



Introduction to Python course – part II

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Defining functions

Often you need to repeat the same operation multiple times with different input variables. Instead of writing the same code each time, you can use a **function**, which is defined by the keyword “**def**”, followed by the function name and the input variables, enclosed in round brackets.

For sake of clarity, functions should be accompanied by a short but comprehensive description of what they do. This can be a comment on a single or multiple lines, preceded and closed by **triple quotes**.

Variables declared outside of a function are called **global** and are anywhere in the code. Variables declared inside a function are called **local** and are only accessible to the current scope.

```
[2]: def introduction(inpname):  
    """  
    This function prints the input global variable 'inpname'  
    and combines it with the local variable 'age' in a output  
    string of greeting.  
    """  
  
    age = 33  
    print('Hello, my name is', first_name, 'and I am', age, 'years old')
```

```
[3]: whoami = 'Silvia'  
    introduction(whoami)  
  
Hello, my name is Silvia and I am 33 years old
```

```
[4]: whoami = 'John'  
    introduction(whoami)  
  
Hello, my name is John and I am 33 years old
```

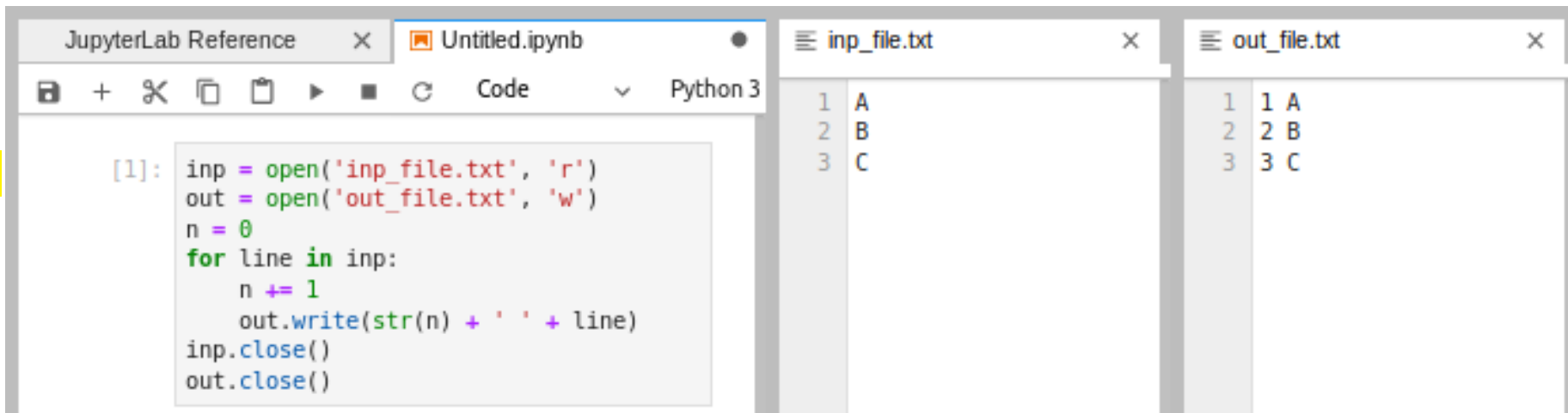
```
[5]: print(age)  
  
-----  
NameError                                Traceback (most recent call last)  
<ipython-input-5-5f7a7c5b2c60> in <module>  
----> 1 print(age)  
  
NameError: name 'age' is not defined
```

Reading and writing files

The `open()` built-in function takes as input a file name and a mode ('*r*' for reading, '*w*' for writing, '*a*' for appending to an existing file, and others...), and returns a file object. By default, files are opened in text mode ('*t*'), but binary files could be opened too, using the '*b*' mode. Once opened, a file must be closed with the `close()` function (**mode A**).

Alternatively, it can be opened using the `with` and `as` statements, which will close it once the code block is finished (**mode B**).

mode A →



The screenshot shows a JupyterLab interface with three tabs: 'JupyterLab Reference', 'Untitled.ipynb', and two file tabs 'inp_file.txt' and 'out_file.txt'. The 'Untitled.ipynb' tab contains the following Python code:

```
[1]: inp = open('inp_file.txt', 'r')
      out = open('out_file.txt', 'w')
      n = 0
      for line in inp:
          n += 1
          out.write(str(n) + ' ' + line)
      inp.close()
      out.close()
```

