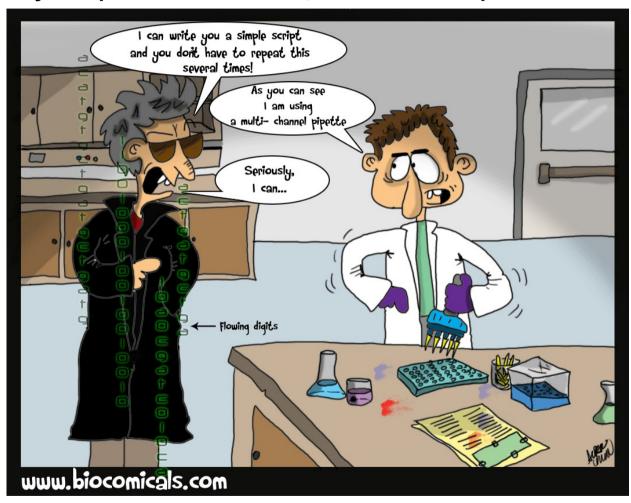
Part II: Python

If you repeat more than once, then write a script to do it!





Part II is a basic introduction to programming

What is Python? Why Python?



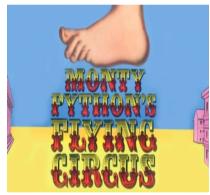
Anaconda

- A relatively new (~90') high-level scripting language
- Python is a interpreted language: we write our code in a text file (script) and then the Python interpreter translates it to machine code for the cpu on the fly.
- Python2.x vs Python 3.x
- Anaconda is a distribution of Python for large-scale data processing, predictive analytics, and scientific computing
- Why python?
 - Easy to use.
 - Widely spread in the bioinformatic community (has recently outranked Perl) as well as in many other fields:

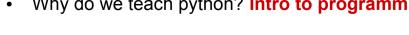
Nature vol 518 (feb 2015) pp 125: "What matters most in the early stages is having a good support network. "Pick the programming language based on what people around you are using,".... Increasingly, that language is Python"

- Expressive (in this context means that a single line of Python code can do more than a single line of code in most other languages) and readable
- Cross-platform
- Free
- Why do we teach python? Intro to programming









Programs and programming languages

- A computer program is a detailed, step-by-step set of instructions telling a computer exactly what to do.
- The instructions are executed sequentially
- Programming languages are human readable notations to express computations in a exact non unambiguous way.
 - Programs are written in some high-level human-readable/writable code (the source code)
 - Then the instructions are translated (compiled) to a machine-readable code (the compiled file or machine code).
 - Note: the source code is independent on the type of computer and OS (platform independent) but the compiled file will only run in a specific machine/OS (platform-specific).
- Programming languages can be classified into two broad types, based on how and when the compilation of the human readable code (source code) into machine code is done:
 - Compilled languages--> compilled programs are stored as binary files
 - Interpreted languages (scripting languages)--> interpreted programs are text files (scripts)
- Each programming language has a specific:
 - Lexicon: the set of words that can be used in a program.
 - Syntax: valid ways of combining the lexicon elements.
 - Semantics: meaning of the syntactically correct expressions
 - In spite of this, all programming languages have a common set of elements or structures.

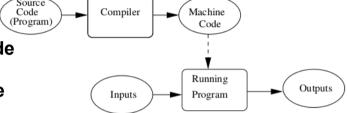


Figure 1.2: Compiling a High-Level Language

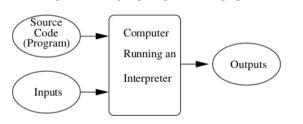
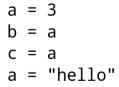


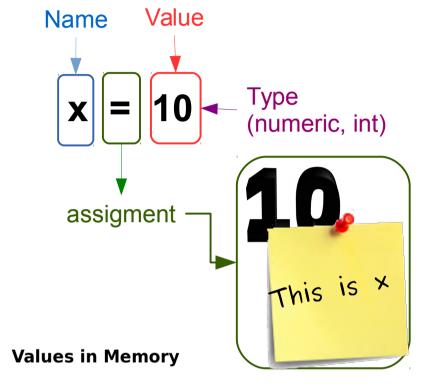
Figure 1.3: Interpreting a High-Level Language.

