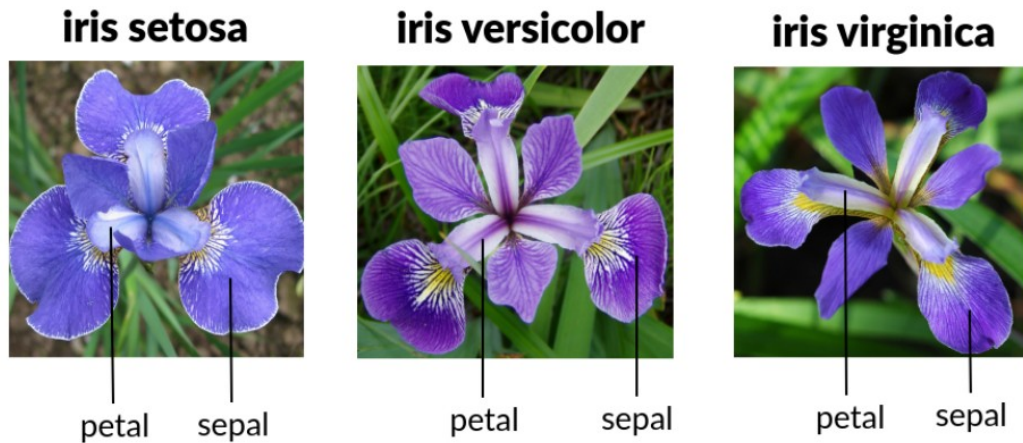


Sakshi Kharat

Oasis Infobyte (Data Science) - Task-1

Iris Flower Classification



Importing the libraries

```
import pandas as pd
import matplotlib.pyplot as plt
```

Loading the Data

```
from google.colab import files
```

```
uploaded = files.upload()
```

```
<IPython.core.display.HTML object>
```

```
Saving Iris.csv to Iris (1).csv
```

```
import numpy as np
```

```
df = pd.read_csv('Iris.csv')
```

Viewing the Dataset

```
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
2	3	4.7	3.2	1.3	0.2
3	4	4.6	3.1	1.5	0.2
4	5	5.0	3.6	1.4	0.2
...
145	146	6.7	3.0	5.2	2.3
146	147	6.3	2.5	5.0	1.9
147	148	6.5	3.0	5.2	2.0
148	149	6.2	3.4	5.4	2.3
149	150	5.9	3.0	5.1	1.8

	Species
0	Iris-setosa
1	Iris-setosa
2	Iris-setosa
3	Iris-setosa
4	Iris-setosa
...	...
145	Iris-virginica
146	Iris-virginica
147	Iris-virginica
148	Iris-virginica
149	Iris-virginica

[150 rows x 6 columns]

Displaying the Dataset

df.info

```
<bound method DataFrame.info of
PetalLengthCm  PetalWidthCm  \
0      1      5.1      3.5      1.4      0.2
1      2      4.9      3.0      1.4      0.2
2      3      4.7      3.2      1.3      0.2
3      4      4.6      3.1      1.5      0.2
4      5      5.0      3.6      1.4      0.2
...    ...    ...    ...    ...    ...
145   146      6.7      3.0      5.2      2.3
146   147      6.3      2.5      5.0      1.9
147   148      6.5      3.0      5.2      2.0
148   149      6.2      3.4      5.4      2.3
149   150      5.9      3.0      5.1      1.8
```

	Species
0	Iris-setosa
1	Iris-setosa
2	Iris-setosa
3	Iris-setosa
4	Iris-setosa

```

..      ...
145  Iris-virginica
146  Iris-virginica
147  Iris-virginica
148  Iris-virginica
149  Iris-virginica

```

```
[150 rows x 6 columns]>
```

Modifying the dataset by removing any Missing Values using fillna() method

```
df.isnull().sum()
```

```

Id                0
SepalLengthCm     0
SepalWidthCm      0
PetalLengthCm     0
PetalWidthCm      0
Species           0
dtype: int64

```

The Values are 0 meaning it has no Null Values all over the dataset

Viewing the Columns in the dataset

```
df.columns
```

```

Index(['Id', 'SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm',
      'PetalWidthCm',
      'Species'],
      dtype='object')

```

```
df.describe
```

```

<bound method NDFrame.describe of      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
2           3           4.7           3.2           1.3           0.2
3           4           4.6           3.1           1.5           0.2
4           5           5.0           3.6           1.4           0.2
..      ...      ...      ...      ...      ...
145  146           6.7           3.0           5.2           2.3
146  147           6.3           2.5           5.0           1.9
147  148           6.5           3.0           5.2           2.0
148  149           6.2           3.4           5.4           2.3
149  150           5.9           3.0           5.1           1.8

      Species
0  Iris-setosa
1  Iris-setosa
2  Iris-setosa

```

```

3      Iris-setosa
4      Iris-setosa
..
145   Iris-virginica
146   Iris-virginica
147   Iris-virginica
148   Iris-virginica
149   Iris-virginica

```

