

# ASSIGNMENT 1

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Class : BE - A

Roll no : 49

Problem Statement : Download the Iris flower dataset or any other dataset into a DataFrame. (eg <https://archive.ics.uci.edu/ml/datasets/Iris> (<https://archive.ics.uci.edu/ml/datasets/Iris>) ) Use Python/R and



## Print the Iris Dataset

In [61]:

```
iris_read.head()
```

```
Id Sepal Length Sepal Width Petal Length Petal Width Species 0 1 5.1
3.5 1.4 0.2 Iris-setosa 1 2 4.9 3.0 1.4 0.2 Iris-setosa 2 3 4.7 3.2
1.3 0.2 Iris-setosa 3 4 4.6 3.1 1.5 0.2 Iris-setosa 4 5 5.0 3.6 1.4
0.2 Iris-setosa
```

## 1. How many features are there and what are their types (e.g., numeric, nominal)?

[62]:

```
print
( "Number of features in the dataset : "
, len ( iris_read .
columns )) print ("Type of features present in the dataset:" )
iris_read . dtypes
```

Number of features in the dataset : 6 Type  
of features present in the dataset:

Out[62]:

In

Id int64  
Sepal Length float64  
Sepal Width float64  
Petal Length float64  
Petal Width float64  
Species object  
dtype: object

## 2. Compute and display summary statistics for each feature available in the dataset. (eg. minimum value, maximum value, mean, range, standard deviation, variance and percentiles

In [63]:

```
iris_read.describe()
```

Out[63]:

	Id	Sepal Length	Sepal Width	Petal Length	Petal Width
count	150.000000	150.000000	150.000000	150.000000	150.000000
mean	75.500000	5.843333	3.054000	3.758667	1.198667
std	43.445368	0.828066	0.433594	1.764420	0.763161
min	1.000000	4.300000	2.000000	1.000000	0.100000
25%	38.250000	5.100000	2.800000	1.600000	0.300000
50%	75.500000	5.800000	3.000000	4.350000	1.300000
75%	112.750000	6.400000	3.300000	5.100000	1.800000



	Id	Sepal Length	Sepal Width	Petal Length	Petal Width	Species
	0	1	5.1	3.5	1.4	0.2 Iris-setosa
	1	2	4.9	3.0	1.4	0.2 Iris-setosa
	2	3	4.7	3.2	1.3	0.2 Iris-setosa
	3	4	4.6	3.1	1.5	0.2 Iris-setosa
	4	5	5.0	3.6	1.4	0.2 Iris-setosa ... ..
145	146	6.7	3.0	5.2	2.3	Iris-virginica
146	147	6.3	2.5	5.0	1.9	Iris-virginica 147 148 6.5 3.0 5.2 2.0

Iris-virginica

**148** 149 6.2 3.4 5.4 2.3 Iris-virginica [In](#)

**149** 150 5.9 3.0 5.1 1.8 Iris-virginica **150** rows × 6 columns

[In](#)



	Id	Sepal Length	Sepal Width	Petal Length	Petal Width	Species
--	----	--------------	-------------	--------------	-------------	---------

<b>0</b>	1	5.1	3.5	1.4	0.2	1
----------	---	-----	-----	-----	-----	---

<b>1</b>	2	4.9	3.0	1.4	0.2	1
----------	---	-----	-----	-----	-----	---

<b>2</b>	3	4.7	3.2	1.3	0.2	1
----------	---	-----	-----	-----	-----	---

<b>3</b>	4	4.6	3.1	1.5	0.2	1
----------	---	-----	-----	-----	-----	---

<b>4</b>	5	5.0	3.6	1.4	0.2	1
----------	---	-----	-----	-----	-----	---

... ..

