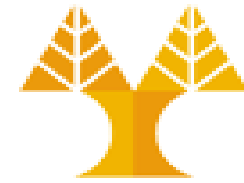


# EPL448: Data Mining on the Web – Lab 4



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# Python Libraries for Data Science

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Many popular Python toolboxes/libraries:

## Data Manipulation (Lab 4)

- NumPy
- SciPy
- **Pandas**

## Data Visualization (Lab 5)

- **matplotlib**
- **seaborn**

## Data Pre-processing (Labs 6-7)

## Machine Learning (Labs 8-9)

- **SciKit-Learn**

# Introduction to Pandas

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- Pre-installed in Anaconda
    - No need to install it separately
    - import it to a notebook using: `import pandas as pd`
      - `pd` is the de facto abbreviation for Pandas used by the data science community
  - Primary data structures in Pandas:
    - Series
    - DataFrames
-

# Pandas: Series & DataFrames Examples



- A **Series** is a one-dimensional array with axis labels

	0	50
axis 0	1	90
	2	100
	3	45

dtype: int64

index

Axis labels are stored in the index. Series support both integer-based and string-based indexing. The default, if index is not set, is integer-based starting from 0.

<https://pandas.pydata.org/pandas-docs/stable/reference/api/pandas.Series.html>

- A **DataFrame** is a two-dimensional tabular dataset with labeled axes

		names	grades
axis 0	0	bob	50
	1	ken	90
	2	art	100
	3	joe	45

Index (row labels)

Column labels

<https://pandas.pydata.org/pandas-docs/stable/reference/api/pandas.DataFrame.html>

# Read data using Pandas



```
import pandas as pd # load pandas library and create alias pd

# Load dataset iris_data.csv and create dataframe object
df = pd.read_csv('iris_data.csv') # you can read a file from url as well:
df = pd.read_csv('http://www.cs.ucy.ac.cy/courses/EPL448/labs/LAB04/iris_data.csv')
```

- There is a number of pandas commands to read other data formats:

```
pd.read_excel('myfile.xlsx', sheet_name='Sheet1', index_col=None, na_values=['NA'])
pd.read_xml('myfile.xml')
pd.read_sql_query('select * from iris', conn)
```

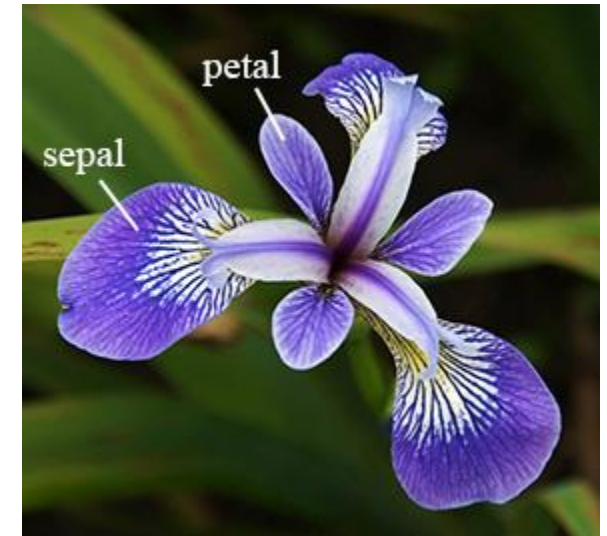
- A file may not always have a header row
  - pandas to assign default column names (but not informative)
  - you can specify (column) names yourself

```
df = pd.read_csv('https://www.cs.ucy.ac.cy/courses/EPL448/labs/LAB04/iris_data2.csv',
                 names=['sepal-length', 'sepal-width', 'petal-length', 'petal-width', 'class'])
```

# Dataset: The iris dataset



- iris flowers dataset
  - “hello world” dataset in machine learning and statistics
- Small dataset with 150 observations of iris flowers
  - each observation has 4 columns of measurements (or variables or features) of the flowers (in centimeters)
  - 5th column is the species (class) of the flower observed
  - all observed flowers belong to one of three species



```
sepal-length, sepal-width, petal-length, petal-width, class  
5.1, 3.5, 1.4, 0.2, Iris-setosa  
5.9, 3.0, 4.2, 1.5, Iris-versicolor  
5.8, 2.7, 5.1, 1.9, Iris-virginica  
4.6, 3.1, 1.5, 0.2, Iris-setosa
```