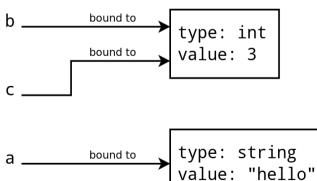
First element common to all programming languajes: Variables

- What is a variable
- Variable attributes:
 - Name (identifier)
 - Value
 - Type

Executed Code: Variable Assignment







Variables

Several ways to run Python programs

A) Python interactive interpreter

- 1) Open an interactive sesion with the python interpreter by typing "python3" in a terminal. Note that the prompt changes to ">>>" indicating we're now in a python shell.
- 2) Type the python commands to be executed (one statement at a time)
- 3) Exit the python shell by typing pressing ctrl key+D or typing:

>>> exit()

Note that all the code and results are now gone!

Terminal: text input/output environment through which you could interact with a computer

- On linux: Ctr+T
- On Mac OS: Command+space (search for "terminal")
- On Windows: Start→Search "powershell"

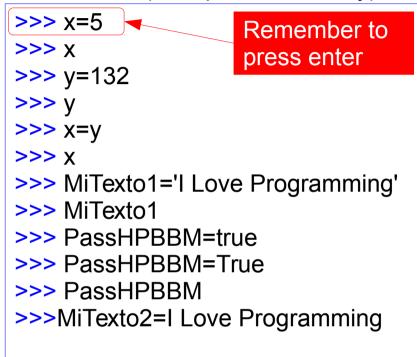
```
luis@luis-SATELLITE-PRO-C70-B: ~
luis@luis-SATELLITE-PRO-C70-B:~$ python3
Python 3.5.2 (default, Jul 5 2016, 12:43:10)
[GCC 5.4.0 20160609] on linux
Type "help"__"copyright", "credits" or "license" for more information.
>>> exit()
luis@luis-SATELLITE-PRO-C70-B:~$
```

Basic data types

Basic types of variable:

- Integer number (*int*). Stores whole numbers (between -2147483648 and +2147483648) without fractional component
- Large or fractional number (float).
 Holds numbers that are extremely large or have decimal fractions.
- Text (string). store a sequence of text characters. They are defined using quotation marks (either single or double)
- Boolean. True or False (0 or 1, respectively).

In a Python shell type the following commands (then press enter key):



Basic data types

Prompt (used in a command-line interface to indicate readiness to accept commands)

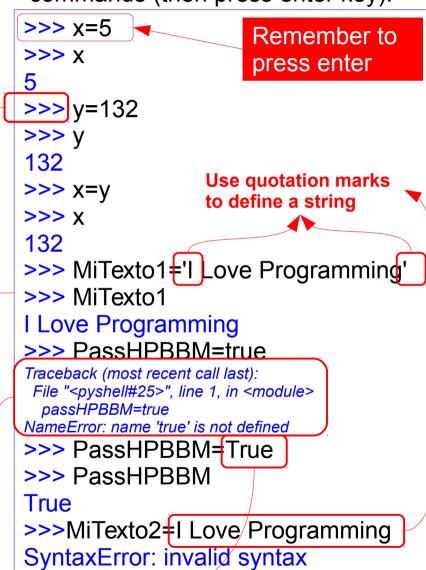
Basic types of variable:

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- Large or fractional number (float).
 Holds numbers that are extremely large or have decimal fractions.
- Text (string). store a sequence of text characters. They are defined using quotation marks (either single or double)
- Boolean. True or False (0 or 1, respectively).

Interactive Python sesion (in a python shell)
In the following slides I used the following color code:
What you type is in black
The computer output in blue

Error message. Python doesn't understand what "true" is

In a Python shell type the following commands (then press enter key):



Operators

Mathematical operators

Operator	Symbol	Action (numbers)	Action (strings)
Addition	+	Sum	Concatenate
Subtraction	-	Subtraction	none
Multiplication	*	Multiplication	Concatenate copies
Division	1	Division	none
Power	**	Power	none
Modulo	%	Remainder	none
Truncated division	//	Non-fractional part of result	none
Assignment	=	Assigns a number to variable	Assigns a string to variable

Comparative operators

Operator	Symbol
Equal to	==
Not equal to	!=, <>
Greater than	>
Less than	<
Greater or equal	>=
Less or equal	<=

Logical operators

Operator	Symbol
And	And, &(bitwise)
Or	or, (bitwise)
Not	Not, !

Do not mistake the assignment operator (=) for the equality operator (==)

Precedence. Python follows the same precedence rules for its mathematical operators that mathematics does: parentheses, exponentiation, Multiplication & Division, Addition and subtraction.

