Homework 1

- This is an introductory homework to familiarize you with handling datasets and visualizing them by using different libraries.
- Please go through the Pandas tutorial in Canvas module 1, prior to attempting this homework.
- There is a Notebook on Canvas called ISLRChapter2Prob8.ipynb which has solutions to Problem 8 from Chapter 2 in the book. Use that as a guide to learn about some Pandas functions.
- Contact the TA regarding any doubts (details available in the office hours tab in Canvas)
- All the data required for all the homeworks will be available in Canvas -Files/Homeworks/Data
- In this homework we will explore the Iris dataset .

Importing relevant libraries

· We start off by importing required libraries

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn import metrics
sns.set()
```

Loading data

- The data is in a csv file: 'Iris.csv'
- Import this file using "pd.read_csv()" command
- When importing the file make sure the notebook is in the same location as the file or specify the path of the file like $data \setminus iris.csv$

```
iris_df = pd.read_csv('Iris.csv')
print(iris_df)

Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm \
0    1    5.1    3.5    1.4    0.2
1    2    4.9    3.0    1.4    0.2
```

```
✓ 1s
                           completed at 11:57 PM
                                                                          X
149 IJU
                    J.9
                                                 J. I
                                                               1.0
           Species
0
       Iris-setosa
1
       Iris-setosa
       Iris-setosa
3
      Iris-setosa
      Iris-setosa
145 Iris-virginica
146 Iris-virginica
147 Iris-virginica
148 Iris-virginica
149 Iris-virginica
[150 rows x 6 columns]
```

Gaining information from data

It is always a good idea to first understand what is your data set. What are the different features etc.

- Use "df.info()" command to get basic information about the datafarme
- Use "df.describe()" command to get statistical information about the dataframe
- · df here refers to dataframe

- use an auphreated) command to get the indices of auphreate rows
- Use " $df['column_name']$.value_counts()" command to get the counts of different species in the dataset.

```
#ommiting the ID column
ID ommited = iris df.loc[:,['SepalLengthCm','SepalWidthCm','PetalLengthCm','PetalWi
print(ID_ommited[ID_ommited.duplicated(keep=False)])
iris df['Species'].value counts()
        SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm Species

1.5 0.1 Iris-setosa
    9
                                                      0.1
                                                             Iris-setosa
    34
                 4.9
                            3.1
                                          1.5
                             3.1
                                                       0.1 Iris-setosa
                                           1.5
    37
                 4.9
                5.8
                            2.7
                                                      1.9 Iris-virginica
    101
                                          5.1
        5.8
                             2.7
                                           5.1
                                                    1.9 Iris-virginica
    142
    Iris-versicolor 50
    Iris-setosa
                    50
    Iris-virginica 50
    Name: Species, dtype: int64
```

Data visualization

In this section you will be required to generate certain plots/graphs from the data and provide your comments/insights from the plots

a) Species count

- For this you are required to produce a histogram about the counts of different species in the datasets
- Use "sns.countplot()"

```
plt.figure(figsize=( 8 , 4 ))
```

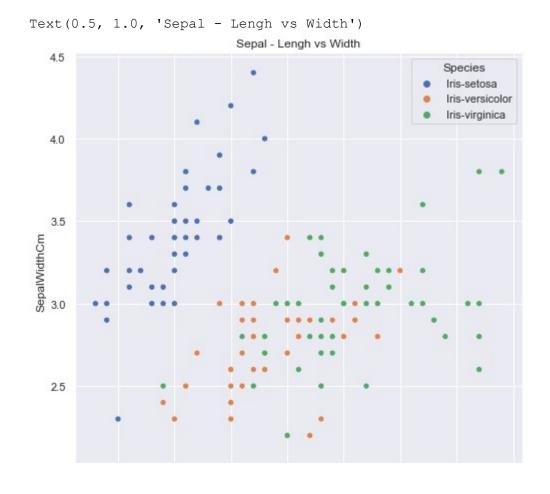
Species

Type your insights here

• __

- b) Uni-variate analysis : Comparison between various species based on sepal length and width
 - For this you are required to produce a **scatter plot between sepal length and sepal width for different species** in the dataset
 - Use "sns.scatterplot()"
 - Set the hue parameter to be the species column of the dataframe

```
plt.figure(figsize=( 8 , 8 ))
sns.scatterplot(
    x = iris_df['SepalLengthCm'],
    y = iris_df['SepalWidthCm'],
    hue = iris_df['Species']
)
plt.title('Sepal - Lengh vs Width')
```

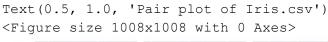


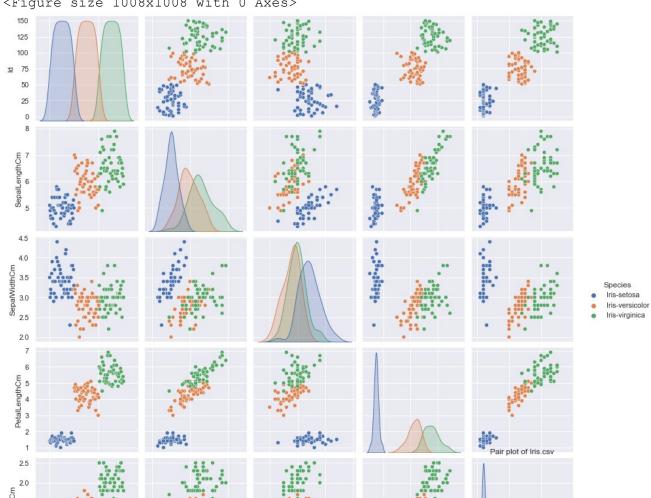
I saw that the iris-setosa had was much shorter on average than iris-versicolor and iris-virginca, but had a larger width. They all seemed to have larger sepa widths when the length of the sepal increased. Iris-versicolor and Iris-virginica clustered together pretty much, showing that they have similar sepal sizes, virginica had slightly larger, but overall they were close.

c) Bi-variate analysis: Plot pairwise relationships

- For this you are required to produce a *pair plot between all the features for different species* in the dataset
- Use "sns.pairplot()"
- Set the hue parameter to be the species column of the dataframe

```
plt.figure(figsize=( 14 , 14 ))
sns.pairplot(iris_df, hue = "Species")
plt.title('Pair plot of Iris.csv')
```

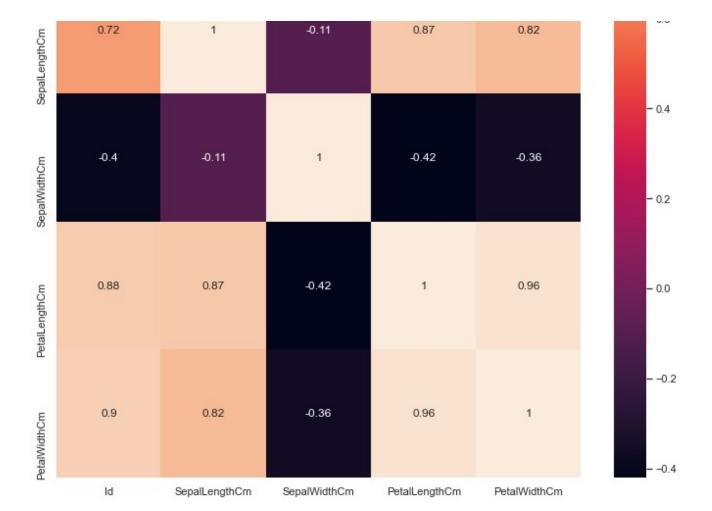




Type your insights here

• looking at the data I saw common groupings that presented of the 3 species of flowers.

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