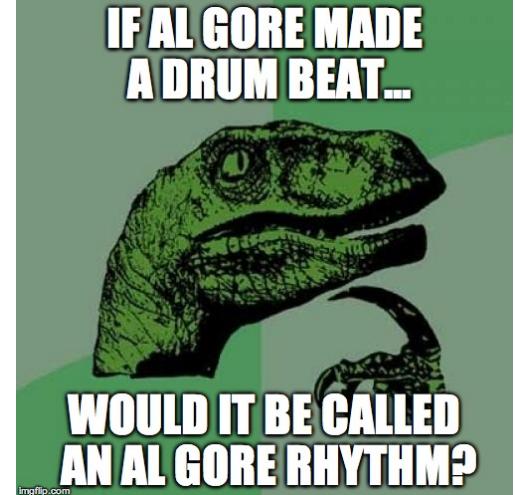
TCTGTCC

Which one is “better”? Does it depend what you’re trying to do?

# Algorithms

# What is an algorithm?

A step by step list of instructions that, if followed exactly, will solve the problem under consideration



The instructions can be carried out or *implemented* in different ways:

- Programmed to be run by a computer
- Carried out yourself

Like a recipe!

# Properties of algorithms

- Is an unambiguously defined series of steps
- Works for all inputs in a defined set
- Always produces a defined set of outputs, for those inputs
- Is guaranteed to produce a correct result, for those inputs

Often written in “pseudocode”

# Example algorithm: Find the smallest number

Input: three numbers A, B, and C

Output: the largest number

```
current_smallest <- A
if B < current_smallest:
    current_smallest <- B
else:
    [do nothing]
if C < current_smallest:
    current_smallest <- C
else:
    [do nothing]
return current_smallest
```

What set of inputs is this algorithm defined for?

# Which of these is an algorithm?

- Instructions for how to find the reverse complement of a DNA sequence
- A program that finds the reverse complement for any DNA sequence

# Programming with Python

# Why are we learning to program?

This class is designed for you to **understand** and **use** bioinformatics algorithms

You won't learn to **implement** all of them, but understanding them requires *programmatic thinking*

Plus, if you do want to implement an algorithm or otherwise code anything, you will be off to a good start!

# What is a program?

A series of instructions that performs a specific task when executed by a computer

## Why are programs useful?

# A note for those with programming experience

- Some of this will be review
- It's fine to use Python tricks and modules beyond what I show in quiz section
  - But please don't, for ex, use a BioPython function to do an entire homework problem in one command

# What is a program?

A series of instructions that performs a specific task when executed by a computer

subject      verb                            object

x = 4 #A line of code...

y = 8 #is like a sentence

z = x + y

print(z)

# Variables and operators

subject    verb                      object

The diagram illustrates the components of the assignment statement `x = 4`. It is labeled with three parts: "subject", "verb", and "object". A red arrow points from "subject" to the variable `x`. Another red arrow points from "verb" to the assignment operator `=`. A third red arrow points from "object" to the value `4`.

