# Python 101 Functions

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## Day overview:

#### **Functions**

- 1. Purpose
- 2. The basic recipe and calling a function
- 3. Arguments
- 4. Variable scopes
- 5. Returning values from a function
- 6. Lambda (anonymous) function

#### I/O Input output

- 1. Input from user/keyboard
- 2. Reading files
- 3. Writing files
- 4. Closing files

# **Purpose**

<u>Functions</u> - pieces of code that are written one time and reused as much as desired within the program. They:

- Are the simplest callable object in python
- Perfom single related actions that can handle repetitive tasks
- Significantly reduce code redundancy and complexity, while providing a clean structure
- Decompose complex problems into simpler pieces

## **Purpose**

• Supose you have a protein sequence and want to find out the frequency of the "W" amino acid and all its positions in the sequence.  $01/10/2014\ 10:41\ AM$ 

```
for i in range (sequence_length):
    if aa_sequence[i] == "w":
        position_list.append(str(i))

p_count = float(aa_sequence.count("w"))
p_frequency = p_count/sequence_length
print "The aa 'w' has a frequency of %s and is found in the following sites: %s" % (p_frequency," ".join(position_list))
```

The aa 'w' has a frequency of 0.058823529411764705 and is found in the following sites: 13 19  $\,$ 

## **Purpose**

• Now you may want to know the same information about, say "**P**". You would need to re-write your entire code again for "**P**"...

```
aa_sequence = "mgagkvikckaafwagkplwegevappkakapca"
 2
      position list = []
 3
      sequence_length = float(len(aa_sequence))
 4
      for i in range (sequence_length):
 5
          if aa_sequence[i] == "p":
 6
              position_list.append(str(i))
 7
 8
      p_count = float(aa_sequence.count("p"))
 9
      p_frequency = p_count/sequence_length
      print "The aa 'p' has a frequency of %s and is found in the
10
      following sites: %s" % (p_frequency," ".join(position_list))
11
```

```
The aa 'p' has a frequency of 0.11764705882352941 and is found in the following sites: 17 25 26 31
```

And 19 more times to accomodate all other amino acids!!

## **Purpose**

• Using a function, the problem can be easily solved like this:

```
1
     aa_sequence = "mgagkvikckaafwagkplwegevappkakapca"
2
     def aa_statistics(sequence,aa):
3
         sequence_length,aa_positions = len(sequence),[]
4
         aa_frequency = (lambda
5
     count,length:float(count)/float(length))
6
         for i in range (sequence_length):
7
             if sequence[i] == aa:
8
                 aa_positions.append(str(i))
9
     (aa_frequency(sequence.count(aa), sequence_length), aa_positio
```

With only 7 lines of code, we are now able to provide the required information for all amino acids and for any input sequence.

## The basic recipe

• The basic steps when defining a function:

```
def name ():
    "Documentation string of the function"
    [statements]
4
```

1. "def" - Functions must start with the "def" keyword.

- Python 101: Ar 'bialog's the backers' by b
  - 3. "()" Parenthesis enclose input parameters or arguments
  - 4. ":" The code block within every function starts with a **colon** and is **indented**
  - 5. Documentation [optional] It is good practice to document your function
  - 6. "statements" The actual code block of your function

# **Function calling**

 After a function is defined, it represents nothing more than an idle piece of code, unless called. It is only when we call a function that the statements inside the function body are executed.

```
1 def print_me ():
2    "This function prints something"
3    print "Hello World"
4
```

## **Arguments**

A function can be created without arguments,

```
def print_me():
    "Example of a simple function without arguments"
    print "Hello World"

print_me()

print_me()
```

Hello World

or using the following types of arguments:

- Required arguments
- Default arguments
- Variable length arguments

## **Arguments**

## **Required arguments**

```
def aa_frequency (sequence,aa):
    "This function takes exactly two arguments"
    sequence_length = len(sequence)
    aa_frequency =
    float(sequence.count(aa))/float(sequence_length)
    print aa_frequency
```

• When calling for a function with required arguments, the **exact** same number of arguments must be specified, no more and no less.

```
▼ 1 def aa_frequency (sequence,aa): #folded
6 aa_frequency ("AWKLCVPAMAKNENAW","K")
7

0.125
```

## **Arguments**

#### **Required arguments**

• It is also possible to provide previously named variables as arguments

• If you specify a different number of arguments, however

```
1 H_sapiens_aa = "AWKLCVPAMAKNENAW"
▼ 2 def aa_frequency (sequence,aa): #folded
```

```
TypeError: aa_statistics() takes exactly 2 arguments (3 given)
```

## **Arguments**

#### Variable length arguments

• Placing an asterisk (\*) before the variable name will store the arguments in a tuple

```
def concatenate (*sequences):
 2
          " This one can take a variable number of arguments,
3
      even 0"
 4
          concatenated_sequences = ""
 5
          for i in sequences: # You can iterate over the tuple,
 6
              concatenated_sequences += i
 7
          if len(sequences) >= 2:
8
              first_sequences = sequences[:2] # and slice its
9
      items
10
              print concatenated_sequences, first_sequences
11
      concatenate("GTCCG","AGTCG","AGTAG","AGTGA")
13
      concatenate() # In this case the tuple "sequences" is empty
```

```
GTCCGAGTCGAGTAGAGTGA ('GTCCG', 'AGTCG')
```

## **Arguments**

#### **Default arguments**

• Arguments can also have default values, by assigning those values to the argument keyword with the assign ("=") symbol.

```
1 def codon_count (Sequence,
2 StopCodon="TAA",StartCodon="ATG"):
```

• The function will assume the default value if the argument keyword is not specified when calling the function.