<b>145</b>	146	6.7	3.0	5.2	2.3	3
------------	-----	-----	-----	-----	-----	---

<b>146</b>	147	6.3	2.5	5.0	1.9	3
------------	-----	-----	-----	-----	-----	---

<b>147</b>	148	6.5	3.0	5.2	2.0	3
------------	-----	-----	-----	-----	-----	---

<b>148</b>	149	6.2	3.4	5.4	2.3	3
------------	-----	-----	-----	-----	-----	---

<b>149</b>	150	5.9	3.0	5.1	1.8	3
------------	-----	-----	-----	-----	-----	---

**150** rows × 6 columns [In \[47\]:](#)



**Out[47]:**

Id int64

Sepal Length float64

Sepal Width float64

Petal Length float64

Petal Width float64

Species int64

dtype: object [In \[21\]:](#)

```
iris_read.print_statistics("Variance is:\n",iris_read.var())
```

Variance is:

```

Id 1887.500000
Sepal Length 0.685694
Sepal Width 0.188004
Petal Length 3.113179
In
Petal Width 0.582414
Species 0.223714
dtype: float64 [36]:

```

```

print(iris_read["Petal Length"], np.percentile(iris_read["Petal Length"], 95))

```

Percentile is : 105.049999999999995 Out[36]:

```

Id Sepal Length Sepal Width Petal Length Petal Width Species
0 1 5.1 3.5 1.4 0.2 1
1 2 4.9 3.0 1.4 0.2 1
2 3 4.7 3.2 1.3 0.2 1
3 4 4.6 3.1 1.5 0.2 1
4 5 5.0 3.6 1.4 0.2 1
... ..
145 146 6.7 3.0 5.2 2.3 1
146 147 6.3 2.5 5.0 1.9 1
147 148 6.5 3.0 5.2 2.0 1
148 149 6.2 3.4 5.4 2.3 1 149 150 5.9 3.0 5.1 1.8 1 150 rows x 6 columns

```

### 3. Data Visualization-Create a histogram for each feature in the dataset to illustrate the feature distributions. Plot each histogram.

```

[44]:
print("Histogram") sns.displot(iris_read['Sepal Length'],kde = False) sns.displot(iris_read['Sepal Width'],kde = False) sns.displot(iris_read['Petal Length'],kde = False) sns.displot(iris_read['Petal Width'],kde = False)
sns.displot(iris_df['Species'],kde = False)

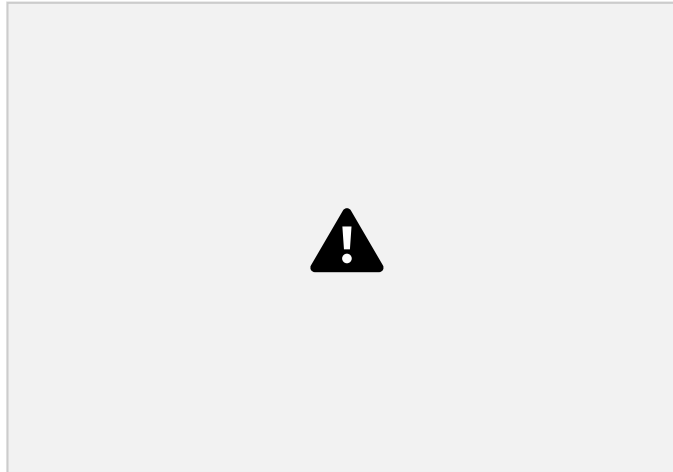
```

Histogram

Out[44]:

<seaborn.axisgrid.FacetGrid at 0x151c9ef8>

In



**4. Create a boxplot for each feature in the dataset. All of the boxplots should be combined into a single plot. Compare distributions and identify outliers. [54]:**

```
plot = sns.countplot(x = 'Sepal Length', data = iris_read)
sns.boxplot(x = 'Sepal Length', y = 'Sepal Width', data = iris_read)
plot.set_xticklabels(plot.get_xticklabels(), rotation=40)
```

Out[54]:

```
[Text(0, 0, '4.3'),
Text(1, 0, '4.4'),
Text(2, 0, '4.5'),
Text(3, 0, '4.6'),
Text(4, 0, '4.7'),
Text(5, 0, '4.8'),
Text(6, 0, '4.9'),
Text(7, 0, '5.0'),
Text(8, 0, '5.1'),
Text(9, 0, '5.2'),
Text(10, 0, '5.3'),
Text(11, 0, '5.4'),
Text(12, 0, '5.5'),
Text(13, 0, '5.6'),
Text(14, 0, '5.7'),
Text(15, 0, '5.8'),
Text(16, 0, '5.9'),
Text(17, 0, '6.0'),
Text(18, 0, '6.1'),
Text(19, 0, '6.2'),
Text(20, 0, '6.3'),
Text(21, 0, '6.4'),
Text(22, 0, '6.5'),
Text(23, 0, '6.6'),
Text(24, 0, '6.7'),
```