- More info: [https://en.wikipedia.org/wiki/Iris\\_flower\\_data\\_set](https://en.wikipedia.org/wiki/Iris_flower_data_set)

# Dataset Overview

---



- Features:
  - sepal length in cm
  - sepal width in cm
  - petal length in cm
  - petal width in cm
  
- Classes (labels):
  - Iris Setosa
  - Iris Versicolour
  - Iris Virginica

# Read data using Pandas: Example



```
In [ ]: # Read csv file
df = pd.read_csv('iris_data.csv')
print(df)
```

```
Out[ ]:      sepal-length  sepal-width  petal-length  petal-width      class
0           5.1         3.5         1.4         0.2  Iris-setosa
1           4.9         3.0         1.4         0.2  Iris-setosa
2           4.7         3.2         1.3         0.2  Iris-setosa
3           4.6         3.1         1.5         0.2  Iris-setosa
4           5.0         3.6         1.4         0.2  Iris-setosa
..          ...         ...         ...         ...         ...
145         6.7         3.0         5.2         2.3  Iris-virginica
146         6.3         2.5         5.0         1.9  Iris-virginica
147         6.5         3.0         5.2         2.0  Iris-virginica
148         6.2         3.4         5.4         2.3  Iris-virginica
149         5.9         3.0         5.1         1.8  Iris-virginica
```

By default, only 10 rows (first & last 5 rows) of the DataFrame are printed. If the number of columns is also large, only 4 columns (first & last 2) are printed.



# Read data using Pandas: Example



```
In [ ]: # Read csv file
df = pd.read_csv('iris_data.csv')
df # jupyter can print the result of the last evaluation of a cell without
a print() function in a more visually-appealing way
```

```
Out[ ]:
```

	sepal-length	sepal-width	petal-length	petal-width	class
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa
...	...	...	...	...	...
145	6.7	3.0	5.2	2.3	Iris-virginica
146	6.3	2.5	5.0	1.9	Iris-virginica
147	6.5	3.0	5.2	2.0	Iris-virginica
148	6.2	3.4	5.4	2.3	Iris-virginica
149	5.9	3.0	5.1	1.8	Iris-virginica

150 rows × 5 columns

By default, only 10 rows (first & last 5 rows) of the DataFrame are printed. If the number of columns is also large, only 4 columns (first & last 2) are printed.

# DataFrames attributes



Python objects have *attributes* and *methods*.

<b>df.attribute</b>	<b>description</b>
dtypes	list the types of the columns
columns	list the column names
axes	list the row labels and column names
ndim	number of dimensions
size	number of elements (number of all values)
shape	return a tuple representing the dimensionality (rows, columns)
values	numpy representation of the data without index and column names

# Data Frames methods



Unlike attributes, methods have *parentheses*. See some methods below.

All attributes and methods can be listed with a *dir()* function: `dir(df)`

<b>df.method()</b>	<b>description</b>
head( [n] ), tail( [n] )	first/last n rows
describe()	generate summary statistics (for numeric columns only)
max(), min()	return max/min values. Select all numeric columns using <code>numeric_only=True</code>
sum()	return the sum of values. Select all numeric columns using <code>numeric_only=True</code>
mean(), median(), std()	return mean/median/standard deviation. Select all numeric columns using <code>numeric_only=True</code>
dropna()	drop all rows with at least one missing value (set axis param to 1 to drop columns)
drop()	drop specified rows or columns
count()	count non-NA cells
value_counts()	returns counts of unique values in descending order so that the first element is the most frequently-occurring element
apply()	applies a function along an axis of the DataFrame
replace()	replaces values with other values
cut()	bin (group) values into discrete intervals
astype(type)	casts to a specified data type (e.g. <code>astype(int)</code> )

# Explore DataFrames



- Dimensions of Dataset

```
# shape
```

```
print(df.shape)
```

(150, 5)

- Peek at the Data

```
# head (10 first lines)
```

```
print(df.head(10))
```

	sepal-length	sepal-width	petal-length	petal-width	class
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa
5	5.4	3.9	1.7	0.4	Iris-setosa
6	4.6	3.4	1.4	0.3	Iris-setosa
7	5.0	3.4	1.5	0.2	Iris-setosa
8	4.4	2.9	1.4	0.2	Iris-setosa
9	4.9	3.1	1.5	0.1	Iris-setosa