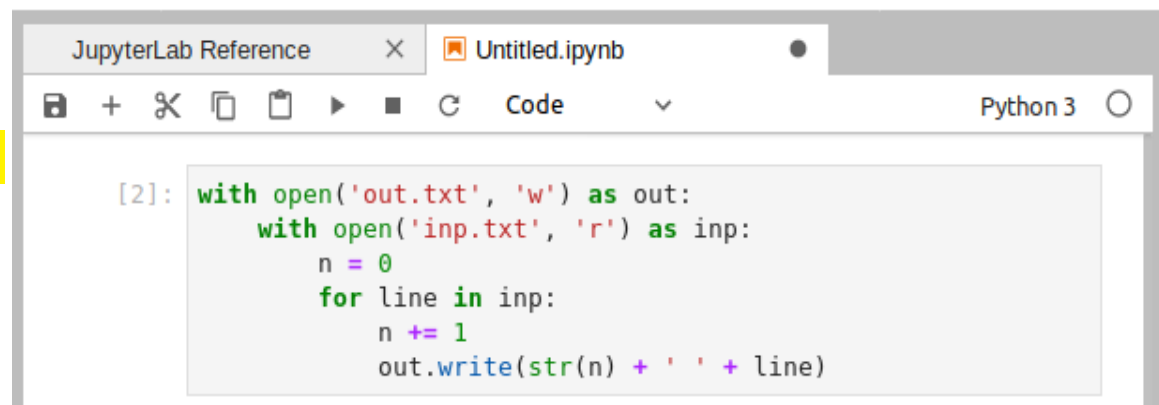
The 'inp_file.txt' tab shows the input file content:

```
1 A
2 B
3 C
```

The 'out_file.txt' tab shows the output file content:

```
1 1 A
2 2 B
3 3 C
```

mode B →



The screenshot shows a JupyterLab interface with two tabs: 'JupyterLab Reference' and 'Untitled.ipynb'. The 'Untitled.ipynb' tab contains the following Python code:

```
[2]: with open('out.txt', 'w') as out:
      with open('inp.txt', 'r') as inp:
          n = 0
          for line in inp:
              n += 1
              out.write(str(n) + ' ' + line)
```

Understanding errors

Programmers spend most of the time debugging **errors**, so it's crucial to know and understand them.

These are some of the errors you will find more frequently.

```
[1]: while True
      print('Hello!')
```

```
File "<ipython-input-1-06120368fd1d>", line 1
      while True
            ^
SyntaxError: invalid syntax
```

```
[2]: with open('inp.txt') as i:
      pass
      i.readline()
```

```
-----
ValueError                                Traceback (most recent call last)
<ipython-input-2-3508ab0af352> in <module>
      1 with open('inp.txt') as i:
      2     pass
----> 3 i.readline()

ValueError: I/O operation on closed file.
```

```
[3]: l = [5, 6, 7]
      l[3]
```

```
-----
IndexError                                Traceback (most recent call last)
<ipython-input-3-e72eec77f208> in <module>
      1 l = [5, 6, 7]
----> 2 l[3]

IndexError: list index out of range
```

```
[4]: d = {'a': 1, 'b': 5}
      print(d['c'])
```

```
-----
KeyError                                  Traceback
<ipython-input-4-c5b7615ea0e1> in <module>
      1 d = {'a': 1, 'b': 5}
----> 2 print(d['c'])

KeyError: 'c'
```

```
[5]: 5 * (3 / 0)
```

```
-----
ZeroDivisionError                        Traceback (most recent call last)
<ipython-input-5-4a4bc964b9b4> in <module>
----> 1 5 * (3 / 0)

ZeroDivisionError: division by zero
```

```
[6]: 5 + '3'
```

```
-----
TypeError                                Traceback (most recent call last)
<ipython-input-6-89b0d6b4af2b> in <module>
----> 1 5 + '3'

TypeError: unsupported operand type(s) for +: 'int' and 'str'
```

Handling exceptions

Even if a given piece of code is syntactically correct, it may still cause an error when you attempt to execute it. These errors are called exceptions and can be handled using the **try** and **except** statements.

The code first attempts to run the block in the try clause. If no exception occurs, the except clause is skipped, otherwise it is executed. The **finally** clause is executed in any case.

If another exception, not captured by the except clause, occurs, then the unhandled exception will be displayed and the execution stopped.

```
[1]: def add_two(inp_num):  
      """This function adds 2 to the input number. """  
  
      try:  
          print(inp_num + 2)  
      except TypeError:  
          print("Oops, that was not a valid number")  
      finally:  
          print('Done!')
```

```
[2]: add_two(3)
```

```
5  
Done!
```

```
[3]: add_two('3')
```

```
Oops, that was not a valid number  
Done!
```

Practical session



Practical session 3 – exercises

- 1) Write a function that takes as input a DNA sequence (e.g. **'ATGGTCA'**) and prints the string **'This DNA sequence is N base pair long'**, where N is the length of your input sequence (for the example above, it would be 7).
- 2) Create a file **inp.txt** containing the following RNA sequences, one per line: **'UGAAAC'**, **'GGGUCUUUU'**, **'GUUAAAACAACCCU'**. Read this file and write a new file **out.txt** containing the length of each sequence in **inp.txt**, one per line. Tip: each input line contains the extra character **'\n'**, which does not have to be counted, but that has to be written in the output file, in order to have separate lines.
- 3) Given the dictionary **bases = {'A': 'A', 'C': 'C', 'G': 'G', 'T': 'U'}** and the DNA sequence **dna = 'ATCKNGA'**, convert it to a new RNA sequence called **rna** using **bases**. Catch the **KeyError** exception raised by **'K'** and **'N'** and print the error message **'Wrong base found!'** together with the base that raised the exception.

Practical session 3 – solutions

1) `def seq_length(dna):`

`"""This function prints a string saying the length of 'dna'. """`

`print('This DNA sequence is', len(dna), 'base pair long')`

`seq_length('AAATTGGGG') # the result must be: This DNA sequence is 9 base pair long`

2) `inp = open('inp.txt', 'r')`

`out = open('out.txt', 'w')`

`for row in inp:`

`out.write(str(len(row) - 1) + '\n')`

`inp.close()`

`out.close()`

3) `bases = {'A': 'A', 'C': 'C', 'G': 'G', 'T': 'U'}`

`dna = 'ATCKNGA'`

`rna = ''`

`for letter in dna:`

`try:`

`rna += bases[letter]`

`except KeyError:`

`print('Wrong base found!', letter)`

`print(rna)`

Coding style

From its invention, Python's philosophy emphasized **code readability**, making it easier to be read and understood by others, when code is shared.

The **Python Enhancement Proposals (PEPs)**, reviewed by the Python community and steering council, are the primary mechanism for proposing major new features and taking new design decisions.

One of this proposals, **PEP 8**, focuses on coding style and is a style guide that promotes readable and eye-pleasing code development. Here are some of the main points:

- Use **4-space indentation** instead of tabs (which introduce confusion)
- Code lines should not exceed **80 characters** (to help users with small displays)
- Functions and classes should be separated by **blank lines**
- **Comments** should be on separate lines, rather than with code
- Use **docstrings** as much as possible (to enhance code documentation and readability)
- Use **spaces** around operators and after commas
- **Name classes and functions consistently**, using UpperCamelCase for classes and lowercase_with_underscores for functions and methods.
- Code should always use **UTF-8** (or **ASCII** in Python 2), don't use other characters (to make code readable by people speaking other languages and using different keyboards)
- **Imports** should be on separate lines

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Python >>> Python Developer's Guide >>> PEP Index >>> PEP 8 -- Style Guide for Python Code

PEP 8 -- Style Guide for Python Code

PEP:	8
Title:	Style Guide for Python Code
Author:	Guido van Rossum <guido at python.org>, Barry Warsaw <barry at python.org>, Nick Coghlan <ncoghlan at gmail.com>
Status:	Active
Type:	Process
Created:	05-Jul-2001
Post-History:	05-Jul-2001, 01-Aug-2013

Useful resources

The topics presented in this course can be found in more detail at <https://docs.python.org/3/tutorial/> , although I would really recommend diving deeper into Python. Here is a list of some useful resources for learning Python:

- **LinkedIn Learning:** free for Uni Oxford members! <https://www.linkedin.com/learning/>
- **Learn Python:** <https://www.learnpython.org/>
- **DataCamp:** <https://www.datacamp.com/>
- **Coursera:** <https://www.coursera.org/>
- **edX:** <https://www.edx.org/learn/python>

...and for practicing Python (problems with solutions):

- **Rosalind:** <http://rosalind.info/problems/list-view/>
- **Leetcode:** <https://leetcode.com/problemset/all/>
- **CodingBat:** <https://codingbat.com/python>
- **W3 Resource:** <https://www.w3resource.com/python-exercises/>
- **Practice Python:** <https://www.practicepython.org/>

Thank you for your attention!

Questions?

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