```
[150 rows x 6 columns]>
```

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

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	
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
5	6	5.4	3.9	1.7	0.4	Iris-
						setosa
6	7	4.6	3.4	1.4	0.3	Iris-
						setosa
7	8	5.0	3.4	1.5	0.2	Iris-
						setosa
8	9	4.4	2.9	1.4	0.2	Iris-
						setosa
9	10	4.9	3.1	1.5	0.1	Iris-
						setosa

```
df.shape
```

```
(150, 6)
```

```
print(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	
2	3	4.7	3.2	1.3	0.2	
3	4	4.6	3.1	1.5	0.2	
4	5	5.0	3.6	1.4	0.2	
..	
145	146	6.7	3.0	5.2	2.3	
146	147	6.3	2.5	5.0	1.9	

147	148	6.5	3.0	5.2	2.0
148	149	6.2	3.4	5.4	2.3
149	150	5.9	3.0	5.1	1.8

```

    Species
0      Iris-setosa
1      Iris-setosa
2      Iris-setosa
3      Iris-setosa
4      Iris-setosa
...
145 Iris-virginica
146 Iris-virginica
147 Iris-virginica
148 Iris-virginica
149 Iris-virginica

```

[150 rows x 6 columns]

```
print(df[10:21])
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
Species					
10	11	5.4	3.7	1.5	0.2
	Iris-setosa				
11	12	4.8	3.4	1.6	0.2
	Iris-setosa				
12	13	4.8	3.0	1.4	0.1
	Iris-setosa				
13	14	4.3	3.0	1.1	0.1
	Iris-setosa				
14	15	5.8	4.0	1.2	0.2
	Iris-setosa				
15	16	5.7	4.4	1.5	0.4
	Iris-setosa				
16	17	5.4	3.9	1.3	0.4
	Iris-setosa				
17	18	5.1	3.5	1.4	0.3
	Iris-setosa				
18	19	5.7	3.8	1.7	0.3
	Iris-setosa				
19	20	5.1	3.8	1.5	0.3
	Iris-setosa				
20	21	5.4	3.4	1.7	0.2
	Iris-setosa				

```
sliced_data=df[10:21]
```

```
print(sliced_data)
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
Species					

10	11	5.4	3.7	1.5	0.2
Iris-setosa					
11	12	4.8	3.4	1.6	0.2
Iris-setosa					
12	13	4.8	3.0	1.4	0.1
Iris-setosa					
13	14	4.3	3.0	1.1	0.1
Iris-setosa					
14	15	5.8	4.0	1.2	0.2
Iris-setosa					
15	16	5.7	4.4	1.5	0.4
Iris-setosa					
16	17	5.4	3.9	1.3	0.4
Iris-setosa					
17	18	5.1	3.5	1.4	0.3
Iris-setosa					
18	19	5.7	3.8	1.7	0.3
Iris-setosa					
19	20	5.1	3.8	1.5	0.3
Iris-setosa					
20	21	5.4	3.4	1.7	0.2
Iris-setosa					

```
df.iloc[5]
```

Id		6
SepalLengthCm		5.4
SepalWidthCm		3.9
PetalLengthCm		1.7
PetalWidthCm		0.4
Species		Iris-setosa

Name: 5, dtype: object

```
df.loc[df["Species"] == "Iris-setosa"]
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
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					
5	6	5.4	3.9	1.7	0.4
Iris-setosa					
6	7	4.6	3.4	1.4	0.3
Iris-setosa					

7	8	5.0	3.4	1.5	0.2
Iris-setosa					
8	9	4.4	2.9	1.4	0.2
Iris-setosa					
9	10	4.9	3.1	1.5	0.1
Iris-setosa					
10	11	5.4	3.7	1.5	0.2
Iris-setosa					
11	12	4.8	3.4	1.6	0.2
Iris-setosa					
12	13	4.8	3.0	1.4	0.1
Iris-setosa					
13	14	4.3	3.0	1.1	0.1
Iris-setosa					
14	15	5.8	4.0	1.2	0.2
Iris-setosa					
15	16	5.7	4.4	1.5	0.4
Iris-setosa					
16	17	5.4	3.9	1.3	0.4
Iris-setosa					
17	18	5.1	3.5	1.4	0.3
Iris-setosa					
18	19	5.7	3.8	1.7	0.3
Iris-setosa					
19	20	5.1	3.8	1.5	0.3
Iris-setosa					
20	21	5.4	3.4	1.7	0.2
Iris-setosa					
21	22	5.1	3.7	1.5	0.4
Iris-setosa					
22	23	4.6	3.6	1.0	0.2
Iris-setosa					
23	24	5.1	3.3	1.7	0.5
Iris-setosa					
24	25	4.8	3.4	1.9	0.2
Iris-setosa					
25	26	5.0	3.0	1.6	0.2
Iris-setosa					
26	27	5.0	3.4	1.6	0.4
Iris-setosa					
27	28	5.2	3.5	1.5	0.2
Iris-setosa					
28	29	5.2	3.4	1.4	0.2
Iris-setosa					
29	30	4.7	3.2	1.6	0.2
Iris-setosa					
30	31	4.8	3.1	1.6	0.2
Iris-setosa					
31	32	5.4	3.4	1.5	0.4
Iris-setosa					

32	33	5.2	4.1	1.5	0.1
Iris-setosa					
33	34	5.5	4.2	1.4	0.2
Iris-setosa					
34	35	4.9	3.1	1.5	0.1
Iris-setosa					
35	36	5.0	3.2	1.2	0.2
Iris-setosa					
36	37	5.5	3.5	1.3	0.2
Iris-setosa					
37	38	4.9	3.1	1.5	0.1
Iris-setosa					
38	39	4.4	3.0	1.3	0.2
Iris-setosa					
39	40	5.1	3.4	1.5	0.2
Iris-setosa					
40	41	5.0	3.5	1.3	0.3
Iris-setosa					
41	42	4.5	2.3	1.3	0.3
Iris-setosa					
42	43	4.4	3.2	1.3	0.2
Iris-setosa					
43	44	5.0	3.5	1.6	0.6
Iris-setosa					
44	45	5.1	3.8	1.9	0.4
Iris-setosa					
45	46	4.8	3.0	1.4	0.3
Iris-setosa					
46	47	5.1	3.8	1.6	0.2
Iris-setosa					
47	48	4.6	3.2	1.4	0.2
Iris-setosa					
48	49	5.3	3.7	1.5	0.2
Iris-setosa					
49	50	5.0	3.3	1.4	0.2
Iris-setosa					

