

Data Handling in Python

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RDM Support | University Library | IT Services



Today's schedule:

09.30-09.40	Raw data handling, setting the stage! [presentation]
09.40-10.50	Practical Data handling in Python [Jupyter notebook]
	- 10 minute coffee break -
11.10-11.30	Formatting and commenting practices [presentation]
11.20-11.30	Anaconda: ultra quick tour [Jupyter and Spyder]
11.30-12.00	Final Exercise: comment and format the code [Spyder]

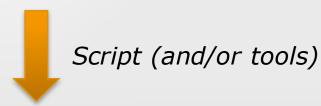


Raw data handling

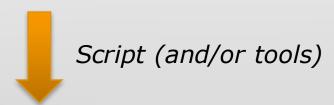


Data analysis pipeline

RAW DATA



Clean data



Analyzed data (results)



Keep copies of your data and your scripts at every step!

Data1 -> RAW DATA

Script (and/or tools) <- Script1

Data2 -> Clean data

Script (and/or tools) <- Script2

Data3 -> Analyzed data (results)



Cleaning your data

Load and view

Are data types correctly encoded?

Complete dataset

- Stack to make tidy
- Merge with other datasets

Quality control

- Explore basic statistics: mean, stdev, etc.
- Are the values consistent and in range?
- Make some plots and look at the data

Save clean dataset



Tidy data

- Each variable is a column and contains values
- Each observation is a row
- Each type of observational unit forms a table

Messy:

	Treatment A	Treatment B
John Smith	-	2
Jane Doe	16	11
Mary Johnson	3	1

Tidy:

Name	Treatment	Result
John Smith	a	-
Jane Doe	a	16
Mary Johnson	a	3
John Smith	b	2
Jane Doe	b	11
Mary Johnson	b	1



Tools & dataset for this workshop

- We will work in Python...
- Which is embedded in a Jupyter notebook...
- Which runs on Surf facility.
- Dataset: we will work with a small fictitious patient dataset, with some general measurements (heart rate, blood pressure) and some factors per patient (i.e. hospital code, date).



Practical Data Handling in Python: your turn!

Part 1:

- Go to the SURF environment at https://uu-its.jove.surfsara.nl
- Log in with the details on your handout
- Open Jupyter notebook

Part 2:

- Install Anaconda if necessary
- Open Anaconda Launcher
- The files (notebook, data1, data2, .py script) can be found in the SURF environment.
- Store these in a folder on your computer
- Open file 'datahandling.py' in Spyder on your computer



Best practices for Python coding



In this (short) overview:

- Script setup
- Script layout and formatting
- Commenting
- File naming conventions



How to set up your python script

- 1. Header with information on the script:
 - Date
 - Context
 - Author
 - Purpose
 - Dependencies (what is needed to use the script)
- 2. Load modules (existing code you will use in your script)
- 3. Functions (reusable code that you have written yourself)
- 4. ...the rest of the script.

```
111
 1
                                                                                                        Header
 2
    This script can be used to detect distinct C2H2 zinc finger motifs
 3
    in files with protein sequences.
    The output is:
 4
    (1) a fasta file with domains in order;
 5
    (2) a csv file with all info that can be used for visualization.
 6
    (3) a stats file with numbers of motifs and the amount of double motifs, and a heatmap of these stats
    (4) fasta files for all motifs with sequences that were found
 8
    Author: Barbara Vreede
 9
    Contact: b.vreede@gmail.com
10
    Date: 10 October 2014
11
12
13
                                                                                                      Modules
14
    import re, sys,config,os.path
15
    import seaborn as sns
16
17
                                                                                                   Functions
18
    def makeheatmap(matrix, labelsx, labelsy, name):
19
            Makes a heatmap of a matrix, using Seaborn.
20
21
            sns.heatmap(matrix, xticklabels=labelsx,yticklabels=labelsy,linewidths=.5)#,cmap="YlOrBr")
22
            pl.savefig("%s-%s.svg" %(heatmapfig,name))
23
            pl.clf()
24
25
            pl.close()
26
27
                                                                                                           Script
28
    fastadict = config.fastadicter(fastadb) # translate the fasta file into a dictionary
    hmmdict = hmmdicter(hmmdb) # translate the hmmer output into a dictionary
29
    segdict = {} # dictionary for [start position]: sequence
30
    motdict = {} # dictionary for [start position]: motif type
31
32
33
    # go through the sequences
    for key in fastadict:
34
            # get info for the first columns (ID and sequence length)
35
            ids = key.split('|')
36
            seqlen = len(fastadict[key]) #length of the sequence
37
```



Python formatting conventions and best practices

- Humans should be able to read your code
- Indentation is a functional part of your code
- Variable names: avoid capitalization, punctuation, ...
- Write robust code: specify constants and variables
- Lines should be short (less than 80 characters)
- Use blank lines sparingly, but wisely

Style guide: https://www.python.org/dev/peps/pep-0008



Humans should be able to read your code

Bad

```
from pandas import to_datetime, read_csv
x =
read_csv("/home/student3/Desktop/backupsSaturdayDesktopHomeC
omputer/courses/2010/August/summercoursePython/data/PatientD
ATA2.txt")
to_datetime(x[5], format="%d-%m-%Y")
x[5].diff()
```



Indentation is a functional part of your code

Bad

```
for date in df_pd2["VISIT"]:
    print(date)

File "<stdin>", line 2
    print(date)
    ^
IndentationError: expected an indented block
```

Good

```
for date in df_pd2["VISIT"]:
    print(date)

2-3-2016
9-8-2015
17-4-2016
```

! Do not mix tabs and spaces! (1 indent = 4 spaces)



Variable names: avoid capitalization, punctuation...

