Utrecht
Bioinformatics
Center

Functions and Modules

Day 3 – Introduction to Python

Functions

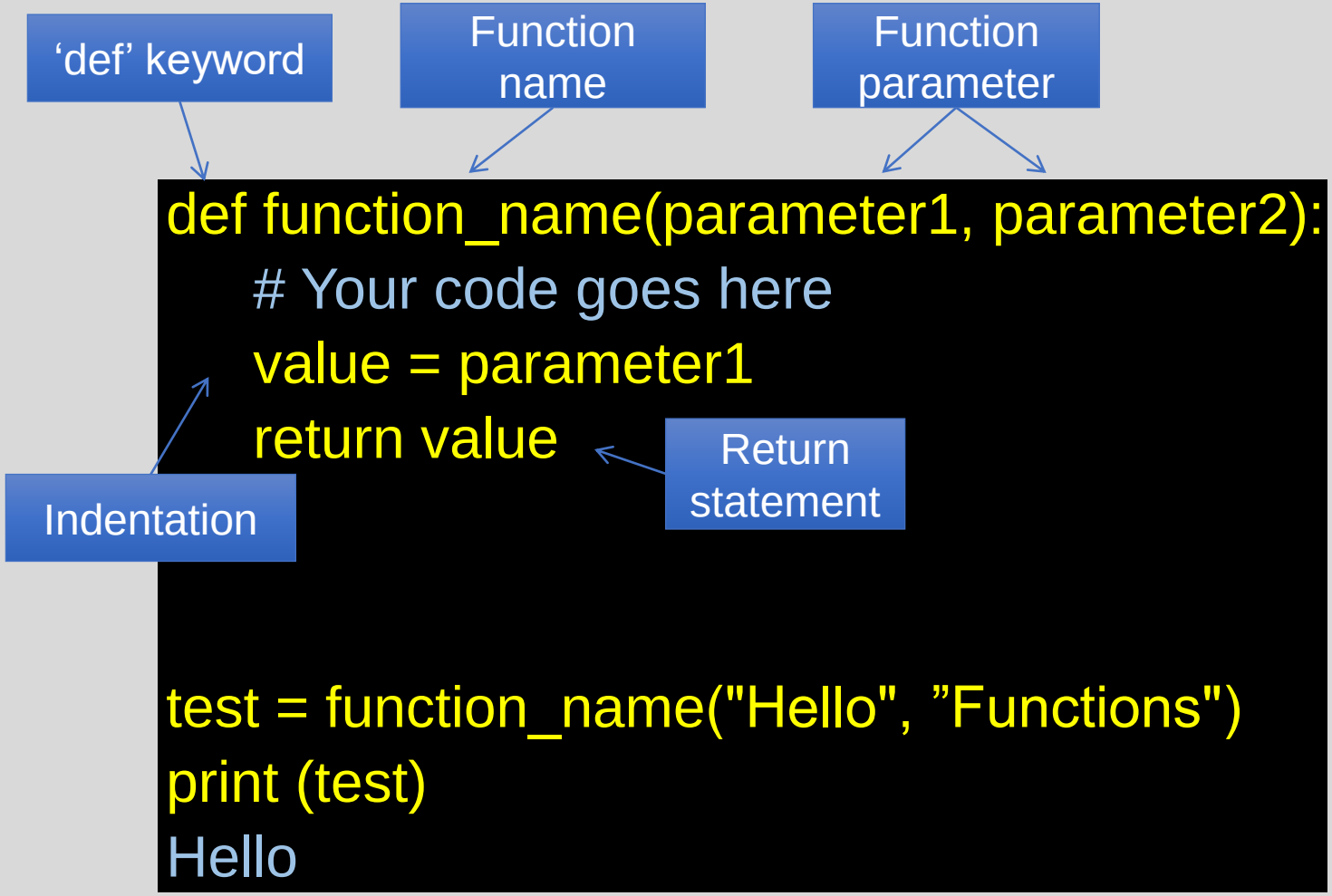
- Pieces of code to execute a specific task

- Advantages:

  - Reduce duplication of code
  - Complex problems into simpler pieces
  - Improve clarity of the code

Functions

- Pieces of code to execute:




```
def reverse_it(stuff_i_type):
        text_out = stuff_i_type[::-1]
        return text_out

andersom = reverse_it(“Apple”)
print (andersom)
epplA
```

Functions

- also work without parameters:

```
def hello_functions():
    value = "Hello Functions!"
    return value

hello = hello_functions()
print (hello)
Hello Functions!
```

- can include loops:

```
def print_list(my_list):
    for item in my_list:
    print (item)

test_list = [1,2,3]
print_list(test_list)
1
2
3
```

```
def print_even(list_of_numbers):
        for each_number in list_of_numbers
            if(each_number%2.0 == 0):
                print (each_number)

some_numbers = range(10)
print_even(some_numbers)
2
4
6
8
10
```

```
def sum_list(list):  # define function sum_list, with parameter “list”
        total = 0 # make variable “total” with value 0
        for item in list: # loop over the list and
            total += item # add each value of the list to total
        return total # return total

test = [1,2,3]
print (sum_list(test))
6
```

