

# Python Dataframe

pandas library

# Read from a CSV file

- `import pandas as pd`
- 
- `df = pd.read_csv('data/iris.csv')`
- `print(type(df))`
- `---`
- `<class 'pandas.core.frame.DataFrame'>`
-

# dataframe

- Print first 5 rows
  - `print(df.head())`
- Print name of the columns
  - `print(df.columns)`
- Print Number rows and columns
  - `print(df.shape)`

# dataframe

- `print(df.index)` - rows index
- `print(df.iloc[1])` - index a row
- `print(df.iloc[1,1])` – index a entry
- `print(df.dtypes)` – types of the columns
  - columns all have the same type

# Input and Output dataframes

- `X =  
df[["sepal_length", "sepal_width", "petal_length", "  
petal_width"]]`
- `y = df["species"]`
- `print(type(X))`
- `print(type(y))`
- `print(X.shape)`
- `print(y.shape)`

# Classification

- The iris dataset is typically used for classification.
- Input the values of the columns, "sepal\_length", "sepal\_width", "petal\_length", "petal\_width"
- Output – classification (setosa, versicolor, virginica)
- X used as input (multiple columns)
- y output (single column)