Bad

```
# forbidden: never start with numbers or use .&*%()$#@-...
2df = pandas.read_csv("PatientDATA2.txt")
df.2 = pandas.read_csv("PatientDATA2.txt")

# bad manners: capitalization or obscure names
Df2 = pandas.read_csv("PatientDATA2.txt")
a = pandas.read_csv("PatientDATA2.txt")
```

```
df_pd2 = pandas.read_csv("PatientDATA2.txt")
patientdata2 = pandas.read_csv("PatientDATA2.txt")

# even more explicit
patientdata_visits = pandas.read_csv("PatientDATA2.txt")
```



Write robust code

Bad

```
# print the time between each subsequent visit
print(df_pd2[5].diff())

# we want to divide the patients into 5 groups
# define group size
print(100 / 5)
```

```
# print the time between each subsequent visit
print(df_pd2["VISITS"].diff())

# we want to divide the patients into 5 groups
# define group size
npatients = df_pd2.shape[1]
ngroups = 5
groupsize = npatients / ngroups
```



Lines should be short

Bad

```
# this is a very long comment, and you will have to scroll endlessly to re
print(pandas.to_datetime(df_pd2["VISIT"], dayfirst=False, format="%d-%m-%Y
```

```
# keep your comments short, too!
df_pd2["VISIT"] = pandas.to_datetime(
    df_pd2["VISIT"],
    format="%d-%m-%Y"
)
print(df_pd2["VISIT"].diff())
```



Use blank lines sparingly, but wisely

Bad

```
import pandas as pd
df_pd2 = pd.read_csv("PatientDATA2.txt")

df_pd2["VISIT"] = pd.to_datetime(
    df_pd2["VISIT"],
    format="%d-%m-%Y"
)
print(df_pd2["VISIT"])
```

```
import pandas as pd

df_pd2 = pd.read_csv("PatientDATA2.txt")
df_pd2["VISIT"] = pd.to_datetime(
         df_pd2["VISIT"],
         format="%d-%m-%Y"
)

print(df_pd2["VISIT"])
```



Python commenting best practices

- Headers and comments and docstrings, oh my...
- Humans should be able to read your code!
- Comment your intent, not the code's function
- Comments can be a great coding tool!



Headers and comments and docstrings, oh my...

Comment

```
# import the data
df_pd2 = pd.read_csv("PatientDATA2.txt")
```

```
df_pd2 = pd.read_csv("PatientDATA2.txt") # import the data
```

Docstring

```
This block of text is a docstring. It is a long(er) piece of text, that may run over several lines.

Triple quotation marks are used to indicate the beginning and the end of a docstring.

"""
```



Humans should be able to read your code

Bad

```
import pandas as pd

df_pd2 = pd.read_csv("PatientDATA2.txt")

df_pd2["VISIT"] = pd.to_datetime(
          df_pd2["VISIT"],
          format="%d-%m-%Y"
)
print(df_pd2["VISIT"].diff())
```

```
import pandas as pd

# import the data
df_pd2 = pd.read_csv("PatientDATA2.txt")

# print the time between each subsequent visit
df_pd2["VISIT"] = pd.to_datetime(
    df_pd2["VISIT"],
    format="%d-%m-%Y"
)
print(df_pd2["VISIT"].diff())
```



Comment your intent, not the code's function

Bad

```
df_pd2 = pd.read_csv("PatientDATA2.txt") # read the csv
df_pd2["VISIT"] = pd.to_datetime(df_pd2["VISIT"], format="%d-
%m-%Y") # the "VISIT" column is a date
print(df_pd2["VISIT"].diff()) # print the difference
```

```
# import the patient dataset that includes visit dates
df_pd2 = pd.read_csv("PatientDATA2.txt")

# print the time between each subsequent visit
df_pd2["VISIT"] = pd.to_datetime(
         df_pd2["VISIT"],
         format="%d-%m-%Y"
)
print(df_pd2["VISIT"].diff())
```



Comments can be a great coding tool

Starting your script...

```
# import the patient data
# make sure the dataset includes visits
# ensure proper formatting of the visit column
# print the time between each subsequent visit
```

... write the code, revise your comments



File naming conventions

Use fixed elements in your file name:

I.e. description of content, project number, name researcher/team.

DON'T →

- Use special characters (&%\$#) or periods or spaces
- Start with numbers (e.g. 2018 patientanalysis.py)

$DO \rightarrow$

- Keep names short, relevant, but descriptive (stand-alone!)
- Go from generic to specific (handy with sorting and finding)
- Use '_' when combining elements to a filename



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