Scientific Programming Practical 7

Introduction

Functions: just a reminder

A function is a block of code that has a name and that performs a task. A function can be thought of as a box that gets an input and returns an output.

```
The basic definition of a function is:

def function_name(input) :
    #code implementing the function
    ...
    return return_value
```

- 1. **Reduce code duplication**: put in functions parts of code that are needed several times in the whole program so that you don't need to repeat the same code over and over again;
- 2. **Decompose a complex task**: make the code easier to write and understand by splitting the whole program in several easier functions

Passing parameters from command line

Python provides the module **sys** to interact with the interpreter.

sys.argv is a list representing all the arguments passed to the python script from the command line.

```
"""Test input from command line"""
      import sys
      if len(sys.argv) != 4: #3 + 1 (the command!)
          print("Dear user, I was expecting three parameters. You gave me ", len(sys.argv)-
          sys.exit(1)
      else:
          for i in range(0, len(sys.argv)):
              print("Param {}: {} ({})".format(i, sys.argv[i], type(sys.argv[i])))
                                                                       1: bash
                                  TERMINAL
biancol@bluhp:~/Google Drive/work/courses/sciprolab1/exercises$
```

Check out: https://docs.python.org/3/

Passing parameters from command line

```
"""Test input from command line"""
      import sys
      if len(sys.argv) != 4: #3 + 1 (the command!)
          print("Dear user, I was expecting three parameters. You gave me ", len(sys.argv)-1)
          sys.exit(1)
      else:
          for i in range(0, len(sys.argv)):
              print("Param {}: {} ({})".format(i, sys.argv[i], type(sys.argv[i])))
                                                                         2: Python
                    DEBUG CONSOLE
                                   TERMINAL
/usr/bin/python3.6 "/home/biancol/Google Drive/work/courses/sciprolab1/exercises/systest.py"
biancol@bluhp:~/Google Drive/work/courses/sciprolab1/exercises$ /usr/bin/python3.6 "/home/biancol/Google Dri
ve/work/courses/sciprolab1/exercises/systest.py"
Dear user, I was expecting three parameters. You gave me 0
biancol@bluhp:~/Google Drive/work/courses/sciprolab1/exercises$ /usr/bin/python3.6 "/home/biancol/Google Dri
ve/work/courses/sciprolab1/exercises/systest.py" param1 2 param3
Param 0: /home/biancol/Google Drive/work/courses/sciprolab1/exercises/systest.py (<class 'str'>)
Param 1: param1 (<class 'str'>)
Param 2: 2 (<class 'str'>)
Param 3: param3 (<class 'str'>)
biancol@bluhp:~/Google Drive/work/courses/sciprolab1/exercises$
```

Argparse is a command line parsing module which deals with **positional arguments** and **optional arguments**.

```
biancol@bluhp:~/Google Drive/work/courses/sciprolab1$ ls --help
Usage: ls [OPTION]... [FILE]...
List information about the FILEs (the current directory by default).
Sort entries alphabetically if none of -cftuvSUX nor --sort is specified.
Mandatory arguments to long options are mandatory for short options too.
                            do not ignore entries starting with .
 -A, --almost-all
                            do not list implied . and ..
     --author
                            with -l, print the author of each file
 -b, --escape
                            print C-style escapes for nongraphic characters
     --block-size=SIZE
                            scale sizes by SIZE before printing them; e.g.,
                              '--block-size=M' prints sizes in units of
                              1,048,576 bytes; see SIZE format below
                            do not list implied entries ending with ~
 -B, --ignore-backups
```

Six steps:

1. Import the module

import argparse

2. Create the parser object

parser = argparse.ArgumentParser(description="This is the description of the program")