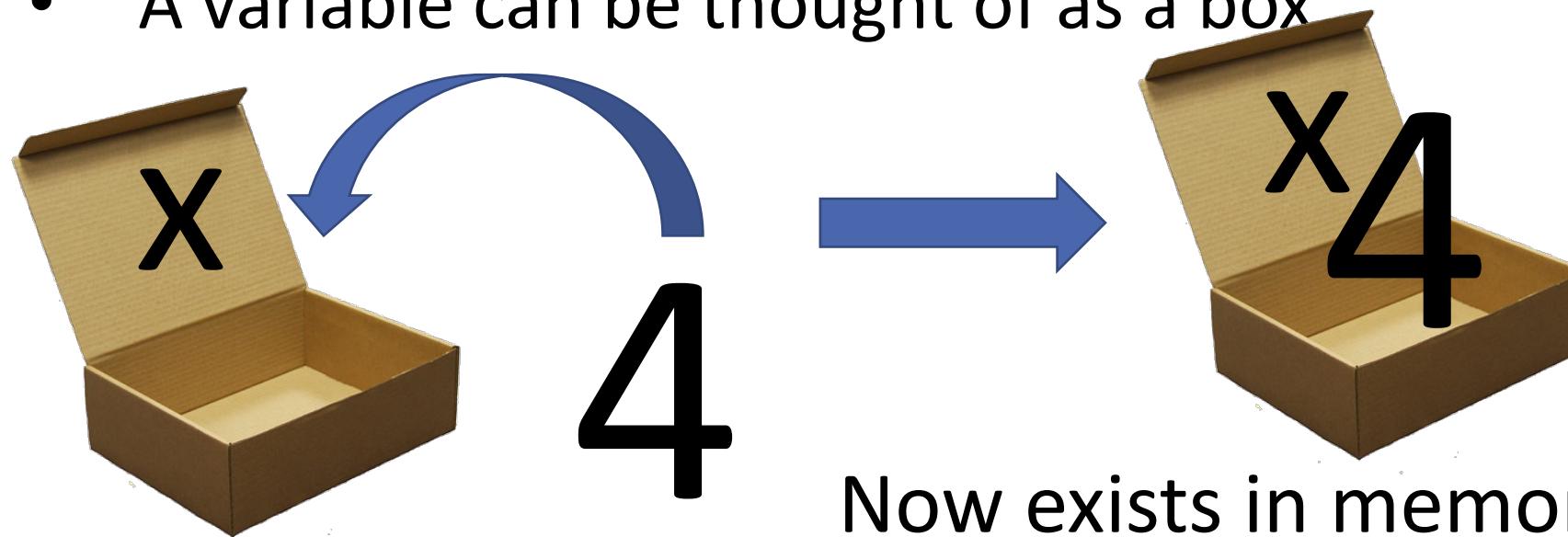
x = 4

# Variables and operators

variable operator data, value



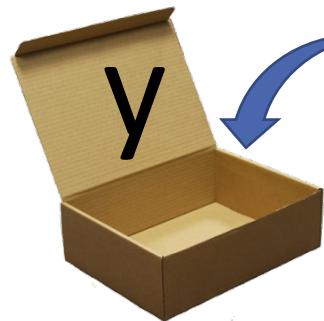
- An operator is the verb
- “=” assigns values to variables
- A variable can be thought of as a box



# Variables and operators

x = 4

y = 8



8

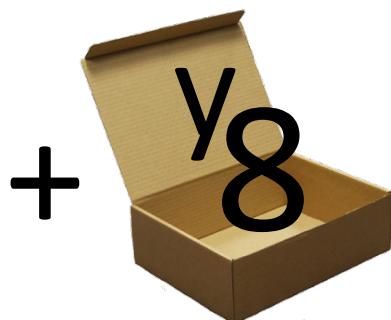


# Variables and operators

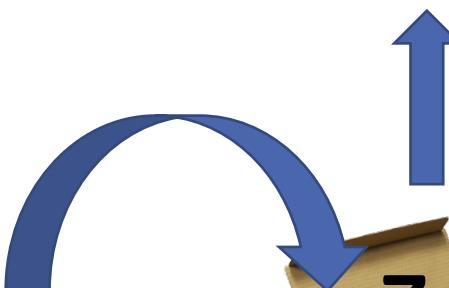
$x = 4$

$y = 8$

$z = x + y$



$= 12$



# Let's use Python!

1. Open a new text file and save it as “myfirstprogram.py”
2. Type the text below and save.

```
x = 4  
y = 8  
z = x + y  
print(z)
```

3. Open terminal and type “python myfirstprogram.py”

# Comments!

Any text followed by a “#” in the same line  
is not read by the computer

```
x = 4 # This is a line of code
y = 8 # This is another
z = x + y # z is the sum of x and y
# print(z)
```

# Why are comments useful?

- For when you look back later
- If other people are trying to read, use, or understand your code
  - E.g. your grader!
- To help make sure your thinking is clear

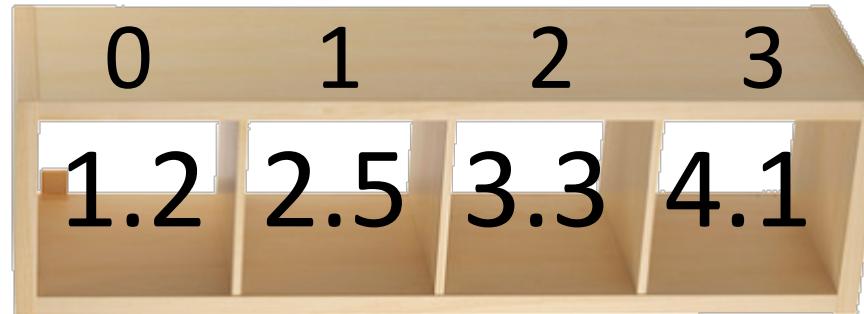
# You can also use Python interactively

- Open a terminal and type “python”
  - OR: Install Jupyter and open a notebook
- Now you can type lines of code, one at a time, and view the result in real time

```
>>> x = 1  
>>> print x  
1  
>>> x  
1
```