```
df["Species"].value_counts()
```

```
Iris-setosa      50
Iris-versicolor  50
Iris-virginica   50
Name: Species, dtype: int64
```

```
sum = df["SepalLengthCm"].sum()
```

```
print(sum)
```

```
876.5
```

```
mean = df["SepalLengthCm"].mean()
```



```
print(mean)
5.843333333333334
min = df["SepalLengthCm"].min()
print(min)
4.3
max = df["SepalLengthCm"].max()
```

Preprocessing The Data

```
df.isnull()

```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
Species					
0	False	False	False	False	False
False					
1	False	False	False	False	False
False					
2	False	False	False	False	False
False					
3	False	False	False	False	False
False					
4	False	False	False	False	False
False					
..
...					
145	False	False	False	False	False
False					
146	False	False	False	False	False
False					
147	False	False	False	False	False
False					
148	False	False	False	False	False
False					
149	False	False	False	False	False
False					

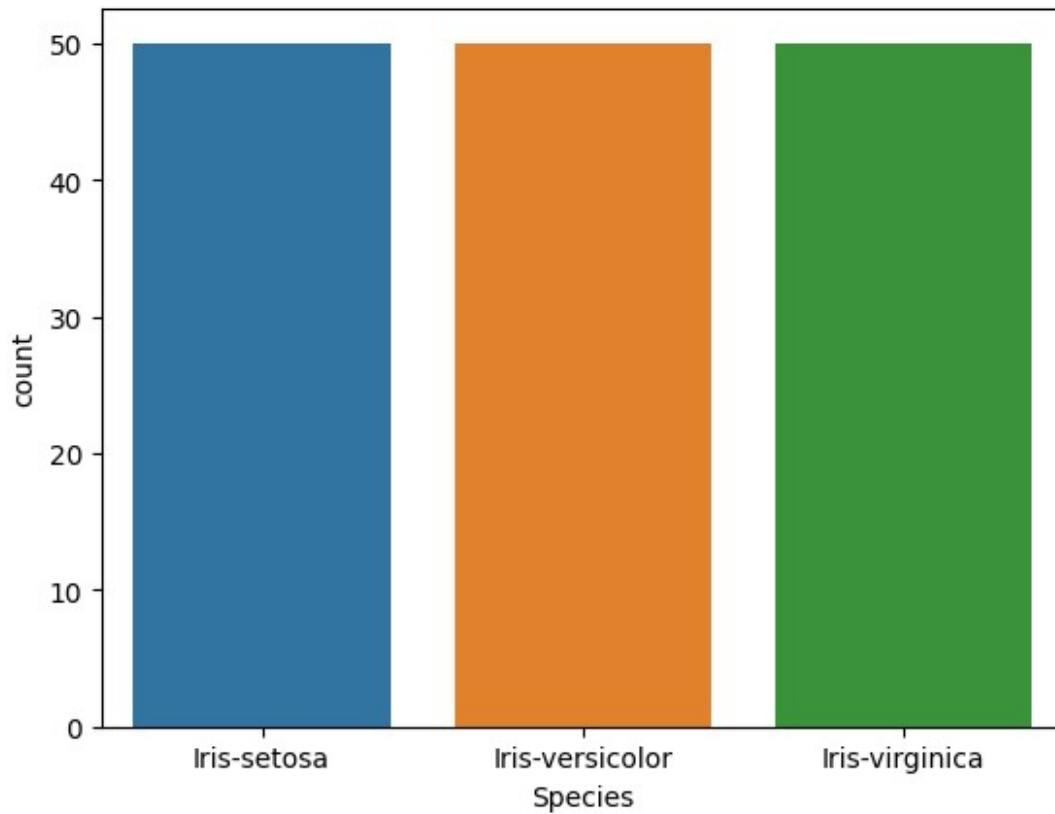
```
[150 rows x 6 columns]
```

Data Visualization

