## What is Pandas?

**Pandas** is one of the most popular data science libraries in Python. Easy to use, it is built on top of **NumPy** and shares many functions and properties.

With Pandas, you can read and extract data from files, transform and analyze it, calculate statistics and correlations, and much more!

To start using pandas, we need to import it first:



pd is a common short name used when importing the library.

Pandas is derived from the term "panel data", an econometrics term for data sets that include observations over multiple time periods for the same individuals.

## Series & DataFrames

The two primary components of pandas are the Series and the DataFrame.

A **Series** is essentially a column, and a **DataFrame** is a multi-dimensional table made up of a collection of Series.

For example, the following DataFrame is made of two Series, ages and heights

ages	heights		
14	165		
18	180		
24	176		
42	184		

You can think of a **Series** as a one-dimensional array, while a **DataFrame** is a multi-dimensional array.

#### **DataFrames**

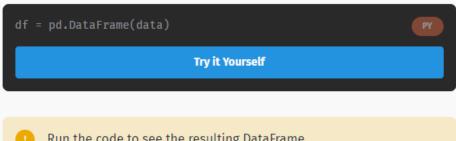
Before working with real data, let's first create a DataFrame manually to explore its functions.

The easiest way to create a DataFrame is using a dictionary:

```
data = {
   'ages': [14, 18, 24, 42],
   'heights': [165, 180, 176, 184]
```

Each key is a column, while the value is an array representing the data for that column.

Now, we can pass this dictionary to the DataFrame constructor:



Run the code to see the resulting DataFrame.

## **DataFrames**

The DataFrame automatically creates a numeric **index** for each row. We can specify a custom index, when creating the DataFrame:



Now we can access a row using its index and the **loc[]** function:



This will output the row that corresponds to the index "Bob".

1 Note, that **loc** uses square brackets to specify the index.

# **Indexing**

We can select a single column by specifying its name in square brackets:

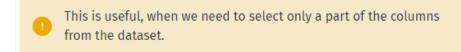


The result is a Series object.

If we want to select multiple columns, we can specify a list of column names:



This time, the result is a DataFrame, as it includes multiple columns.



# **Slicing**

Pandas uses the **iloc** function to select data based on its numeric index. It works the same way indexing lists does in Python.

#### For example:

```
# third row
print(df.iloc[2])

#first 3 rows
print(df.iloc[:3])

# rows 2 to 3
print(df.iloc[1:3])

Try it Yourself
```

1 iloc follows the same rules as slicing does with Python lists.

## **Conditions**

We can also select the data based on a condition.

For example, let's select all rows where **age** is greater than 18 and **height** is greater than 180:



# **Reading Data**

It is quite common for data to come in a file format. One of the most popular formats is the **CSV** (comma-separated values).

Pandas supports reading data from a CSV file directly into a DataFrame.

For our examples, we will use a CSV file that contains the COVID-19 infection data in California for the year 2020, called 'ca-covid.csv'.

The **read\_csv()** function reads the data of a CSV file into a DataFrame:



# **Reading Data**

Once we have the data in a DataFrame, we can start exploring it. We can get the first rows of the data using the head() function of the DataFrame:



	date	state	cases	deaths
0	25.01.20	California	1	0
1	26.01.20	California	1	0
2	27.01.20	California	0	0
3	28.01.20	California	0	0
4	29.01.20	California	0	0

By default it returns the first 5 rows. You can instruct it to return the number of rows you would like as an argument (for example, **df.head(10)** will return the first 10 rows).

We can see that our DataFrame contains the **date**, **state**, number of **cases** and **deaths** for that date.



Similarly, you can get the last rows using the tail() function.

## **Reading Data**

The **info()** function is used to get essential information about your dataset, such as number of rows, columns, data types, etc:



#### Run the code to see the result!

From the result, we can see that our dataset contains 342 rows and 4 columns: date, state, cases, deaths.

We also see that Pandas has added an auto generated index.
We can set our own index column by using the **set\_index**() function:

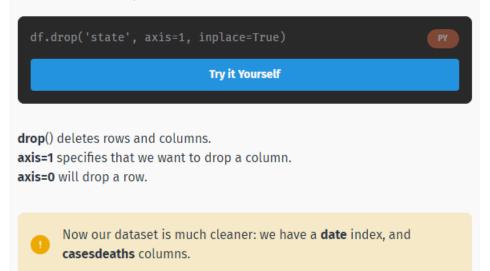


The **date** column is a good choice for our index, as there is one row for each date.

The **inplace=True** argument specifies that the change will be applied to our DataFrame, without the need to assign it to a new DataFrame variable.

# **Dropping a Column**

Since our data is only for the state of California, we can remove that column from our DataFrame, as it contains the same value for all rows:



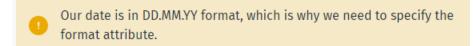
# **Creating Columns**

Pandas allows us to create our own columns.

For example, we can add a **month** column based on the **date** column:



We do this by converting the **date** column to datetime and extracting the month name from it, assigning the value to our new **month** column.



# **Summary Statistics**

Now that our dataset is clean and set up, we are ready to look into some stats!

The **describe**() function returns the summary statistics for all the numeric columns:



This function will show main statistics for the numeric columns, such as **std**, **mean**, **min**, **max** values, etc.

#### Run the code to see the result!

From the result, we see that the maximum cases that have been recorded in a day is **64987**, while the average daily number of new cases is **6748**.



We can also get the summary stats for a single column, for example: df['cases'].describe()

#### Grouping

Since we have a **month** column, we can see how many values each month has, by using the **value\_counts**() functions:





We can see that, for example,  ${\bf January}$  has only 7 records, while the other months have data for all days.



value\_counts() returns how many times a value appears in the dataset, also called the frequency of the values.

# Grouping

Now we can calculate data insights!

For example, let's determine the number of total infections in each month. To do this, we need to group our data by the month column and then calculate the sum of the cases column for each month:



The **groupby**() function is used to group our dataset by the given column.

We can also calculate the number of total cases in the entire year:



We can see that California had 2,307,769 infection cases in 2020.