Variables & operators

Type the following statements in an interactive python interpreter and hit enter (shown just in the first one)

IMPORTANT: before hitting enter, try to guess the result you will get.

```
>>> 5+2
>>> "viva"+"luis"
>>> 5*3
>>> "luis"*3
>>> "viva"*"luis"
>>> "luis"/3
>>> "luis"+3
>>> type(5)
>>> type(3.1416)
>>> type(4/2)
>>> 10/3
```

type is a function (more about this soon) that, within this context, gives the type of the variable

```
>>> 5+2
>>> "viva"+"luis"
'vivaluis'
>>> 5*3
15
>>> "luis"*3
'luisluisluis'
>>> "viva"*"luis"
Traceback (most recent call last):
 File "<stdin>", line 1, in <module>
TypeError: can't multiply sequence by non-int of type 'str'
>>> "luis"/3
Traceback (most recent call last):
 File "<stdin>", line 1, in <module>
TypeError: unsupported operand type(s) for /: 'str' and 'int'
>>> "luis"+3
Traceback (most recent call last):
 File "<stdin>", line 1, in <module>
TypeError: Can't convert 'int' object to str implicitly
>>> type(5)
<class 'int'>
>>> type(3.1416)
<class 'float'>
>>> type(4/2)
<class 'float'>
>>> 10/3
3.3333333333333333
```

Note the final "5". There is a limit to the size and precision of the stored numbers.

Operators

Mathematical operators			
Operator	Symbol	Action (numbers)	Action (strings)
Addition	+	Sum	Concatenate
Subtraction	-	Subtraction	none
Multiplication	*	Multiplication	Concatenate copies
Division	1	Division	none
Power	**	Power	none
Modulo	%	Remainder	none
Truncated division	//	Non-fractional part of result	none
Assignment	=	Assigns a number to variable	Assigns a string to variable

RETURN A STRING OR NUMBER

Comparative operator		
Operator	Symbol	

Equal to ==

Not equal to !=, <>

Greater than >

Less than <

Greater or equal >=

Less or equal <=

Logical operators

Operator	Symbol
And	and,&(bitwise)
Or	or, (bitwise)
Not	Not,!
Is	is
In	in

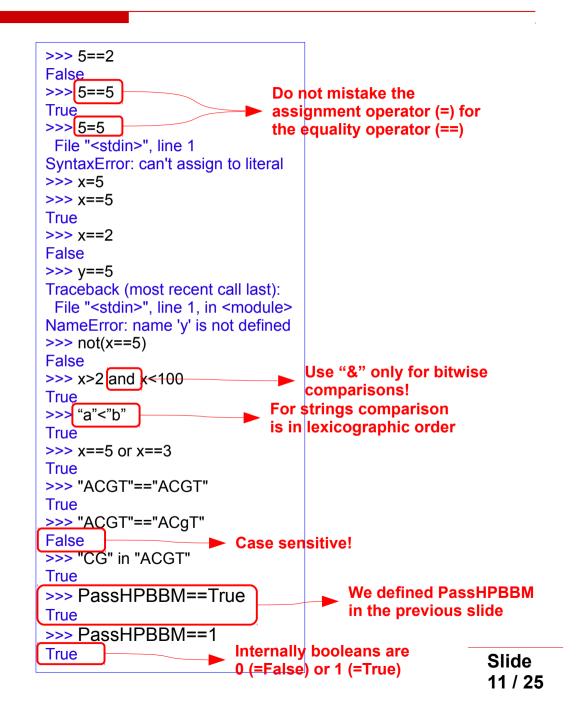
Do not mistake the assignment operator (=) for the equality operator (==)



Operators (comparative and logical)

Type the following statements in an interactive python interpreter and hit enter (shown just in the first one) IMPORTANT: try to guess the result you will get before hitting enter)

```
>>> 5==2
>>> 5==5
>>> 5=5
>>> x=5
>>> x==5
>>> x==2
>>> y==5
>>> not(x==5)
>>> x>2 and x<100
>>> "a"<"b"
>>> x==5 \text{ or } x==3
>>> "ACGT"=="ACGT"
>>> "ACGT"=="ACgT"
>>> "CG" in "ACGT"
>>> PassHPBBM==True
>>> PassHPBBM==1
```



Functions and methods

Functions can be thought of as little stand-alone programs that are called and executed within your program and return a specific calculation or carry out a specific task. Many function for common tasks come built into programming languages.

function_name ([parameter],[parameter],.....)