```
import seaborn as sns
iris = sns.load_dataset("iris")
```

Count Plot

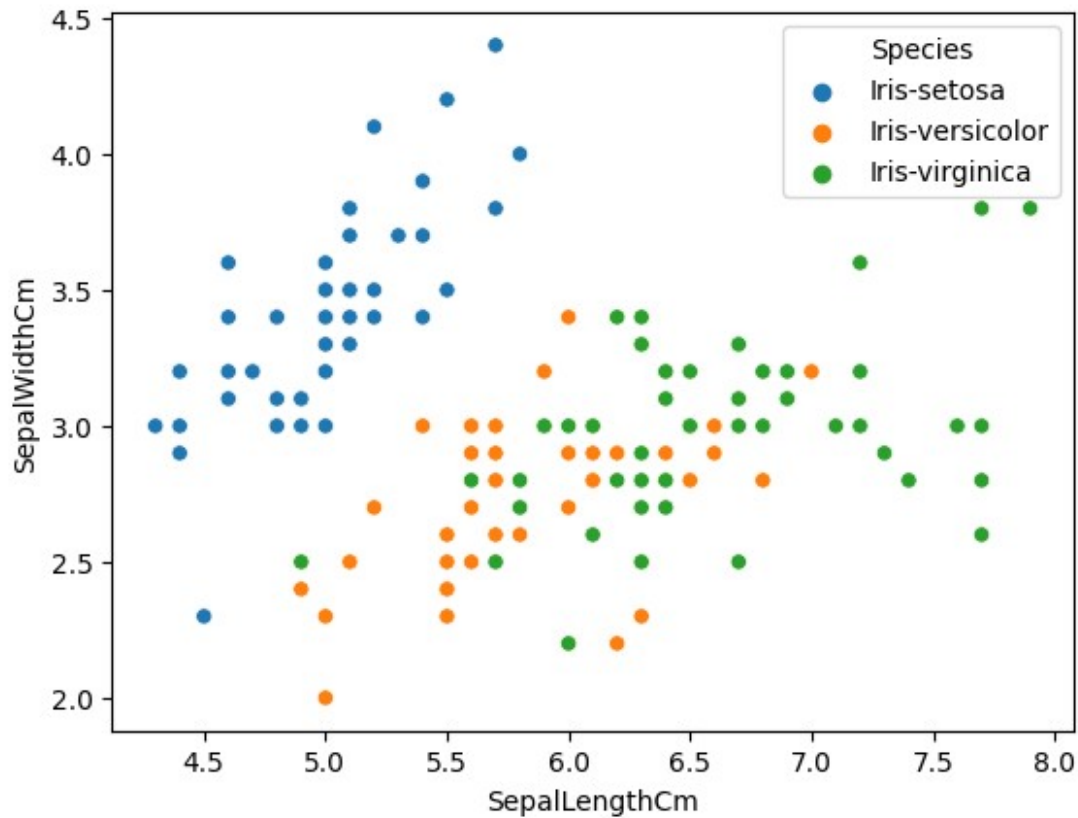
```
sns.countplot(x='Species', data=df, )
plt.show()
```



According to the Plot the total rows are 150 and on which 50 are Iris-Setosa, 50 are Iris-Versicolor and 50 are Iris-Virginica

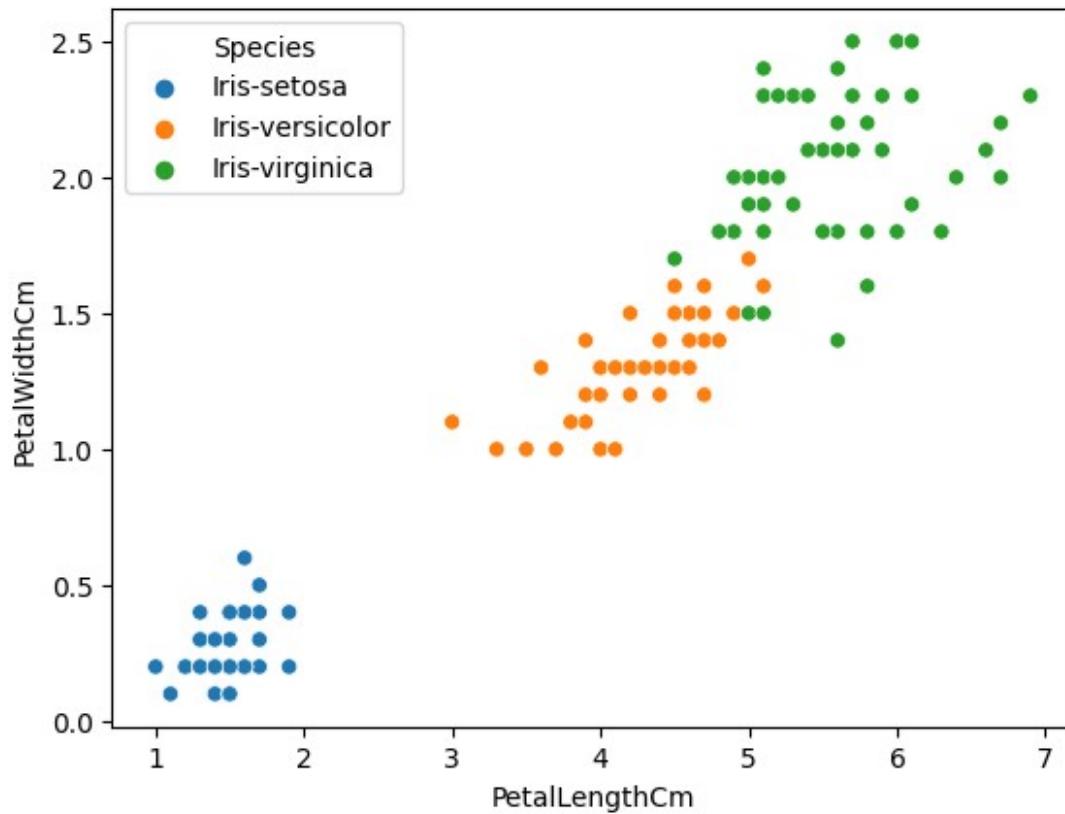
Scatter Plot