# A list is like a bookshelf of variables accessible by position in the sequence

```
x = [ 1.2, 2.5, 3.3, 4.1 ]
```



```
>>> print x[0]
```

```
1.2
```

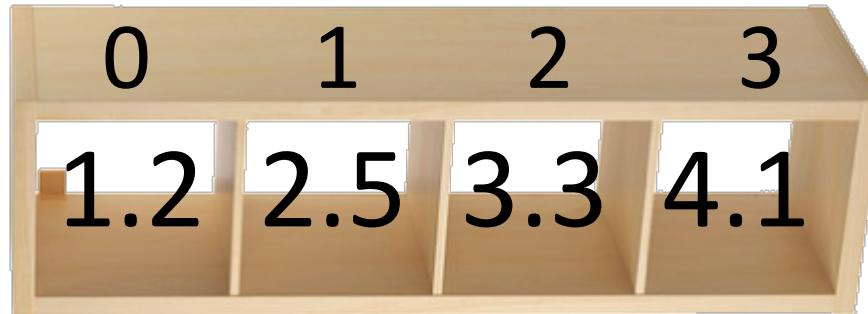
```
>>> print x[2]
```

```
3.3
```

```
>>> print x[-1] ?
```

# You can “slice” a list into a smaller piece with notation below

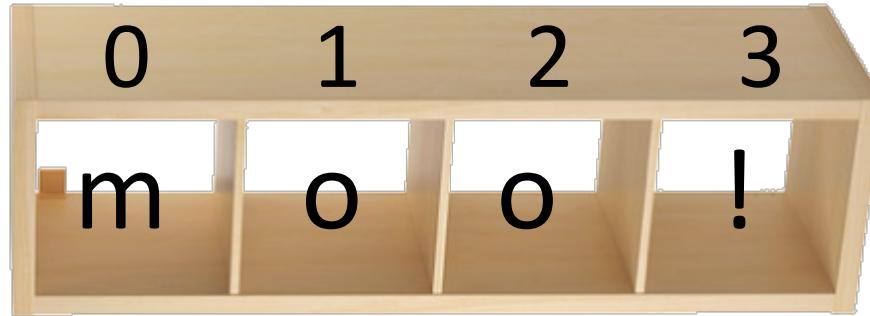
```
x = [ 1.2, 2.5, 3.3, 4.1 ]
```



```
>>> print x[0:2]  
[1.2, 2.5]  
>>> print x[1:3]  
[2.5, 3.3]  
>>> print x[1:]
```

# A string is like a list of characters

```
x = 'moo!'
```



```
>>> print x[0]
>>> print x[2]
>>> print x[-1]
```

# Variables have *types*

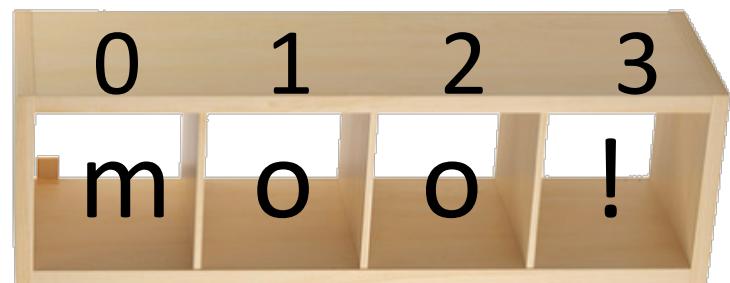
- Boolean
    - True or False
  - Int
    - 1, 12, -46, 0
  - Float
    - 1.24, 12.0, -0.5
- Simple types



# Variables have *types*

- List
  - [True, False, 1, 12]
- String
  - ‘hello how are you?’
- Hash/Dictionary
  - [True:12, False:1]

Complex types



# Common Boolean operators

```
x = 4 # not boolean! (assignment)
```

x == 4

x != 4

x > 4

x <= 3

x > 2 **and** x < 5

x == 4 **or** x != 4

# We can use Boolean operators in If/else statements

x = 4

Only things that evaluate to a Boolean go here

```
if x == 5:
    print 'x is 5!'
else:
    print 'x is not 5!'
'x is not 5!'
```

# Review and practice problems

<http://interactivepython.org/runestone/static/thinkcsby/index.html> Sections 1, 2, 4

Homework on Catalyst tomorrow...