3. Add positional arguments

4. Add optional arguments

5. Parse the arguments

```
args = parser.parse args()
```

6. Retrieve and process the arguments

```
myArgName = args.arg_name
myOptArg = args.optional arg
```

Example: Let's write a program that reads and prints to screen a text file specified by the user. Optionally, the file might be compressed with gzip to save space. The user should be able to read also gzipped files. Hint: use the module gzip which is very similar to the standard file management method

```
import argparse
import gzip
parser = argparse.ArgumentParser(description="""Reads and prints a text file""")
parser.add argument("filename", type=str, help="The file name")
parser.add argument("-z", "--gzipped", action="store true",
                    help="If set, input file is assumed gzipped")
args = parser.parse args()
inputFile = args.filename
fh = ""
if(args.gzipped):
    fh = gzip.open(inputFile, "rt")
else:
    fh = open(inputFile, "r")
for line in fh:
    line = line.strip("\n")
    print(line)
fh.close()
```

```
import argparse
import gzip
parser = argparse.ArgumentParser(description="""Reads and prints a text file""")
parser.add argument("filename", type=str, help="The file name")
parser.add argument("-z", "--gzipped", action="store true",
                    help="If set, input file is assumed gzipped")
args = parser.parse args()
inputFile = args.filename
fh = ""
if(args.gzipped):
    fh = gzip.open(inputFile, "rt")
else:
    fh = open(inputFile, "r")
for line in fh:
    line = line.strip("\n")
    print(line)
fh.close()
```

```
import argparse
import gzip
parser = argparse.ArgumentParser(description="""Reads and prints a text file""")
parser.add argument("filename", type=str, help="The file name")
parser.add argument("-z", "--gzipped", action="store true",
                    help="If set, input file is assumed gzipped")
args = parser.parse args()
inputFile = args.filename
fh = ""
if(args.gzipped):
    fh = gzip.open(inputFile, "rt")
else:
    fh = open(inputFile, "r")
for line in fh:
    line = line.strip("\n")
    print(line)
fh.close()
```

```
biancol@bluhp:~/Google Drive/work/courses/sciprolab1$ python3 exercises/readFile_gz.py file_samples/textFile.txt
Hi everybody,
This is my first file
and it contains a total of
four lines!
```

```
biancol@bluhp:~/Google Drive/work/courses/sciprolab1$ python3 exercises/readFile_gz.py file_samples/textFile.gz -z
Hi everybody,
This is my first file
and it contains a total of
four lines!
```

https://docs.python.org/3/library/index.html







Previous topic

10. Full Grammar specification

Next topic

1. Introduction

This Page

Report a Bug Show Source

The Python Standard Library

While The Python Language Reference describes the exact syntax and semantics of the Python language, this library reference manual describes the standard library that is distributed with Python. It also describes some of the optional components that are commonly included in Python distributions.

Python's standard library is very extensive, offering a wide range of facilities as indicated by the long table of contents listed below. The library contains built-in modules (written in C) that provide access to system functionality such as file I/O that would otherwise be inaccessible to Python programmers, as well as modules written in Python that provide standardized solutions for many problems that occur in everyday programming. Some of these modules are explicitly designed to encourage and enhance the portability of Python programs by abstracting away platform-specifics into platform-neutral APIs.

The Python installers for the Windows platform usually include the entire standard library and often also include many additional components. For Unix-like operating systems Python is normally provided as a collection of packages, so it may be necessary to use the packaging tools provided with the operating system to obtain some or all of the optional components.

In addition to the standard library, there is a growing collection of several thousand components (from individual programs and modules to packages and entire application development frameworks), available from the Python Package Index.

- 1. Introduction
- · 2. Built-in Functions
- · 3. Built-in Constants
 - 3.1. Constants added by the site module
- · 4. Built-in Types
 - 4.1. Truth Value Testing
 - 4.2. Boolean Operations and, or, not
 - 4.3. Comparisons
 - 4.4. Numeric Types int, float, complex
 - 4.5. Iterator Types
 - 4.6. Sequence Types list, tuple, range
 - 4.7. Text Sequence Type str
 - 4.8. Binary Sequence Types bytes, bytearray, memoryview
 - 4.9. Set Types set, frozenset

http://qcbprolab.readthedocs.io/en/latest/practical7.html

Exercises

 Modify the program of Exercise 4 of Practical 6 in order to allow users to specify the input and output files from command line. Then test it with the provided files. The text of the exercise follows:

Write a python program that reads two files. The first is a one column text file (contig_ids.txt) with the identifiers of some contigs that are present in the second file, which is a fasta formatted file (contigs82.fasta). The program will write on a third, fasta formatted file (e.g. filtered_contigs.fasta) only those entries in contigs82.fasta having identifier in contig_ids.txt.

Show/Hide Solution

2. Cytoscape is a well known tool to perform network analysis. It is well integrated with several online databases housing for example protein-protein interactions like EBI's IntAct. It is also able to read and write a very simple text file called self to represent interactions between the nodes of a network. Sif formatted files are tab separated (\tau t) and each line represents a connection between the nodes of the network. For example:

```
nodel interaction1 node2
node1 interaction2 node3
node2 interaction1 node3
```

represents two types of interactions between node1, node2 and node3. Normally nodes are represented as circles in a network (graph) and interactions as lines (that can be of different kinds) connecting nodes (edges). The following is an extract from the file pka.sif that has been downloaded by Cytoscape from the database IntAct and represents the interactions of the Protein Kinase A (PKA) of E.coli:

P75742 EBI-9168813 P76594