```
sns.scatterplot(x='SepalLengthCm', y='SepalWidthCm',  
                hue='Species', data=df, )  
<Axes: xlabel='SepalLengthCm', ylabel='SepalWidthCm'>
```



According to the above plot 1) Iris-Setosa has smaller Sepal Length and larger Sepal Widths 2) Iris-Versicolor is the medium range of Sepal Length and Sepal Width 3) Iris-Virginica has larger sepal lengths and larger sepal width

```
sns.scatterplot(x='PetalLengthCm', y='PetalWidthCm',
                hue='Species', data=df, )
<Axes: xlabel='PetalLengthCm', ylabel='PetalWidthCm'>
```



This is formatted as code

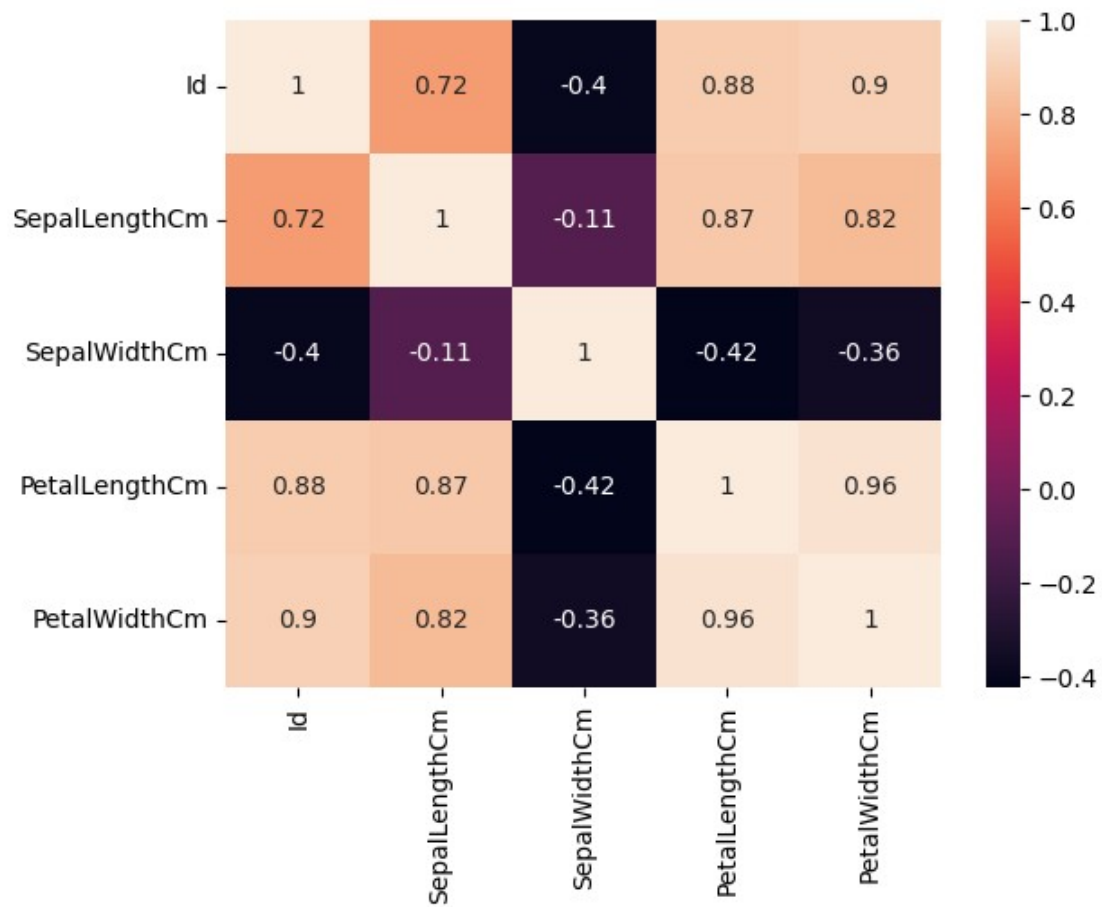
This plot explains that Iris Setosa has smaller Petal Length and Smaller Petal width, while Iris- Versicolor lies in the middle and Iris- Virginica has Larger Petal Width and Smaller Petal Length

Heat Map