In

```
Text(25, 0, '6.8'),
Text(26, 0, '6.9'),
Text(27, 0, '7.0'),
Text(28, 0, '7.1'),
Text(29, 0, '7.2'),
Text(30, 0, '7.3'),
Text(31, 0, '7.4'),
Text(32, 0, '7.6'),
Text(33, 0, '7.7'),
```

```
Text(34, 0, '7.9')]
```



[55]:

```
plot = sns.countplot(x = 'Petal Length', data = iris_read)
sns.boxplot(x = 'Petal Length', y = 'Petal Width', data = iris_read)
plot.set_xticklabels(plot.get_xticklabels(), rotation
40)
```

Out[55]:

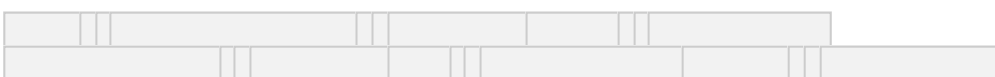
```
[Text(0, 0, '1.0'),
Text(1, 0, '1.1'),
Text(2, 0, '1.2'),
Text(3, 0, '1.3'),
Text(4, 0, '1.4'),
Text(5, 0, '1.5'),
Text(6, 0, '1.6'),
Text(7, 0, '1.7'),
Text(8, 0, '1.9'),
Text(9, 0, '3.0'),
Text(10, 0, '3.3'),
Text(11, 0, '3.5'),
Text(12, 0, '3.6'),
Text(13, 0, '3.7'),
Text(14, 0, '3.8'),
```

```
Text(15, 0, '3.9'),  
Text(16, 0, '4.0'),  
Text(17, 0, '4.1'),  
Text(18, 0, '4.2'),  
Text(19, 0, '4.3'),  
In  
Text(20, 0, '4.4'),  
Text(21, 0, '4.5'),  
Text(22, 0, '4.6'),  
Text(23, 0, '4.7'),  
Text(24, 0, '4.8'),  
Text(25, 0, '4.9'),  
Text(26, 0, '5.0'),  
Text(27, 0, '5.1'), Text(28, 0, '5.2'),  
Text(29, 0, '5.3'),  
Text(30, 0, '5.4'),  
Text(31, 0, '5.5'),  
Text(32, 0, '5.6'),  
Text(33, 0, '5.7'),  
Text(34, 0, '5.8'),  
Text(35, 0, '5.9'),  
Text(36, 0, '6.0'),  
Text(37, 0, '6.1'),  
Text(38, 0, '6.3'),  
Text(39, 0, '6.4'),  
Text(40, 0, '6.6'),  
Text(41, 0, '6.7'), Text(42,  
0, '6.9')]
```



```
[Text(0, 0, 'Iris-setosa'),
```

```
Text(1, 0, 'Iris-versicolor'), Text(2,  
0, 'Iris-virginica')]  
In [70]:
```



```
plot =  
sns.countplot(x = 'Species',data = iris_df) sns.boxplot(x =  
'Species', y = 'Sepal Width', data = iris_df)
```

```
plot.set_xticklabels(plot.get_xticklabels(),rotation = 40)
```

```
Out[70]:  
[Text(0, 0, 'Iris-setosa'), Text(1,  
0, 'Iris-versicolor'),  
Text(2, 0, 'Iris-virginica')]
```





In [71]:

```
plot = sns.countplot(x = 'Species',data = iris_df)
sns.boxplot(x = 'Species', y = 'Petal Length', data
= iris_df)
plot.set_xticklabels(plot.get_xticklabels(),rotation=40)
```

Out[71]:

```
[Text(0, 0, 'Iris-setosa'),
 Text(1, 0, 'Iris-versicolor'), Text(2,
0, 'Iris-virginica')]
```



In [72]:

```
plot = sns.countplot(x = 'Species',data = iris_df)
sns.boxplot(x = 'Species', y = 'Petal Width', data
= iris_df)
plot.set_xticklabels(plot.get_xticklabels(),rotation=40)
```

Out[72]:

```
[Text(0, 0, 'Iris-setosa'),
```

```
Text(1, 0, 'Iris-versicolor'), Text(2,  
0, 'Iris-virginica')]
```



In [ ]: