

# **BINF2111 - Introduction to Bioinformatics Computing**

**To function or not to function that is the question?**



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# Learning Objectives

- Import
- Functions
- AT\_content function
- Function or to not to function?
- Testing and importing
- Quiz 19

# Imports

Time/date time import

```
>>> import time  
>>> from datetime import date  
>>> today = date.today()  
>>> today
```

Output?

# Imports

Time/date time import

```
>>> import time  
>>> from datetime import date  
>>> today = date.today()  
>>> today
```

**Output >>> datetime.date(2021, 11, 9)**

Today == date.fromtimestamp(time.time())

Output ?

# Imports

Time/date time import

```
>>> import time  
>>> from datetime import date  
>>> today = date.today()  
>>> today
```

**Output >>> datetime.date(2021, 11, 9)**

```
Today == date.fromtimestamp(time.time())
```

**Output True**

# Imports

Time/date time import

```
>>> import datetime
```

```
>>> datetime.datetime.now()
```

```
>>>
```

Output?

# Imports

## Time/date time import

```
>>> import datetime  
>>> datetime.datetime.now()  
>>>
```

Output >>> datetime.datetime(2021, 11, 8, 20, 25, 48, 294368)

In this example you import the datetime **class**. You have to explicitly call the datetime **type** and the **function** now.

# Common Imports

**These are some commonly used imports**

**import zlib**

Allows you to compress your data

**import math**

Allows for mathematical functions

**import re**

string matching patterns (will talk about more in regular expressions)



# Common Imports

**These are some commonly used imports**

**import sys**

Creates a list of the system arguments.

**import os**

Allows you to send commands to the shell.

**Import argparse**

makes it easy to write user-friendly command-line interface

# Common Imports

**These are some commonly used imports**

## **import numpy**

Comprehensive mathematical functions, random number generators, linear algebra routines, Fourier transforms, and more.

## **import pandas**

For data manipulation and analysis. In particular, it offers data structures and operations for manipulating numerical tables and time series.

# Common Imports

**These are some commonly used imports**

**import scikit-learn**

machine learning library, tools for predictive data analysis

**import matplotlib, seaborn, plotly**

Plotting imports

# Functions

**There are two kinds of functions in Python.**

- 1) **Built-in functions** that are provided as part of Python -  
`input()`, `type()`, `float()`, `int()` ...
- 2) **Custom Functions** that we define ourselves

In Python a function is some reusable code that takes argument(s) as input, does some computation, and then returns a result or results

As we showed earlier you can import functions that have been previously written

# AT vs. GT content function

```
#!/usr/bin/python3
```

```
my_dna = 'ACTGATCCATATATATTTT'  
length = len(my_dna)  
a_count = my_dna.count('A')  
t_count = my_dna.count('T')  
g_count = my_dna.count('G')  
c_count = my_dna.count('C')  
at_content = (a_count + t_count) / (length)  
gc_content = (1 - at_content)  
print('AT content is: ' + str(at_content))  
print('GC content is: ' + str(gc_content))
```

# AT vs. GT content function

This script counts the AT content of a piece of dna. But if we want to do it later, we have to reassign the variables, and re-run.

If we write a function, we can count the AT content of any string as we encounter it or read it in, without reassigning `a_count` and `t_count` etc.

# Functions

Define a function using : **def**

Call/invoke the function by using the function name, parenthesis and arguments in an expression

The first statement can be optional documentation of the function

The code block within the defined function is indented and starts with a colon :

The statement `return [expression]` exits the function, and can pass back an expression when used.

Example

```
def printme(str):  
    "This prints a passed string into this function"  
    print str  
    return str
```

# Functions

The function will not execute on its own, call the function to make it run.

#Function definition is here

```
def printme( str ):  
    "This prints a passed string into this function"  
    print str  
    return str
```

#Calling the printme function

```
printme("First call of user-defined function!")  
printme("Second call of user-defined function")
```

Once we have defined a function, we can invoke it as many times as we like



# Arguments

Arguments are passed into the function as input when the function is called.

Arguments are usually variables but can be values input directly.

The argument is a placeholder for the values we want to pass to the function.