```
import seaborn as sns
import matplotlib.pyplot as plt
```

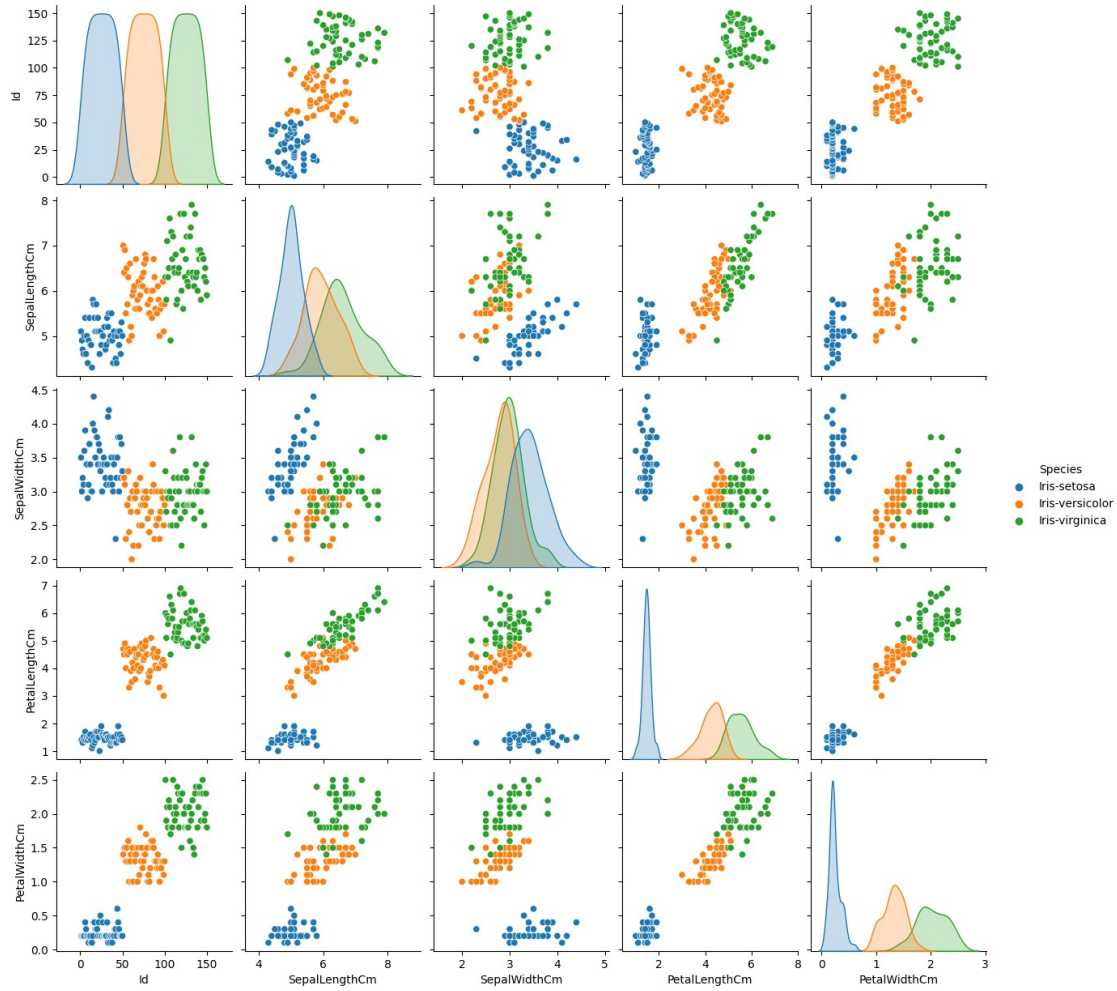
```
sns.heatmap(df.corr(method='pearson'),
             annot = True);
```

