

### **Biotecnologia**

# Introduction to Programming with Python

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#### **Tutorial Outline**

- interactive "shell"
- basic types: numbers, strings
- list
- variables
- control structures
- functions & procedures
- files & standard library



## Código em Python:

print "Hello World!"

```
Código em Java:

public class Hello

{

public static void main(String[] args) {

    System.out.printf("Hello World!");
    }
}
```



#### Interactive "Shell"

- Great for learning the language
- Great for experimenting with the library
- Great for testing your own modules
- Two variations: IDLE (GUI), python (command line)
- Type statements or expressions at prompt:

```
>>> print "Hello, world"
Hello, world
>>> x = 12**2
>>> x/2
72
>>> # this is a comment
```



#### Interactive "Shell"

```
>>> print "hello world"
hello world
>>> 2+2
>>> 52*345
17940
>>> 2.0 / 3.0
0.666666666666666
>>> a=1
>>> b=2
>>> a+b
3
>>> for a in range(5):
        print a,
01234
>>>
```



## Data Type

#### Numéricos:

- Integer (1,2,3,-1,-2,0,7,234,...)
- Float (1.23,0.3566,3.53e+63)
- Complex (2+3j,2j,5+27j)
- Boolean (True, False)

#### Estruturados:

- Strings ('ATGCCCAATTG')
- Listas ([ 1,5, 0.2, 'A', 'xxx' ])
- Tuplas ( (1, 2, 3) )
- Conjuntos ( set(['A','T','G','C']) )
- Dicionários ( { 'A' : 'Ala', 'V' : 'Val', 'I' : 'Ile', 'L' : 'Leu' } )

Ordenados (sequenciais)

Não Ordenados



## Strings

- "hello"+"world" "helloworld" # concatenation
- "hello"\*3 "hellohello" # repetition
- "hello"[0] "h" # indexing
- "hello"[-1] "o" # (from end)
- "hello"[1:4] "ell" # slicing
- len("hello")5# size
- "hello" < "jello" 1 # comparison</li>
- "e" in "hello" 1 # search
- "escapes: \n etc, \033 etc, \if etc"
- 'single quotes' """triple quotes""" r"raw strings"



'GGC'

## Strings

```
>>> seq1='ATGGGCA'
>>> seq2='AATTAAAT'
>>> seq1 + seq2
'ATGGGCAAATTAAAT'
>>> poly alanine='A'*100
>>> len(poly_alanine)
100
>>> poly alanine
>>> seq1[0]
'A'
>>> seq1[3:6]
```



## Strings (operations)

```
>>> seq1
'ATGGGCA'
>>> 'A' in seq1
True
>>> 'X' in seq1
False
>>> len(seq1)
>>> min(seq1)
'A'
>>> max(seq1)
'T'
>>> seq1*3
'ATGGGCAATGGGCA'
```



## Strings (Methods)

Método	Descrição
str.capitalize()	Primeira letra para maiúscula
str.count(s)	Conta número de ocorrências de s
str.index(s,[start, [end]])	Retorna a posição de <b>s</b>
str.replace(s,r)	Subsitui s por r
str.split(sep)	Separa numa lista usando separador sep
str.lower()	Converte para minúsculas
str.strip()	Remove espaços circundantes
str.isalpha()	True se str é apenas alfabético
str.upper()	Convere para maísculas



## Strings (Methods)

```
>>> myseq = 'ATTGGCCAAACCG'
>>> myseq.count('A')
4
>>> myseq.count('CC')
2
>>> myseq.replace('A','U')
'UTTGGCCUUUCCG'
>>> myseq.capitalize()
'Attggccaaaccg'
>>> myseq.lower()
'attggccaaaccg'
>>> myseq.index('AAA')
>>> '193.136.227.168'.split('.')
['193', '136', '227', '168']
```



#### Lists

- Flexible arrays, not Lisp-like linked lists
  - a = [99, "bottles of beer", ["on", "the", "wall"]]
- Same operators as for strings
  - a+b, a\*3, a[0], a[-1], a[1:], len(a)
- Item and slice assignment
  - a[0] = 98
  - a[1:2] = ["bottles", "of", "beer"]
    - -> [98, "bottles", "of", "beer", ["on", "the", "wall"]]
  - del a[-1] # -> [98, "bottles", "of", "beer"]



## More List Operations

```
>>> a = range(5)
                    # [0,1,2,3,4]
                    # [0,1,2,3,4,5]
>>> a.append(5)
                    # [0,1,2,3,4]
>>> a.pop()
5
>>> a.insert(0, 42) # [42,0,1,2,3,4]
                    # [0,1,2,3,4]
>>> a.pop(0)
5.5
>>> a.reverse()
                     # [4,3,2,1,0]
                     # [0,1,2,3,4]
>>> a.sort()
```



#### Variables

- No need to declare
- Need to assign (initialize)
  - use of uninitialized variable raises exception
- Not typed

```
if friendly: greeting = "hello world" else: greeting = 12**2 print greeting
```

- Everything is a "variable":
  - Even functions, classes, modules



#### Variables

```
>>> 1aa = 12
   SyntaxError: invalid syntax
   >>> Ala+ = 10
   SyntaxError: invalid syntax
   >>> new_seq = 'ATTTGGC'
   >>> Ala = 10
   >>> Ala+new seq
   Traceback (most recent call last):
     File "<pyshell#345>", line 1, in <module>
       new seq+Ala
   TypeError: cannot concatenate 'str' and 'int' objects
   >>> str(Ala)+new_seq
   '10ATTTGGC'
```

#### Reference Semantics

- Assignment manipulates references
  - x = y does not make a copy of y
  - x = y makes x reference the object y references
- Very useful; but beware!
- Example:

```
>>> a = [1, 2, 3]
>>> b = a
>>> a.append(4)
>>> print b
[1, 2, 3, 4]
```



#### **Control Structures**

if *condition*: while *condition*:

statements statements

[elif condition:

statements] ... for var in sequence:

else: statements

statements

break

continue



## **Grouping Indentation**

```
In C:
In Python:
                                for (i = 0; i < 20; i++)
for i in range(20):
   if i\%3 == 0:
                                   if (i\%3 == 0) {
      print i
                                       printf("%d\n", i);
     if i\%5 == 0:
                                      if (i\%5 == 0) {
         print "Bingo!"
                                         printf("Bingo!\n"); }
   print "---"
                                     printf("---\n");
```

```
Bingo!
12
15
Bingo!
18
```



## Functions, Procedures

```
def name(arg1, arg2, ...):
    """documentation"""# optional doc string
    statements
```

```
return # from procedure
return expression # from function
```



#### Instance Variable Rules

- On use via instance (self.x), search order:
  - (1) instance, (2) class, (3) base classes
  - this also works for method lookup
- On assignment via instance (self.x = ...):
  - always makes an instance variable
- Class variables "default" for instance variables
- But...!
  - mutable class variable: one copy shared by all
  - mutable instance variable: each instance its own



#### Modules

- Collection of stuff in foo.py file
  - functions, classes, variables
- Importing modules:
  - import re; print re.match("[a-z]+", s)
  - from re import match; print match("[a-z]+", s)
- Import with rename:
  - import re as regex
  - from re import match as m



## File Objects

- f = open(filename[, mode[, buffersize])
  - mode can be "r", "w", "a" (like C stdio); default "r"
  - append "b" for text translation mode
  - append "+" for read/write open
  - buffersize: 0=unbuffered; 1=line-buffered; buffered

#### methods:

- read([nbytes]), readline(), readlines()
- write(string), writelines(list)
- seek(pos[, how]), tell()
- flush(), close()
- fileno()



## Input / Output

raw\_input() - data type string / input()

```
>>> seq = raw input("Introduza uma sequência: ")
Introduza uma sequência: ATTGGCCCGAA
>>> print seq
ATTGGCCCGAA
>>> n = raw input("Introduza um número: ")
Introduza um número: 5
>>> print "o quadrado do número introduzido é ", n*n
>>> n = int(raw input("Introduza um número: "))
Introduza um número: 5
>>> print "o quadrado do número introduzido é ", n*n
o quadrado do número introduzido é 25
```



## Input / Output

```
>>> x=55.23456
>>> print "O valor de x é %5.2f" % x
O valor de x é 55.23
>>> print "O valor de x é %08.3f" % x
O valor de x é 0055.235
```

```
Forma geral de print formatado:

print "... %n.mc ... %n.mc ... " % (x,y,z,...)

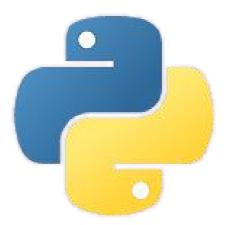
FORMATOS VARIÁVEIS
```



## Python + GUI Interface

IDE geany







## python training exercise

1 - Write a program that translates a DNA sequence into mRNA

#### AAATTGCGCG = AAAUUGCGCG

2 - Write a program that calculates the GC content of a DNA sequence

```
gc_count = g_count + c_count
gc content = (100.0*gc count) / length
```