## **Python for Data Analysis**

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## Intro

### Data types

Basic materials (Wood / Stone)

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#### **Methods**

Associated activities (Wood sawing / Stone chipping)

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#### **Data structures**

Building structures (Wooden wall / Stone wall)

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Associated activities (Wood sawing / Stone chipping)

#### **Data structures**

Building structures (Wooden wall / Stone wall)

#### Flow control

Command builders ('Build me a 10 foot stone wall!')

# **Data types**

### Data types: numbers

- int (Integers): whole numbers
- float: floating-point numbers

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#### **Variables**

- a = 4
- b = 2
- a + b 6

### Data types: Boolean

- Switch with 2 states: True or False
- Answers to a yes/no question

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#### **Comparisons**

- 3 == 4 False
- 2 < 4 True

### **Data types: Strings**

- Strings of letters without inherent meaning
- Always in quotation marks

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#### **String manipulation**

- name = "Andreas"
- last = "Weller"
- name + last"AndreasWeller"
- "II" in last True
- "First name: %s, last name: %s." % (name, last)
   First name: Andreas, last name: Weller.

## **Slicing**

- Ways to retrieve data from an iterable
- Scheme: iterable [ first item : item AFTER last item : step]

```
a n d r e a s
0 1 2 3 4 5 6
```

#### Slicing

- name = "Andreas"
- name[3:5]"re"
- name[3:] "reas"
- name[::2]"Ades"

### **Functions and Methods**

### **Methods**

- Methods are the verbs of Python
- Scheme: object.method(argument)

English	Python	Abstract
"The dog sleeps"	dog.sleep()	obj.method()
"The dog jumps high"	dog.jump("high")	obj.method(arg)
"The dog eats a cat"	dog.eat(cat)	obj.method(obj)

### **Methods II**

Methods are specific to each data type.

#### **String methods**

- name = "Andreas"
- name.upper()"ANDREAS"
- name.lower()"andreas"
- name.strip("as")"Andre"
- name.replace("dre", "XXX") "AnXXXas"

### **Data structures**

#### Lists

- Lists are ordered arrays of objects.
- The items in a list can be of any type.
- Lists are defined in square brackets.

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#### **List methods**

- students = ["Pauline","Carme","Anthony"]
- "Andreas" in students False
- students[1]"Carme"
- students.append("Pauline")
- students ["Pauline","Carme","Anthony","Pauline"]
- students.count("Pauline")2

### **Dictionaries**

- Dictionaries are a collection of unordered {key:value} pairs
- Keys need to be of type str, values can be anything
- Dictionaries defined in curly brackets, but selected in square brackets

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#### **Dictionary methods**

- grades = {"Pauline":2, "Carme":4, "Anthony":1}
- grades["Carme"]4
- grades["Reem"] = 1
- grades {"Pauline":2, "Reem":1, "Carme":4, "Anthony":1}
- grades.keys() ["Pauline","Reem","Carme","Anthony"]

### Flow control

## Simple scripting

- A script is simply a text file ending in ".py"
- Scripts are executed on the commandline
- 'python script.py'

```
# outputs "Hello World!"

message = "Hello_World!"

print message
```

### If statements

- Run some code IF a condition is True
- Code to be run is indented with tabs

```
students = 8
max_students = 5

if (students > max_students):
    print "Too_many_students!"

else:
    print "Right_amount_of_students!"

print "Welcome_to_the_class!"
```

### For loops

- Run some code FOR each item in an iterable
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```
# greet all students in class
students = ["Pauline","Carme","Anthony"]
for name in students:
    message = "Hello_" + name + "!"
    print message
print "Welcome_to_the_class!"
```

# Working with files

## File manipulation I

```
# read file as a list of rows
handle = open("quasar.tsv", "r")
rows = handle.readlines()
handle.close()
# print rows
for row in rows:
    print row
# write file
output = open("test.txt", "w")
output.write("This,,is,,a,,test.")
output.close()
```

## File manipulation II

```
# read file as a list of rows
handle = open("quasar.tsv", "r")
rows = handle.readlines()
handle.close()
# print chromosome and position of the first sample
for row in rows:
    f = row.split("\t")
    sample = f[0]
    chrom = f[1]
    pos = f[2]
    if sample == "Q2PL1 A01 v1 T":
        print chrom, pos
```

### File manipulation III

- The WITH statement is the best way to open files
- Python walks through the rows one by one, without loading the whole file into memory

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- Python walks through the rows one by one, without loading the whole file into memory

```
# print chromosome and position of the first sample
with open("quasar.tsv") as variant file:
    for row in variant file:
        f = row.split("\t")
        sample = f[0]
        chrom = f[1]
        pos = f[2]
        if sample == "Q2PL1 A01 v1 T":
            print chrom, pos
```

#### **Modules**

- Many useful modules are not loaded by default.
- We can use any Python code in our script by importing it.
- The module exists as an object with methods to call

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```
# Use the os (operating system) module
# to read all files in a folder as a list
import os
files = os.listdir("./")
# Use the sys module to parse command line arguments
import sys
argument1 = sys.argv[1]
argument2 = sys.argv[2]
```

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- They are define using the **def** keyword.

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```
# define a function that greets the class
def greet_class(all students):
    for student in all students:
        print "Good, morning, %s!" % student
# use the function
all students = ["Carme", "Suska"]
greet class(all students)
```

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```
# define a function that greets the class
def greet class(all students):
    for student in all_students:
        print "Good, morning, %s!" % student
    return len(all students)
# use the function
all students = ["Carme", "Suska"]
student count = greet_class(all_students)
print "l_have_%s_students." % student count
```

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```
# define a function that greets the class
def greet class(all students):
    for student in all_students:
        print "Good, morning, %s!" % student
    return len(all students)
# use the function
my students = ["Carme", "Suska"]
student_count = greet_class(my_students)
print "I,have, students." % student count
```