- Data type for each column

```
print(df.dtypes)
```

sepal-length	float64
sepal-width	float64
petal-length	float64
petal-width	float64
class	object

- Statistical summary

```
# descriptions
```

```
print(df.describe())
```

	sepal-length	sepal-width	petal-length	petal-width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.054000	3.758667	1.198667
std	0.828066	0.433594	1.764420	0.763161
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.300000
50%	5.800000	3.000000	4.350000	1.300000
75%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	2.500000

# Explore DataFrames



```
#List DataFrame columns  
print(df.columns)
```

```
Index(['sepal-length', 'sepal-width', 'petal-length', 'petal-width', 'class'],  
      dtype='object')
```

```
#Get DataFrame values  
print(df.values)
```

```
[[5.1 3.5 1.4 0.2 'Iris-setosa']  
 [4.9 3.0 1.4 0.2 'Iris-setosa']  
 [4.7 3.2 1.3 0.2 'Iris-setosa']  
 ...  
 [6.5 3.0 5.2 2.0 'Iris-virginica']  
 [6.2 3.4 5.4 2.3 'Iris-virginica']  
 [5.9 3.0 5.1 1.8 'Iris-virginica']]
```

# Select a column in a Data Frame

---



*Method 1:* Use the column name in square brackets:

```
df['sepal-width']
```

Can be used to select more than one column:

```
df[['sepal-width', 'sepal-length']]
```

*Method 2:* Use the column name as an attribute:

```
df.column_name
```

*Note:* column names can be used as attributes when they are valid variable names (starting with a letter followed by a letter, a number or an underscore \_ and do not conflict with a predefined attribute e.g. max, min, etc. For example, we cannot use df.sepal-length since - is considered as the minus symbol.

# Drop rows / columns



- `drop()` method can be used to remove index (rows) or columns by specifying label names and corresponding axis, or by specifying directly index or column names
  - Drops rows if `axis = 0` or `'index'` (default), drops columns of `axis = 1` or `'columns'`
  - Returns a new DataFrame without the removed rows or columns or `None` if `inplace=True`

```
In [ ]: # drop 2 columns
df.drop(columns=['sepal-length', 'sepal-width'])
# alternative command with the same effect
df.drop(['sepal-length', 'sepal-width'], axis=1)
# drop index (rows) with index = 3 and 5
df.drop(index=[3, 5])
# alternative command with the same effect
df.drop([3, 5])
# alternative command with the same effect
df.drop([3, 5], axis=0)
```

# DataFrame processing operations

---



- Group-by method
  - splits data into groups by a categorical col, e.g. group faculty members by rank or gender
- Aggregate (or agg) method
  - an aggregation method is one which takes multiple individual values and returns a summary; in most of the cases, this summary is a single value
  - apply multiple aggregation methods on one or more columns or groups
- Splitting (binning) operation
  - bins (groups) values into discrete intervals; convert numerical to categorical, e.g. split faculty member into categorical groups based on their age: 25-39 young, 40-59 middle, 60+ old
- Filtering operation
  - selects a subset of data based on one or more conditions
- Slicing operation
  - selects a subset of data by row or column position/label
- Sorting operation
  - sorts data by the values of one or more column(s)



# Grouping data using groupby()



- Using "group by" method we can:
  - Split the data into groups based on some criteria (e.g. the values of a column)
  - Once a groupby object is created, we can run aggregation methods (e.g. sum, mean) on each group and combine the results into a data structure