Functions (and methods) may expect zero, one, two,...or any number of values.

These are called *parameters or arguments*.

Trailing set of parentheses is required even when the function takes no parameters

Examples of Built-in Functions abs()
max()
help()
len()
min()
print()
sum()
type()

Examples of string type methods

str.upper()

str.lower()

str.count()

str.find()

str.replace()

Methods can be thought of as functions built into variables (see *objects*). They are used with the *dot notation*:

variable_name.function_name ([parameter],[parameter],.....

String methods are extremely useful in bioinformaticts to process nucleotide and protein sequences.

This are built-in functions but you can create your own functions as we'll see soon.

Part II: Python Variables and operators

Functions

```
>>> abs()
>>> abs(-5)
>>> x=-5
>>> abs(x)
>>> y=abs(x)
>>> y
>>> MyDNA="ACGTGC"
>>> len(MyDNA)
>>> round(2.45687)
>>> round(2.95687)
>>> round(2.95687,2)
>>> z=input("type any number and then press enter")
>>> Z
>>> type(z)
>>> int(z)
>>> help("round")
```

```
Help on built-in function round:

round(...)

round(number[, ndigits]) -> floating point number

Round a number to a given precision in decimal digits (default 0 digits).

This always returns a floating point number. Precision may be negative.

(END)

Press q (quick) to return to command shell
```

```
>>> abs()
Traceback (most recent call last):
 File "<stdin>", line 1, in <module>
TypeError: abs() takes exactly one argument (0)
given)
>>> abs(-5)
>>> x=-5
>>> abs(x)
                    Error, abs() function requires
                    one argument to work and we
                    didn't pass it
>>> y=abs(x)
>>> y
>>> MyDNA="ACGTGC"
>>> len(MvDNA)
>>> z=input("type any number and then press
enter")
>>> 7
(here whatever you typed)
>>>type(z)
                   Input() always returns a string!
<class 'str'>
>>> round(2.45687)
>>> help("round")
>>> round(2.95687,2)
2.96
```

String methods

Type the following statements in an interactive python interpreter and hit enter (shown just in the first one)
IMPORTANT: try to guess the result you will get before hitting enter)

```
>>> MyDNA="AcagTaC"
>>> MyDNA
>>> MvDNA.upper()
>>> MvDNA
>>> MyDNA_Mayus=MyDNA.upper()
>>> MyDNA Mayus
>>> MyDNA.lower()
>>> MyDNA.count("A")
>>> MyDNA Mayus.count("A")
>>> MyDNA.upper().count("A")
>>> MyDNA.count("A").upper()
>>> MyDNA.find("a")
>>> MyDNA.find("a",3)
>>> MyDNA.replace("T","U")
>>> MyDNA.split("a")
>>> MyDNA_Mayus.split("a")
>>> MyDNA.split("a",1)
```

```
>>> MyDNA="AcagTaC"
>>> MvDNA
'AcagTaC'
>>> MyDNA.upper()
'ACAGTAC'
>>> MvDNA
'AcagTaC'
>>>
MyDNA Mayus=MyDNA.upper()
>>> MyDNA Mayus
'ACAGTAC'
>>> MyDNA.lower()
'acagtac'
>>> MyDNA.count("A")
>>> MyDNA Mayusy.count("A")
>>>MyDNA.upper().count("A")
>>>MyDNA.count("A").upper()
Traceback (most recent call last):
 File "<stdin>", line 1, in
<module>
AttributeError: 'int' object has no
attribute 'upper'
>>> MyDNA.find("a")
>>> MyDNA.find("a",3)
```

```
>>> MyDNA.replace("T","U")
'AcagUaC'
>>> MyDNA="ACGTGC"
>>> MyDNA.split("a")
['Ac', 'gT', 'C']
>>> MyDNA_Mayus.split("a")
['ACGTGC']
>>> MyDNA.split("a",1)
['Ac', 'gTaC']
```

Note that you can concatenate several methods in a single statement.