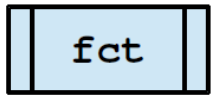
Functions

Function Definition

function name (identifier)

named parameters

```
def fct(x,y,z):
    """documentation"""
    # statements block, res computation, etc.
    return res
```



result value of the call, if no computed result to return: `return None`

parameters and all variables of this block exist only *in* the block and *during* the function call (think of a "black box")

Advanced: `def fct(x,y,z,*args,a=3,b=5,**kwargs):`

**args variable positional arguments (→`tuple`), default values,*
***kwargs variable named arguments (→`dict`)*

Function Call

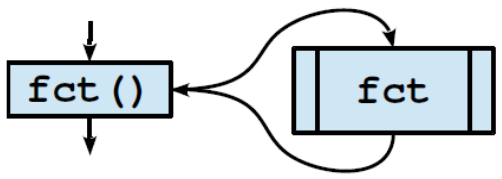
```
r = fct(3,i+2,2*i)
```

storage/use of returned value

one argument per parameter

this is the use of function name *with parentheses* which does the call

Advanced:
**sequence*
***dict*



Functions and Lists

- Now lets try to write a function to get all codons from the sequence: 'ACGATCGATCGTACGATCGATACG’

```
def get_codons(seq):
    # Make an empty list that will contain codons
    # Loop over seq considering 3 letters at a time
    for codon in range(?,len(seq),?):
            # Append your list with the codon from seq
            # You can return the list containing codons
    return # list with codons


sequence = 'ACGATCGATCGTACGATCGATACG'
print (get_codons(sequence))
```

Functions and Dictionaries

- Write a function to calculate number of times a codon is used in the sequence: 'ACGATCGATCGTACGATCGATACG’.

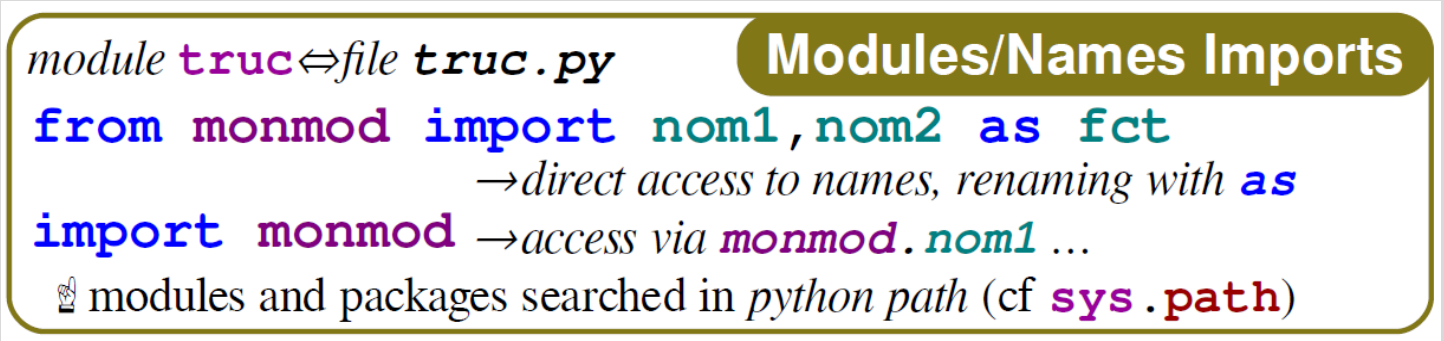
```
def get_counts(seq):
    # Make an empty dictionary to store codons
    for i in range(0,len(seq),3):
            codon = seq[i:i+3]
            # use the codons as keys and the counts as values
            # If the codon is already within your
            # dictionary then increase its count
            # otherwise add it to the dictionary with count = 1
    return # dictionary with codon counts


sequence = 'ACGATCGATCGTACGATCGATACG'
print (get_counts(sequence))
```

Modules

- Toolbox or kit which is a collection of functions
- A file consisting of Python code.
- A module allows you to logically organize your code.
  - To split it into several files for easier maintenance.
  - To reuse that handy function that you've written in several programs, without copying it into every script.
- Package is a collection of modules

Built-in Modules

- Some important (default) modules:
  - math
  - itertools
  - random
  - sys
  - dir
  - And many more: https://docs.python.org/3/library/index.html



Modules – making you own

- You will write a function to count the number of times a codon is used.
- You can store such functions in a file so that we can use them later.
- We have stored get_counts and translate_dna into a file called dna_tools.py

- You can import a module in Python using its file name:

```
import dna_tools #This calls the file dna_tools.py stored in the same folder


dna_tools.get_counts('ATCGATCATGAC')
```

Modules – rename

- You can also rename a module when you import it in Python and call the function inside the module.

```
import dna_tools as tools
tools.get_counts('ATCGATCATGAC')
```

- Now try to load your dna_tools module and try both functions

Exercises