df

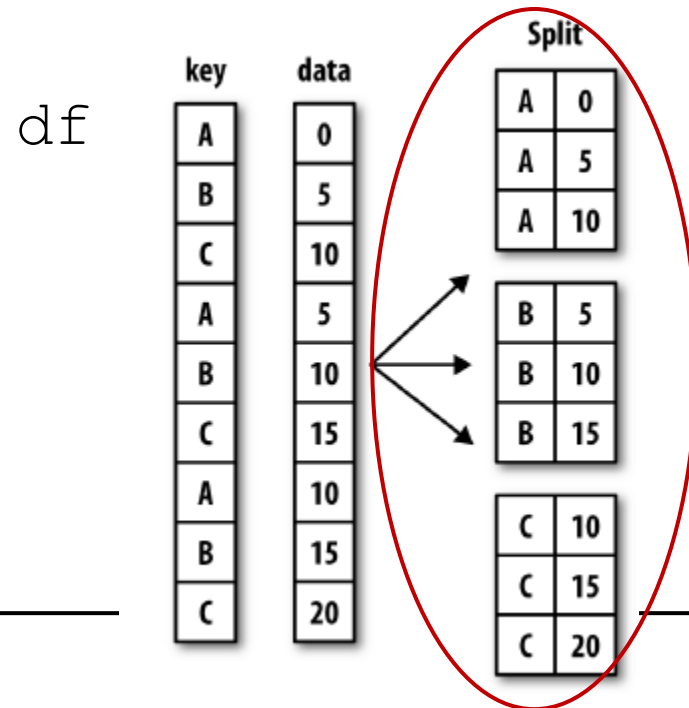
key	data
A	0
B	5
C	10
A	5
B	10
C	15
A	10
B	15
C	20

# Grouping data using groupby()



- Using "group by" method we can:
  - Split the data into groups based on some criteria (e.g. the values of a column)
  - Once a groupby object is created, we can run aggregation methods (e.g. sum, mean) on each group and combine the results into a data structure

```
df.groupby('key')
```

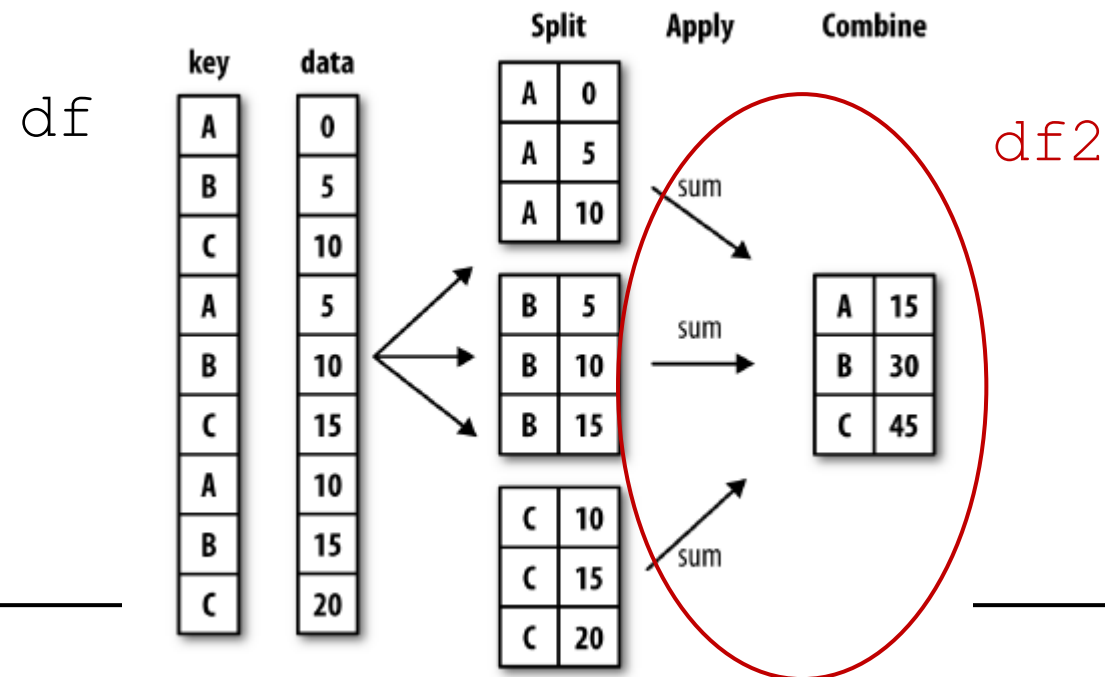


# Grouping data using groupby()



- Using "group by" method we can:
  - Split the data into groups based on some criteria (e.g. the values of a column)
  - Once a groupby object is created, we can run aggregation methods (e.g. sum, mean) on each group and combine the results into a data structure

```
df2 = df.groupby('key').sum()
```



# Grouping data using groupby(): Example



- Using "group by" method we can:
  - Split the data into groups based on some criteria (e.g. the values of a column)
  - Once a groupby object is created, we can run aggregation functions (e.g. Sum, mean) on each group and combine results into a data structure

```
In [ ]: #Group data using class column which has categorical data
df_groupby_class = df.groupby('class')
```

```
In [ ]: #Calculate mean value for each numeric column per each group
df_groupby_class.mean(numeric_only=True)
```

	class	sepal-length	sepal-width	petal-length	petal-width
Groups	Iris-setosa	5.006	3.418	1.464	0.244
	Iris-versicolor	5.936	2.770	4.260	1.326
	Iris-virginica	6.588	2.974	5.552	2.026

When applying groupby, a new **groupby object** is returned having the **specified group** as **index name**

# Grouping data using groupby()



- On the groupby object we can isolate column(s) and then run aggregation functions (statistics) for each group:

```
In [ ]: # Isolate sepal-length and calculate mean for each flower class:  
df_groupby_class[['sepal-length']].mean(numeric_only=True)
```

```
Out[ ]:  
      class  sepal-length  
Group  
names Iris-setosa  91786.230769  
      Iris-versicolor  81362.789474  
      Iris-virginica  123624.804348
```

*Note:* If **single brackets** are used to specify the column (e.g. sepal-length), then the output is **Pandas Series** object. When **double brackets** are used the output is a **DataFrame**.

# Grouping data using groupby()



- `groupby()` can be performed on multiple columns

```
In [ ]: #Group data using class and other column(s) which have categorical data
df_class = df.groupby(['class', '...'])
df_class.mean(numeric_only=True)
```

- `groupby` performance notes:
  - by default, the group names are sorted during the `groupby` operation. You may want to set `sort=False` for potential processing speedup:

```
In [ ]: #Calculate mean sepal-length for each flower class:
df.groupby('class', sort=False)[['sepal-length']].mean()
```

# Aggregations on multiple columns or groups



- `agg()` method allows applying **multiple** aggregation methods **on one or more columns or groups**

```
In [ ]: # get mean values for all columns
df_groupby_class.agg(['mean'])
```

```
Out [ ]:      sepal-length      sepal-width      petal-length      petal-width
          mean          mean          mean          mean
class
Iris-setosa      5.006      3.418      1.464      0.244
Iris-versicolor  5.936      2.770      4.260      1.326
Iris-virginica   6.588      2.974      5.552      2.026
```

```
In [ ]: # multiple aggregations (max, mean, std) for all columns
df_groupby_class.agg(['max', 'mean', 'std'])
```

```
Out [ ]:      sepal-length      sepal-width      petal-length      petal-width
          max mean      std max mean      std max mean      std max mean      std
class
Iris-setosa  5.8  5.006  0.352490  4.4  3.418  0.381024  1.9  1.464  0.173511  0.6  0.244  0.107210
Iris-versicolor  7.0  5.936  0.516171  3.4  2.770  0.313798  5.1  4.260  0.469911  1.8  1.326  0.197753
Iris-virginica  7.9  6.588  0.635880  3.8  2.974  0.322497  6.9  5.552  0.551895  2.5  2.026  0.274650
```

# Split data values into bins (groups)

---



- `cut()` method bins (groups) values into discrete intervals
  - Useful for going from a numerical (continuous) variable to a discrete (categorical) variable
    - For example, `cut()` could convert ages to groups of age ranges e.g.  $(0-12]$  → child,  $(12-18]$  → teenager,  $(18-60]$  → adult,  $(60-\infty]$  → elder)
  - Supports binning **into equal-sized bins, or a pre-specified array of bins**
  - Returns an array-like object representing the respective bin for each value of the column to be split
-



# Split data values into bins (groups)



- Example 1: Split flowers by their sepal-length into 3 equal-sized bins (all bins have the same number of flowers)

```
In [ ]: pd.cut(df['sepal-length'], bins=3)
```

```
Out[ ]: 0      (4.296, 5.5] } belong to the first bin
        1      (4.296, 5.5]
        2      (4.296, 5.5]
        3      (4.296, 5.5]
        4      (4.296, 5.5]
        ...
        145     (5.5, 6.7] } belong to the third bin
        146     (5.5, 6.7]
        147     (5.5, 6.7]
        148     (5.5, 6.7]
        149     (5.5, 6.7]
Name: sepal-length, Length: 150, dtype: category
Categories (3, interval[float64, right]): [(4.296, 5.5] < (5.5, 6.7] < (6.7, 7.9]]
```

# Split data values into bins (groups)



- Example 1: **Split** flowers **by their sepal-length** into 3 equal-sized bins (a label for each split can be defined)

```
In [ ]: pd.cut(df['sepal-length'], bins=3, labels=['low', 'medium', 'high'])
```

```
Out[ ]: 0          low
        1          low
        2          low
        3          low
        4          low
        ...
        145       medium
        146       medium
        147       medium
        148       medium
        149       medium
        Name: sepal-length, Length: 150, dtype: category
        Categories (3, object): ['low' < 'medium' < 'high']
```

belong to the first bin

belong to the third bin

# Split data values into bins (groups)



- Example 2: Split flowers by their sepal-length into **3 pre-specified array of bins** (a label for each split can be defined)

```
In [ ]: pd.cut(df['sepal-length'], bins=[4, 5, 6, np.inf], labels=['small', 'medium', 'large'])
```

```
Out[ ]: 0      medium  belongs to the second bin
        1      small  }
        2      small  } belong to the first bin
        3      small  }
        4      small  }
        ...
        145     large  }
        146     large  } belong to the third bin
        147     large  }
        148     large  }
        149     medium belongs to the second bin
Name: sepal-length, Length: 150, dtype: category
Categories (3, object): ['small' < 'medium' < 'large']
```

# Data Frame: Filtering



- To subset the data we can apply Boolean indexing. This indexing is commonly known as a filter. For example, if we want to subset the rows in which the petal length value is greater than 5.5 cm:

```
#Select only those rows that petal-length is more than 5.5:  
df_sub = df[ df['petal-length'] > 5.5 ]
```

Any Boolean operator can be used to subset the data:

> greater;      >= greater or equal;  
< less;        <= less or equal;  
== equal;      != not equal;

```
0      False  
1      False  
2      False  
3      False  
4      False  
...  
145    False  
146    False  
147    False  
148    False  
149    False  
Name: petal-length,  
Length: 150, dtype:  
bool
```

```
#Select only those rows that contain Iris setosa flowers:  
df_i = df[ df['class'] == 'Iris-setosa' ]
```

# Data Frame: Filtering

---



- Symbol & refers to AND condition which means meeting both the criteria:

```
df_1 = df[(df['class'] == 'Iris-setosa') & (df['petal-length'] > 5.5) ]
```

- Symbol | refers to OR condition which means meeting any of the criteria:

```
df_2 = df[ (df['class'] == 'Iris-virginica') | (df['petal-length'] > 2) ]
```

---

# Data Frame: Filtering



- Select rows which a specific column involves specific values

```
df_3 = df[ df['petal-length'].isin([5,10,15]) ]
```

- Select rows which a specific column contains a specific letter

```
df_4 = df[ df['rank'].str.contains('f') ]
```

- Select rows with NaN values in specific column

```
df_5 = df[ df['class'].isnull() ]
```

- Select rows with NaN values in any column

```
df_6 = df[ df.isnull().any(axis=1) ]
```

# Data Frames: Slicing

---



- There are a number of ways to subset (get a slice of) the DataFrame:
    - select one or more columns
    - select one or more rows
    - select a subset of rows and columns
  
  - Rows and columns can be selected by their position or label
-

# Data Frames: Slicing (selecting columns)



- When selecting one column, it is possible to use single set of brackets, but the resulting object will be a Series (not a DataFrame):

```
In [ ]: #Select sepal-length column:  
df['sepal-length']
```

- When we need to select **more than one columns** and/or make the output to be a DataFrame, we should use double brackets:

```
In [ ]: #Select sepal-length and sepal-width columns:  
df[['sepal-length', 'sepal-width']]
```



# Data Frames: Slicing (selecting rows)



- If we need to select a subset of rows, we can specify the range using :

```
In [ ]: #Select rows by their position:  
df[10:20]
```

- The first row has a position 0, and the last value in the range is not returned:
  - So for 0:10 range the first 10 rows are returned with the positions starting with 0 and ending with 9
- BUT (\*):

```
In [ ]: #Does not show the first row. USE iloc instead (see next slide)  
df[0]
```

---

(\*) The primary purpose of the DataFrame indexing operator, [] is to select columns.

# DataFrames: method `iloc`



- If we need to select a single row using its integer position we can use the method `iloc`:

```
In [ ]: #Select a row by its position:  
df.iloc[0]
```

```
Out[ ]: sepal-length      5.1  
        sepal-width     3.5  
        petal-length    1.4  
        petal-width     0.2  
        class           Iris-setosa  
        Name: 0, dtype: object
```

← Result as Pandas Series

```
In [ ]: #Select a row by its position (returns Dataframe):  
df.iloc[[0]]
```

	sepal-length	sepal-width	petal-length	petal-width	class
0	5.1	3.5	1.4	0.2	Iris-setosa

← Result as Pandas DataFrame

# DataFrames: method `iloc`



- If we need to select a range of rows and/or columns, using their (integer) positions we can use method `iloc`:

```
In [ ]: #Select rows and columns by their positions:  
df.iloc[10:20, [0, 2, 3]]
```

```
Out[ ]:
```

	sepal-length	petal-length	petal-width
10	5.4	1.5	0.2
11	4.8	1.6	0.2
12	4.8	1.4	0.1
13	4.3	1.1	0.1
14	5.8	1.2	0.2
15	5.7	1.5	0.4
16	5.4	1.3	0.4
17	5.1	1.4	0.3
18	5.7	1.7	0.3
19	5.1	1.5	0.3

# DataFrames: method `iloc` (summary)



```
df.iloc[0]    # First row of a data frame  
df.iloc[i]    # (i+1)th row  
df.iloc[-1]   # Last row
```

```
df.iloc[:, 0] # All rows of the first column  
df.iloc[:, -1] # All rows of the last column
```

```
df.iloc[0:7]      # First 7 rows  
df.iloc[:, 0:2]   # All rows of the first 2 columns  
df.iloc[1:3, 0:2] # Second through third rows and first 2 columns  
df.iloc[[0,5], [1,3]] # 1st and 6th rows and 2nd and 4th columns
```

# DataFrames: method loc



- If we need to select a range of rows and/or columns, using their labels we can use method loc:

```
In [ ]: #Select rows and columns by their labels:  
df.loc[10:20, ['sepal-width', 'sepal-length']]
```

```
Out[ ]:
```

	sepal-width	sepal-length
10	3.7	5.4
11	3.4	4.8
12	3.0	4.8
13	3.0	4.3
14	4.0	5.8
15	4.4	5.7
16	3.9	5.4
17	3.5	5.1
18	3.8	5.7
19	3.8	5.1
20	3.4	5.4

# Data Frames: Sorting



- We can sort the data by the values of a specified column. By default, the sorting will occur in ascending order (change using ascending Boolean parameter) and a new dataframe is returned.

```
In [ ]: # Create a new data frame from the original,  
# sorted by the column sepal-length  
df_sorted = df.sort_values(by = 'sepal-length')  
df_sorted.head()
```

```
Out[ ]:      sepal-length  sepal-width  petal-length  petal-width      class  
13           4.3           3.0           1.1           0.1  Iris-setosa  
42           4.4           3.2           1.3           0.2  Iris-setosa  
38           4.4           3.0           1.3           0.2  Iris-setosa  
8            4.4           2.9           1.4           0.2  Iris-setosa  
41           4.5           2.3           1.3           0.3  Iris-setosa
```

# Data Frames: Sorting



- We can sort the data using 2 or more columns:

```
In [ ]: df_sorted = df.sort_values( by =['sepal-length', 'sepal-width'],  
ascending = [True, False])  
df_sorted.head(10)
```

```
Out[ ]:      sepal-length  sepal-width  petal-length  petal-width  class  
13          4.3          3.0          1.1          0.1  Iris-setosa  
42          4.4          3.2          1.3          0.2  Iris-setosa  
38          4.4          3.0          1.3          0.2  Iris-setosa  
8           4.4          2.9          1.4          0.2  Iris-setosa  
41          4.5          2.3          1.3          0.3  Iris-setosa  
22          4.6          3.6          1.0          0.2  Iris-setosa  
6           4.6          3.4          1.4          0.3  Iris-setosa  
47          4.6          3.2          1.4          0.2  Iris-setosa  
3           4.6          3.1          1.5          0.2  Iris-setosa  
2           4.7          3.2          1.3          0.2  Iris-setosa
```