This is a list, we'll see them in

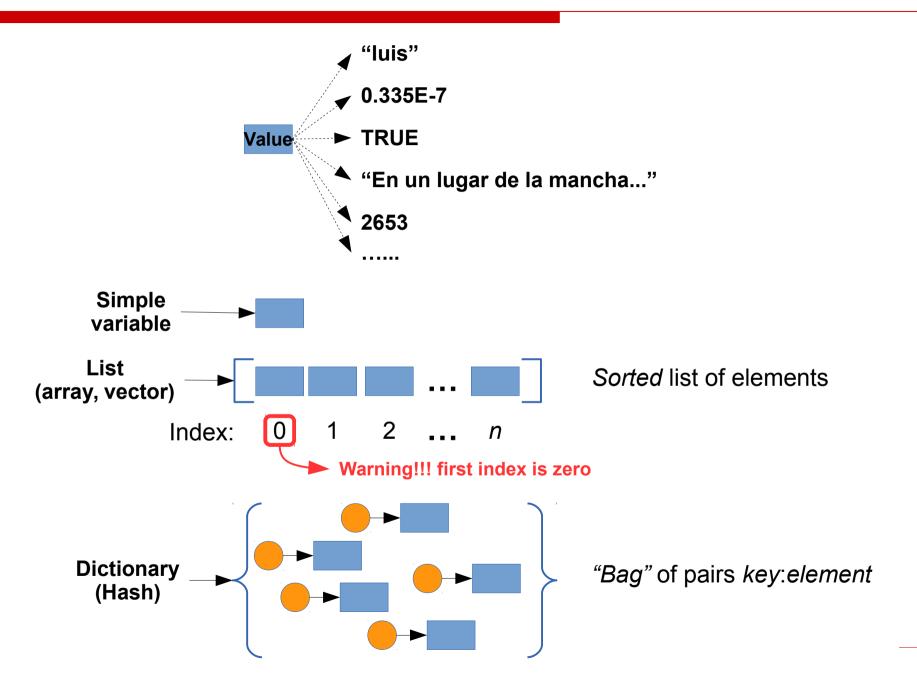
the next slide

However, beaware of the order, they're avaluated from left to right.

By default, finds first occurence

Slide 14 / 25

Lists (arrays) and dictionaries



Lists (arrays) and dictionaries

Storing amino acids MWs into a list

```
>>>aaMW=[57,71,87,101,103,99,113,113,131,97,147,
163,186,115,129,114,128,137,128,156]
>>> aaMW
>>> aaMW[0]
>>> aaMW[3]
>>> len(aaMW)

>>> len(aaMW)
```

How would you calculate the average weight in the class without typing weight values again but using the variable Weight4Bq?

```
>>> REnz={"EcoRI":"GAATTC", "BamHI":"GGATCC",
"HindIII":"AAGCTT", "NotI":"GCGGCCGC"}

>>> REnz["NotI"]
>>> REnz["EcoRI"]
>>> REnz.keys()

We define (declare) a
dictionary by using curly-
backets and pairs
key:value using colon
```

From the course's Moodle page, open the document InClassExercise_Dictionaries.txt and copy its content into the Python shell.

What type of variable is HRAS? and REnz? How do you know it?

Which of the enzymes above cut in the sequence of HRAS? (tip you just need to use the two variables in this exercise and one of the operators we've already seen)

```
>>>aaMW=[57,71,87,101,103,99,113,113,131,97,147,
163,186,115,129,114,128,137,128,156]
>>> aaMW
[57, 71, 87, 101, 103, 99, 113, 113, 131, 97, 147, 163,
186, 115, 129, 114, 128, 137, 128, 156]
>>> aaMW[0]
57
>>> aaMW[3]
101
>>> Weight4Bq[-1]
156
>>> len(Weight4Bg)
20
>>> sum(Weight4Bg)
2375
>>> sum(Weight4Bq)/len(Weight4Bq)
118.75
>>> REnz={"EcoRI":"GAATTC", "BamHI":"GGATCC",
"HindIII": "AAGCTT", "NotI": "GCGGCCGC"}
>>> REnz["Notl"]
'GCGGCCGC'
>>> REnz["EcoRI"]
'GAATTC'
>>> REnz.keys()
dict keys(['BamHI', 'EcoRI', 'NotI', 'HindIII'])
>>> HRAS="GGTCCCGGC......
>>> REnz["Notl"] in HRAS
True
>>> REnz["EcoRI"] in HRAS
                                              Slide
False
```