```
plt.show()
```



Pair Plot

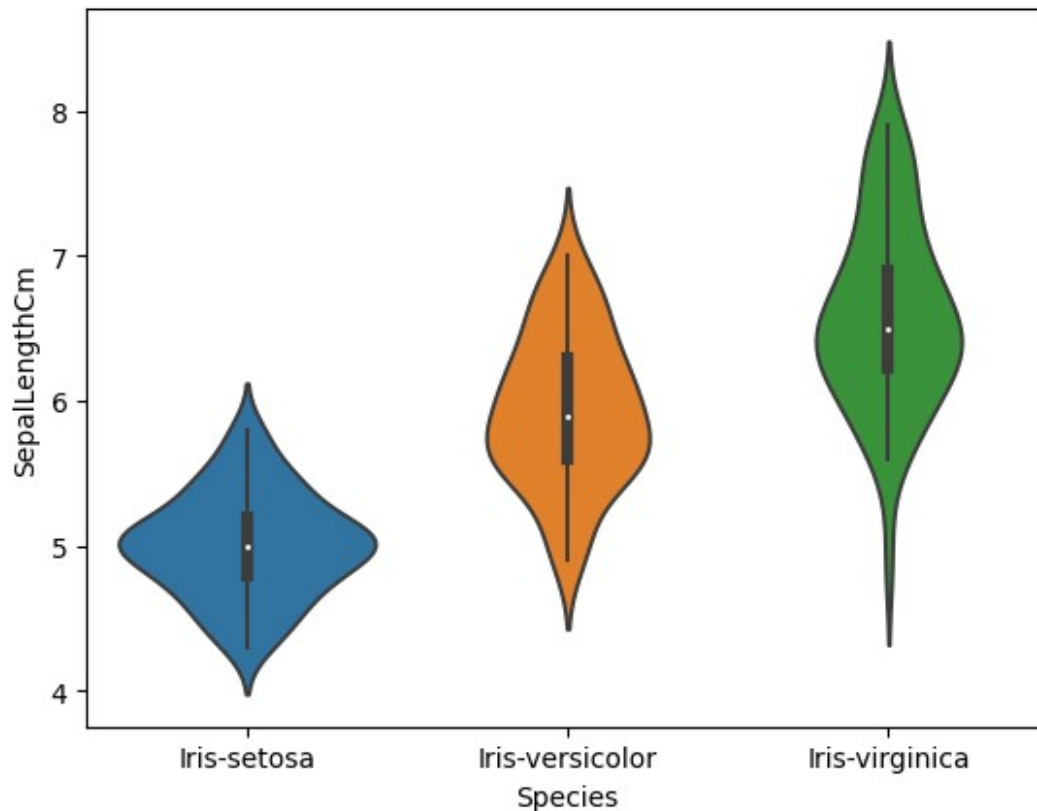
```
g = sns.pairplot(df, hue="Species")
```



Violin Plot

```
sns.violinplot(data=df, x='Species', y='Sepal.LengthCm')
```

```
<Axes: xlabel='Species', ylabel='Sepal.LengthCm'>
```



Histogram

```
import seaborn as sns
import matplotlib.pyplot as plt
```

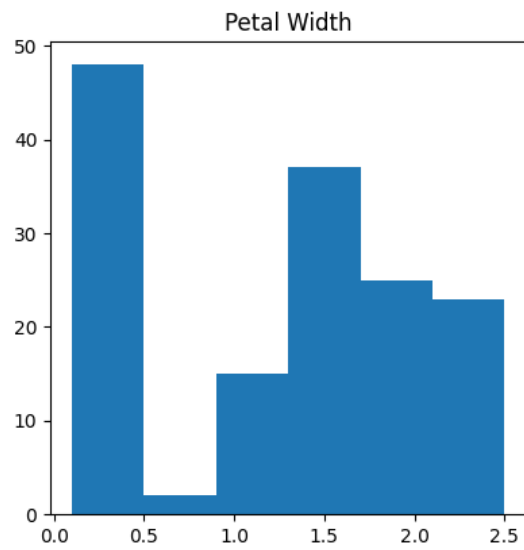
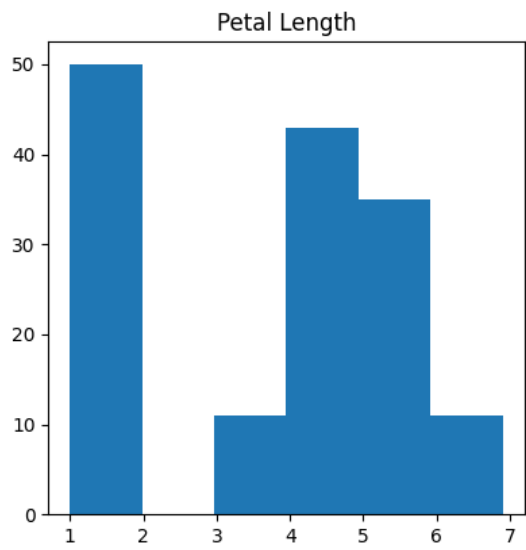
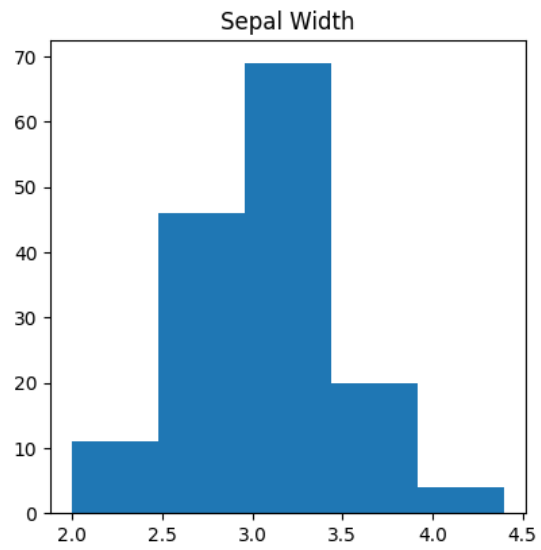
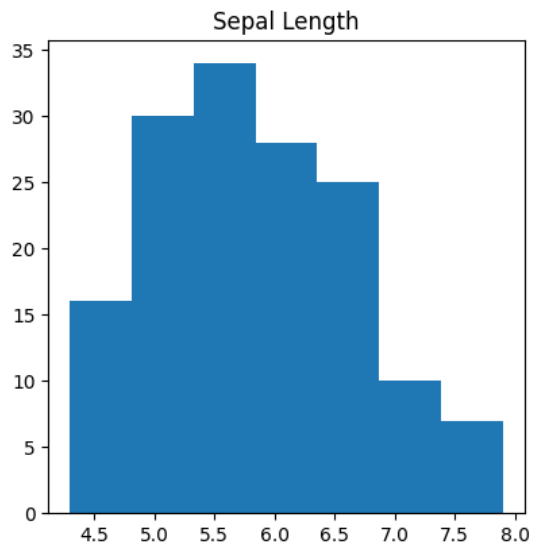
```
fig, axes = plt.subplots(2, 2, figsize=(10,10))
```