## **Arguments**

#### Using argument keywords

• When calling a function, the order of the arguments can be changed by using the argument's keyword and the assign ("=") symbol.

 Note that this is necessary if you would like to change only the second default argument, and leave the first with the default value

```
1 H_sapiens = "AGCTAGTCGTAGCATGATTAACGTAGGCTATACTACTAAATGRC"
    def codon_count (Sequence,
        StopCodon="TAA", StartCodon="ATG"): #folded
        codon_count (H_sapiens, StartCodon="ATT")
```

2 1

## **Arguments**

## Considerations when combining different argument types

• **Default** arguments should come after **required** arguments

```
def name (required,required,
    (...),default=value,default=value,(...)):
    [...code block...]
```

• **Variable length** arguments should be used only once and be always last. There is also no point in using them with **default** arguments.

```
def name (required,required,(...),*varible_length):
     [...code block...]
4
```

#### Namespaces or scope of variables

When writting a program, it is extremely important to know the difference between the **local** and **global** scope of the variables

#### Glogal variables

• Variables defined outside functions or other objects (i.e., classes) are **global** variables - they are accessible throughout most of the program, even by functions.

```
1  sequence = "ACGTGTGC"
2  def print_me():
3    print sequence
4  
5   print_me()
6
```

ACGTGTG

 $\bullet$  To change the contents of a  ${\bf global}$  variable in a function, we can use the global keyword

#### Namespaces or scope of variables

#### Local variables

• By default, all variables defined inside a function (including argument keywords) are **local** variables - they are not accessible by the whole program, only within the function where they are declared.

```
1    sequence = "TTTTT"
2    def print_me():
3        sequence = "AAAAAA"
4        print sequence
5    print_me()
7
AAAAAA
```

#### Return

The  $\it return$  keyword is used to return values from a function, which can then be assigned to new variables that are accessible to the whole program

```
H_sapiens_lc1 =
      "AGCTAGTCGTAGCATGATTAACGTAGGCTATACTACTAAATGRC"
 3
      H_sapiens_lc2 =
 4
5
      \verb"CGTAGTCGTAGTTTGCAGTGCGCTGATCGTAGTCGATGCTGTGT"
 6
      def concatenate (*sequences):
           concatenated_sequence = ""
 7
 8
           for i in sequences:
 9
               concatenated_sequence += i
10
           return concatenated_sequence
11
      new_sequence = concatenate(H_sapiens_lc1,H_sapiens_lc2)
12
           # And now we can use the output of a function, as the
13
      input of another
14
15
      def codon_count (Sequence,
      StopCodon="TAA", StartCodon="ATG"):
    stop_count = Sequence.count(StopCodon)
16
17
18
          start_count = Sequence.count(StartCodon)
19
          print stop_count, start_count
      codon_count (new_sequence)
```

```
2 3
```

#### Return

#### **Returning multiple values**

• Functions can return multiple values

```
def codon_count (Sequence,
    StopCodon="TAA",StartCodon="ATG"):
        stop_count = Sequence.count(StopCodon)
        start_count = Sequence.count(StartCodon)
        return stop_count, start_count # Returns a tuple with
        two items
        # OR
        # return [stop_count, start_count] -> Returns a list
        with two items
```

• And these values can be assigned to multiple variables

```
1 H_sapiens = "AGCTAGTCGTAGCATGATTAACGTAGGCTATACTAAATGRC"
    def codon_count (Sequence,
        StopCodon="TAA", StartCodon="ATG"): #folded
        stop, start = codon_count(H_sapiens)
        print stop, start
        start = codon_count(H_sapiens)[1] # You can even select the
        variable(s) you want
        print start
```

2

## Return

## Functions always return something

If a function does not contain the return keyword, it will return None

```
1  def print_me():
2    a = 2+2
3    print_me() == None
6
True
```

#### Lambda (anonymous) functions

<u>Lambda</u> is an anonymous (unnamed) function that is used primarily to write very short functions that are a hassle to define in the normal way. Where a regular function would do:

```
1 def add(a,b):
2 print a+b
3
4 add(4,3)
5
```

a lambda function:

```
1 print (lambda a,b: a+b)(4,3)
2
```

```
1  RNA = ["AUGAUU","AAUCGAUCG","ACUAUG","ACUAUG"]
2  DNA = map(lambda sequence: sequence.replace("U","T"), RNA)
5  
["ATGATT","AATCGATCG","ACTATG","ACTATG"]
```

# Wrap up

So, we have covered thus far:

- How to define functions using the *def* keyword
- How to call a function
- The three main types of arguments a function can take: Required , variable length and arguments
- The local and global scope of variables
- The usage of the *return* keyword to return values from functions
- Lambda functions