### **Python Summer Course**

Course 5: Pandas

Théophile Gentilhomme July 29, 2025





## Introduction

Pandas is a Python library for **data analysis and manipulation**.

It provides two main data structures:

- Series → 1D labeled array
- DataFrame → 2D labeled table (like Excel or a database)





Pandas





## Why Use Pandas?

- Built on top of NumPy
- Makes it easy to:
  - Load and clean data
  - Filter, transform, and group data
  - Perform statistics and summaries
- Works well with CSV, Excel, SQL, and more





## **Example**

- Each key becomes a column name
- Each value must be a list or array of the same length

Pandas is the perfect tool for **data science**, **bioinformatics**, and any task involving **structured data**.





## DataFrame = 2D Table

A DataFrame is like a table with rows and columns.

```
Python Code Start Over

import pandas as pd

data = {
  "Name": ["Alice", "Bob", "Charlie"],
  "Age": [25, 30, 35]

df = pd.DataFrame(data)
  print(df)
```

- Rows are automatically indexed (0, 1, 2...)
- Columns are labeled (like a dict of Series)





## Series = 1D Column or Row

A Series is like a **single column** with an index:

- Has both values and an index
- Acts like a NumPy array + labels





# Access by Label .loc[row\_label,

## column\_label]

You can access specific element of the table using row/column label with .loc[].

```
Python Code ⊕ Start Over

1  # Access one element
2  print(df.loc[0, "Age"])  # Age of Alice
```





# **Modify an element**

```
Python Code ⊕ Start Over

1 df.loc[2, "Age"] = 31
2 print(df)
```





## Row/column access

```
Python Code ⊕ Start Over

1  print (df.loc[1])  # Entire row (Series)

2  print (df.loc[:, "Name"])  # Entire column (Series)
```





# **Access by Index**

### .iloc[row\_index, column\_index]

```
Python Code ⊕ Start Over

1 print (df.iloc[0, 1]) # Same as above (row 0, column 1)
```

#### Same as .loc[] but with indexes





# Add a Column / Filter Rows

You can add columns just like a dictionnary

```
Python Code ⊕ Start Over

1 df["HES"] = [True, False, False]
```





# Filter with conditions (Boolean mask)

You can filter the row using boolean mask with the same lenght as the columns.

Usually, you filter using value in one or several columns

```
Python Code Start Over

1 print (df.loc[df["Age"] > 30]) # Rows where Age > 30
2
3 # Combine conditions with `&` (and) `|` (or)
```





# Loading CSV and Excel Files

Pandas makes it easy to import real-world data from common formats.

This is usually what your are going to do.

Examples for CSV and Excel files





## **Loading files**

```
1 import pandas as pd
2 # Load a CSV file from a URL or local path
3 df = pd.read_csv("my_data.csv") # or from a URL
4 print(df.head()) # Show first 5 rows
```

- Use sep=";" if using semicolon separator
- Use encoding="utf-8" if needed
- Can use a column as row indexes index\_col="thisCol"

```
1 # Load an Excel File
2 df = pd.read_excel("my_data.xlsx", sheet_name="Sheet1")
```





# **Compute Stats and Explore Data**

Pandas offers many built-in tools to summarize and analyze your data quickly.





# **Quick Exploration**





#### **Basic Statistics**





#### **Data Overview**

```
Python Code ⊕ Start Over

1 print (df.info()) # Structure and types
2 print (df.describe()) # Summary stats for numeric columns
```





# Handling Missing Values in Pandas

Missing values are common in real-world data. Pandas uses NaN (Not a Number) to represent them.





## **Drop or Fill**





# Simple Plots with Pandas

Pandas makes it easy to plot data directly from a DataFrame.

It uses <u>Matplotlib</u>, which is a Python library commuly used for plots.

You can customize the plots using Matplolib functionalities.





#### **Line Plot**

```
▶ Run Code
 1 import pandas as pd
   import matplotlib.pyplot as plt
 3
 4 df = pd.DataFrame({
 5
       "Year": [2020, 2021, 2022, 2023],
       "Revenue": [100, 150, 200, 250]
 6
   })
 8
   df.plot(x="Year", y="Revenue", kind="line", title="Revenue Over Time")
10
11
   # Call pyplot show
12 plt.show() # You normally do need this (only for the presentation)
```





## **Other Plot Types**

```
→ Start Over
Python Code
                                                                                  ▶ Run Code
1 df.plot(kind="bar") # Bar chart
2 plt.show()
Python Code
            → Start Over
                                                                                  ▶ Run Code
1 df.plot(kind="hist")
                                # Histogram
2 plt.show()

→ Start Over

Python Code
                                                                                  ▶ Run Code
1 df.plot(kind="box")
                                # Boxplot
2 plt.show()
            → Start Over
                                                                                  ▶ Run Code
Python Code
 df.plot(kind="scatter", x="Year", y="Revenue") # Scatter plot
2 plt.show()
```





# And many more to cover on Pandas!

- group\_by()
- Time series
- interpolation
- ...





### Your turn!

You are given a dataset gene\_expression\_data.cvs. There is no descriptions, but you need to explore and make some analyses.

- 1. Load the file using Pandas
- 2. Explore it using the functions seen in this course
- 3. Check how many treated vs healthy samples there are using value\_counts() on the column or using boolean condition





- 4. Compute the mean expression for <a href="mailto:Tissue\_B">Tissue\_B</a> (optional: all tissues)
- 5. Calculate the difference between Tissue A and B and store it in a new column
- 6. Filter the genes where Tissue C expression is greater than 9 and Biomarker is True (i.e. two boolean expression combined with &)





7. Plot a scatter plot of Tissue A vs Tissue B, colored with biomarker. Tip: use scatter with

```
colors = ["red" if b else "blue" for b in df["Biomarker"]]
```





# Solution





### 1. Load the CSV





# 2. Explore the Data





### 3. Count Conditions





# 4. Mean Expression per Tissue





# 5. Add Difference Column (Tissue A - B)





# 6. Filter: Tissue C > 9 and Biomarker = True





# 7. Scatter Plot (Tissue A vs B, colored by Biomarker)





## More references

Python course for data analysis

PANDAS-TUTORIAL

<u>Pandas</u>