# Function definition is here

```
def printme( str ):
```

```
    "This prints a passed string into this function"
```

```
    print str
```

```
    return
```

# Now you can call the printme function

```
printme("I'm first call to user defined function!")
```

```
printme("Again second call to the same function")
```

# Arguments

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Arguments are usually variables but can be values input directly.

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```
def printme( str ):
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    "This prints a passed string into this function"
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    print str
```

```
    return
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# Now you can call the printme function

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printme("I'm first call to user defined function!")
```

```
printme("Again second call to the same function")
```

# Arguments/Functions - AT counter

Now lets define a function that calculates A/T content.

```
def get_at_content(dna):  
    length = len(dna)  
    a_count = dna.count('A')  
    t_count = dna.count('T')  
    at_content = (a_count + t_count) / length  
    return at_content
```

```
my_at_content = get_at_content("ATTTACTACATACGCGAGCGAG")  
print(str(my_at_content))
```

**Output?**

# Arguments/Functions - AT counter

Now lets define a function that calculates A/T content.

```
def get_at_content(dna):  
    length = len(dna)  
    a_count = dna.count('A')  
    t_count = dna.count('T')  
    at_content = (a_count + t_count) / length  
    return at_content
```

```
my_at_content = get_at_content("ATTTACTACATACGCGAGCGAG")  
print(str(my_at_content))
```

**Output - 54.54%**

# Arguments/Functions - AT counter

```
#!/usr/bin/python3
```

```
def get_at_content(dna):  
    length = len(dna)  
    a_count = dna.count('A')  
    t_count = dna.count('T')  
    at_content = (a_count + t_count) / length  
    return at_content
```

```
my_at_content = get_at_content("ATTTACTACATACGCGAGCGAG")  
print(str(my_at_content))  
print(get_at_content("ATCGCGCGCATCGTY"))  
print(get_at_content("atatatatcgcgcatcagt"))  
print("AT content is " + str(get_at_content("AATTAGGCGC")))
```

**Outputs?**

# Arguments/Functions - AT counter

**Outputs?**

0.5454545454545454

0.3333333333333333

0.0

AT content is 0.5

**Function?**

# Arguments/Functions - AT counter

```
#!/usr/bin/python3
```

```
def get_at_content(dna):  
    length = len(dna)  
    a_count = dna.upper().count('A')  
    t_count = dna.upper().count('T')  
    at_content = (a_count + t_count) / length  
    return round(at_content,2)  
  
my_at_content = get_at_content("ATTTACTACATACGCGAGCGAG")  
print(str(my_at_content))  
print(get_at_content("ATCGCGCGCATCGTY"))  
print(get_at_content("atatatatcgcgcatcagt"))  
print("AT content is " + str(get_at_content("AATTAGGCGC")))
```

Outputs?

# Arguments/Functions - AT counter

**Outputs?**

0.55

0.33

0.64

AT content is 0.5

**Round up with second argument?**



# Arguments/Functions - AT counter

```
#!/usr/bin/python3
```

```
def get_at_content(dna, sig_figs):  
    length = len(dna)  
    a_count = dna.upper().count('A')  
    t_count = dna.upper().count('T')  
    at_content = (a_count + t_count) / length  
    return round(at_content, sig_figs)
```

```
test_dna = "ATCATACATACAT"  
print(get_at_content(test_dna,1))  
print(get_at_content(test_dna,2))  
print(get_at_content(test_dna,3))
```

Output?

# Arguments/Functions - AT counter

**Outputs?**

0.8

0.77

0.769

# Arguments/Functions - AT counter

```
#!/usr/bin/python3
```

```
def get_at_content(dna, sig_figs):  
    length = len(dna)  
    a_count = dna.upper().count('A')  
    t_count = dna.upper().count('T')  
    at_content = (a_count + t_count) / length  
    return round(at_content, sig_figs)
```

```
test_dna = "ATCATACATACAT"  
print(get_at_content(test_dna,1))  
print(get_at_content(test_dna,2))  
print(get_at_content(test_dna,3))
```

**Sometimes functions have default arguments. Ex. open filename is automatically set to read. You can define default arguments**

# Multiple parameters

More than one parameter can be used when defining a function. When the function is called, we pass the required arguments for the function (as defined) or in correct order.

```
# Function definition is here
```

```
def printInfo ( name, age ):
```

```
print "Name: ", name
```

```
print "Age: ", age
```

```
return;
```

```
# Calling printinfo function
```

```
printInfo( age=50, name="miki" )
```

```
Name: miki
```

```
Age: 50
```

# Multiple parameters

More than one parameter can be used when defining a function. When the function is called, we pass the required arguments for the function (as defined) or in correct order.

```
# Function definition is here
def printInfo ( name, age ):
    print "Name: ", name
    print "Age: ", age
    return;
```

```
# Calling printinfo function
printInfo( name="miki", age=50 )
```

```
Name: miki
Age: 50
```

# Multiple parameters

More than one parameter can be used when defining a function. When the function is called, we pass the required arguments for the function (as defined) or in correct order.

```
# Function definition is here
def printInfo ( name, age ):
    print "Name: ", name
    print "Age: ", age
    return;
```

```
# Calling printinfo function
PrintInfo(50, "miki")
```

```
Name: 50
Age: miki
```

# Return values

**A function will take its arguments, do some computation and return a value to be used.**

**The “return” keyword is used.**

**A return value can be anything. Lists can be return values.**

# Function definition is here

```
def sum( arg1, arg2 ):
```

```
    "Add both the parameters and return them."
```

```
    total = arg1 + arg2
```

```
    print "Inside the function : ", total
```

```
    return total
```

# Now you can call sum function

```
total = sum( 10, 20 )
```

```
print "Outside the function : ", total
```

# Return values

A “fruitful” function is one that produces a result (or return value)

The return statement ends the function execution and “sends back” the result of the function

You don’t have to return a value, you can also print a value at the end of the function, or you can return nothing (void function).



# Importing your own programs

**Import using the from command (don't need the .py) and the function that is imported.**

```
#!/usr/bin/env python
```

```
from atContent import get_at_content
```

```
print  
(get_at_content("ATACATATATAAACCAGGGGATACAG"))
```

# To function or not to the function that is the question?

Organize your code into “paragraphs” - capture a complete thought and “define it”

Don't repeat yourself - make it work once and then reuse it

If something gets too long or complex, break up into logical parts and “encapsulate” with functions.

Make a library of common functions that you will use over and over in bioinformatics.



# To function or not to the function that is the question?

Organize your code into “paragraphs” - capture a complete thought and “define it”

Don't repeat yourself - make it work once and then reuse it

If something gets too long or complex, break up into logical parts and “encapsulate” with functions.

Make a library of common functions that you will use over and over in bioinformatics.

**IMPORT YOUR OWN FUNCTIONS  
WHEN YOU NEED TO USE THEM**



# Read and strip

There are a few ways to open and read lines. This is one of my preferred ways:

```
fh = open("file.txt")
```

```
sequence = ""
```

```
for line in fh:
```

```
    sequence += line.rstrip('\n')
```

```
fh.close
```

```
print(sequence)
```

#without header, can add if statement (class 6)

```
with open('file.txt') as f:
```

```
    sequence = [line.rstrip() for line in f]
```

```
    sequence = "".join(sequence[1:])
```

#if you have header

# Docstrings

Docstrings allow you to print out the first string right after the defined function.

#Function definition is here

```
def printme( str ):
```

```
    "This prints a passed string into this function"
```

```
    print str;
```

```
    return;
```

#Calling the printme function

```
printme("First call of user-defined function!")
```

```
printme("Second call of user-defined function")
```

```
print(printme.__doc__)
```

# Docstrings

Docstrings allow you to print out the first string right after the defined function.

#Function definition is here

```
def printme( str ):
```

```
    """ This prints a passed string into this function  
        I wrote this
```

```
    """
```

```
    print str;
```

```
    return;
```

#Calling the printme function

```
printme("First call of user-defined function!")
```

```
printme("Second call of user-defined function")
```

```
print(printme.__doc__)
```

# Testing functions

You can test your functions using assert. If the function returns correctly, it passes, else there is an assertion error.

#Function definition is here

```
def printme( str ):  
    """This prints a passed string into this function"""  
    print str;  
    return;
```

#Calling the printme function

```
printme("First call of user-defined function!")  
printme("Second call of user-defined function")
```

```
assert printme("this string") == "this string"  
assert printme("this string") == "wrong string"
```

# Quiz 19

- On canvas now