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In class exercises

• The nucleotide frequency in the genome of a newly discovered specie is given in the following dictionary: NucFrq={'A':0.35,'C':0.25,'G':0.3,'T':0.2}. Using the variable NucFrq, what would be the probability of finding the EcoRI restriction site ('GAATTC')? How many EcoRI sites would you expect if the genome is 2.7*10⁵ bases long? Solve it using the interactive python shell and the provided variable (dictionary).

```
>>> NucFrq={'A':0.35,'C':0.25,'G':0.3,'T':0.2}
>>> (NucFrq['G']* (NucFrq['A']**2)* (NucFrq['T']**2)* NucFrq['C'])*270000
99.225
```

• The nucleotide frequency in the genome of a newly discovered specie is given in the following array: NucFrq=[0.4,0.3,0.2,0.1], where the values correspond to the frequencies of "A", "C", "G" and "T" respectively. Using the variable NucFrq, what would be the probability of finding the HindIII restriction site ('AAGCTT')? How many HindIII sites would you expect if the genome is 3.4*109 bases long? Solve it using the interactive python shell and the provided variable (array).

```
>>> NucFrq2=[0.4,0.3,0.2,0.1]
>>> (NucFrq2[1]* (NucFrq2[0]**2)* (NucFrq2[3]**2)* NucFrq2[2])*(3.4e9)
326400.000000002
>>> round((NucFrq2[1]* (NucFrq2[0]**2)* (NucFrq2[3]**2)* NucFrq2[2])*(3.4e9))
326400
```

Part II: Python Variables and operators

Already too much code for an interactive session: save code as an script

A) Python interactive interpreter

- 1) Open an interactive sesion with the python interpreter by typing "python3" in a terminal. Note that the prompt changes to ">>>" indicating we're now in a python shell.
- 2) Type the python commands to be executed (one statement at a time)
- 3) Exit the python shell by typing pressing ctrl key+D or typing: >>> exit()

Note that all the code and results are now gone!

B) IDLE (Integrated DeveLopment Environment for Python)

- 1) The main IDLE shell window is equivalent to the regular interactive interpreterinal with a few improvements
- 2) It includes a python code editor: go to "File-->New file" under the IDLE shell menu
- 3) The code editor allows you to save a list of commands and execute them in batch.
- 4) The code is saved as a plain text file: this is an **script**.

C) Running python from an script

- 1) Write program (code) in your favourite text editor.
- 2) Save it (remember it is a text file!). This is your source code (your script). For example save it as my1program.py
- 3) Run your program. To do this last step type in a terminal:

The general syntax is:

[interpreter script_name of the limit of the l

D) More sophisticated IDEs (Integrated Development Environment)

Spyder, Ninja, SPE, ERIC, many others

Terminal: text input/output environment through which you could interact with a computer

- · On linux: Ctr+T
- On Mac OS: Command+space (search for "terminal")
- On Windows: Start→Search "powershell"

To open IDLE:

- On linux: type idle3 in a terminal
- On Windows/Mac: select IDLE from program menu

This send your source code (saved in the *my1program.py* text file) to the python interpreter (a program that translates your source code into machine code) who compiles it and pass the instructions to the CPU

Your first script: computing GC content of a DNA sequence

1 Open IDLE

```
Python 3.5.2 Shell

File Edit Shell Debug Options Window Help

Python 3.5.2 (default, Jul 5 2016, 12:43:10)

[GCC 5.4.0 20160609] on linux

Type "copyright", "credits" or "license()" for more information.

>>>> |

Ln: 4 Col: 4
```

2 File-→New File

- 3 Type commands
- 4 Save script: File-->save
- Sun script:
 Run-->Run module

```
percentGC.py - /home/luis/Documents/Teaching/HPBBM/2016

File Edit Format Run Options Window Help

DNA=input("Enter your DNA (and hit enter): ")
numberG=DNA.upper().count("G")
numberC=DNA.upper().count("C")
numberA=DNA.upper().count("A")
numberT=DNA.upper().count("T")
percentGC=(numberG+numberC)/(numberG+numberC+numberA+numberT)
print("%GC:")
print(percentGC)
```

```
Eile Edit Shell Debug Options Window Help

Python 3.5.2 (default, Jul 5 2016, 12:43:10)
[GCC 5.4.0 20160609] on linux
Type "copyright", "credits" or "license()" for more information.

>>>
===== RESTART: /home/luis/Documents/Teaching/HPBBM/2016_17/percentGC.py =====
Enter your DNA (and hit enter): aggcgatgacgcgtgca
%GC:
0.6470588235294118
>>> |
```

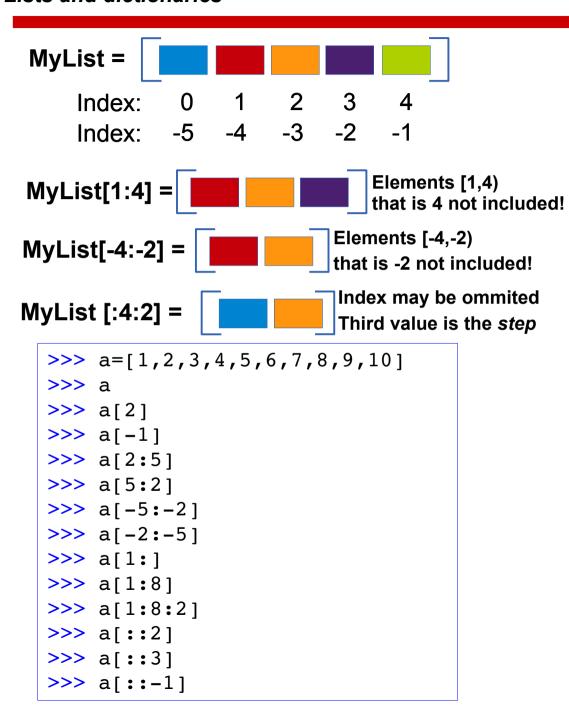
You are now ready to complete programming Assignment 1!

Modifying lists and dictionaries

```
>>> MyList=list(range(2,20,3))
>>> MyList
>>> MyList[3]="x"
>>> MyList
>>> MyList=MyList+["a"]
>>> MyList
>>> MyList=["b","c"]+MyList
>>> MyList
>>> MyList.append("5")
>>> MyList
>>> MyList.insert(4,"new")
>>> MyList
>>> MyList.remove("x")
>>> MyList
>>> MyList.pop(4)
>>> MyList
>>> MyList.clear()
>>> MyList
>>> MyList=[34,5,14,3,1]
>>> MyList.reverse()
>>> MyList
>>> sorted(MyList)
>>> MyList
>>> MyList.sort()
>>> MyList
>>> MyList=[]
>>> MyList
```

```
>>> MyDic={'A':'Ala','G':'Gly','Y':'Tyr'}
>>> MyDic
>>> MyDic['F']='Phe'
>>> MyDic
>>> MyDic.keys()
>>> MyDic.values()
>>> MyDic.get('G')
>>> MyDic.get('S')
>>> MyDic.get('S','not found')
>>> del MyDic['G']
>>> MyDic
>>> del MyDic
>>> MyDic
>>> MyList1=['A','G','Y']
>>> MyList2=['Ala','Gly','Tyr']
>>> MyDic=dict(zip(MyList1,MyList2))
```

List slicing



```
\Rightarrow \Rightarrow a = [1,2,3,4,5,6,7,8,9,10]
>>> a
[1, 2, 3, 4, 5, 6, 7, 8, 9, 10]
>>> a[2]
3
>>> a[-1]
10
>>> a[2:5]
[3, 4, 5]
>>> a[5:2]
[]
>>> a[-5:-2]
[6, 7, 8]
>>> a[-2:-5]
[]
>>> a[1:]
[2, 3, 4, 5, 6, 7, 8, 9, 10]
>>> a[1:8]
[2, 3, 4, 5, 6, 7, 8]
>>> a[1:8:2]
[2, 4, 6, 8]
>>> a[::2]
[1, 3, 5, 7, 9]
>>> a[::3]
[1, 4, 7, 10]
                              Slide
>>> a[::-1]
[10, 9, 8, 7, 6, 5, 4, 3,
```

Strings as lists.

In some respects strings behave as if they were lists of characters: strings can be indexed and sliced. The function list() and the method join() can be used to transform a string into a <u>true</u> list and a list into a string respectively.

```
MyString = " P y t h o n "

Index: 0 1 2 3 4 5

Index: -6 -5 -4 -3 -2 -1
```

```
>>> MyDNA='ACGTGACGACCATGA'
>>> MyDNA[3]
>>> MyDNA[3:5]
>>> MyDNA[-1]
>>> MyDNA[3:]
>>> MyDNA[2:8:2]
>>> MyDNA[::-1]
>>> list(MyDNA)
>>>

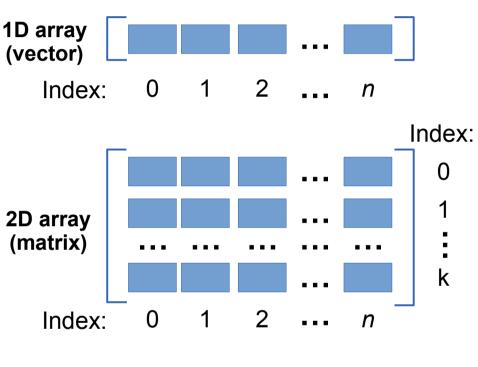
MyList=["B","i","o","c","h","e","m","
i","s","t","r","y"]
>>> "-".join(MyList)
>>> "".join(MyList)
```

```
>>> MyDNA='ACGTGACGACCATGA'
>>> MyDNA[3]
יידי
>>> MyDNA[3:5]
'TG'
>>> MyDNA[-1]
'A'
>>> MyDNA[3:]
'TGACGACCATGA'
>>> MyDNA[2:8:2]
'GGC'
>>> MyDNA[::-1]
'AGTACCAGCAGTGCA'
>>> list(MyDNA)
['A', 'C', 'G', 'T', 'G', 'A',
'C', 'G', 'A', 'C', 'C', 'A',
'T', 'G', 'A']
>>>
MyList=["B","i","o","c","h","e"
,"m","i","s","t","r","y"]
>>> "-".join(MyList)
'B-i-o-c-h-e-m-i-s-t-r-y'
>>> "".join(MyList)
'Biochemistry'
```

Multidimensional arrays

So far we have studied one-dimensional arrays (also known as vectors). However, arrays can have any number of dimensions. In all the cases we use the square-bracket notation to access the array elements.

```
>>> a=[1,2,3,4,5,6]
>>> a
[1,2,3,4,5,6]
>>> a[2]
>>> aa=[[1,2,3],[4,5,6],[7,8,9]]
>>> aa
[[1, 2, 3], [4, 5, 6], [7, 8, 9]]
>>> aa[1]
[4, 5, 6]
>>> aa[1][0]
4
>>> aa[2][2]
9
>>> aaa=[[[1,2],[3,4]],[[5,6],[7,8]]]
>>> aaa
[[[1,2],[3,4]],[[5,6],[7,8]]]
>>> aaa[1]
[[5, 6], [7, 8]]
>>> aaa[1][0]
[5, 6]
>>> aaa[1][0][1]
6
```



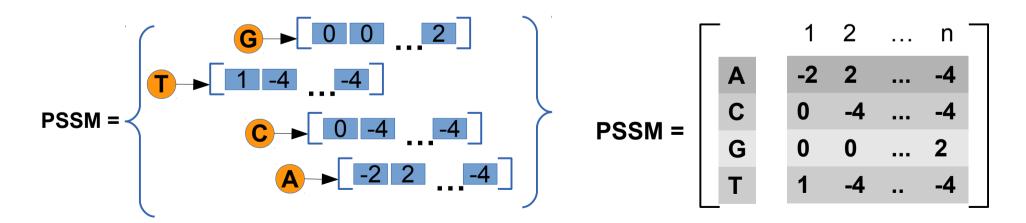
aa =
$$\begin{bmatrix} 1 & 4 & 7 \\ 2 & 5 & 8 \\ 3 & 6 & 9 \end{bmatrix}$$

Not limited to 2D...

Combinations of lists and arrays

Similarly, dictionaries can also consists in more than one dimension. Moreover, we can combine arrays and dictionaries.

```
>>> PSSM={'A':[-2,2,-4,-4,-4],'C':[0,-4,2,-4,-4],'G':[0,0,-4,2,-4,-4],'G':[0,0,-4,2,-4,-4],'T':[1,-4,-4,-4],'B':[0,0,-4,2,-4],'B':[1,-4,-4,-4],'B':[0,0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,-4],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2],'C':[0,-4,2,2],'C':[0,-4,2,2],'C':[0,-4,2],'C':
```



Some on-line programming aids and learning tools

Learning on-line:

- http://software-carpentry.org/
- https://www.codecademy.com/learn
- https://www.edx.org/course/introduction-computer-science-mitx-6-00-1x-8

On-line books:

- https://learnpythonthehardway.org/book/
- http://greenteapress.com/thinkpython/html/index.html

Programming aids

- http://pythontutor.com/
- https://github.com/
- http://blog.coderscrowd.com/real-time-programming-for-bioinformatics-and-for-fun/
- http://scratch.mit.edu/





Online Python Tutor: Embeddable Web-Based Program Visualization for CS Education