```
axes[0,0].set_title("Sepal Length")
axes[0,0].hist(df['SepalLengthCm'], bins=7)
```

```
axes[0,1].set_title("Sepal Width")
axes[0,1].hist(df['SepalWidthCm'], bins=5);
```

```
axes[1,0].set_title("Petal Length")
axes[1,0].hist(df['PetalLengthCm'], bins=6);
```

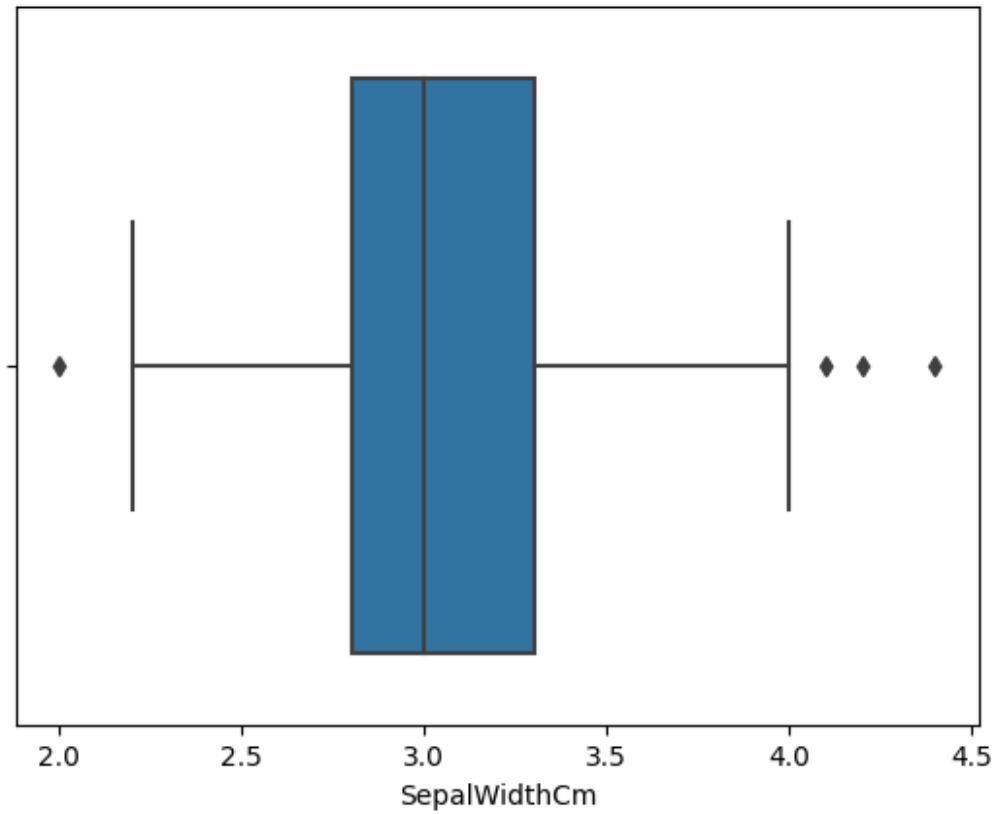
```
axes[1,1].set_title("Petal Width")
axes[1,1].hist(df['PetalWidthCm'], bins=6);
```



Box Plot

```
sns.boxplot(x='SepalWidthCm', data=df)
```

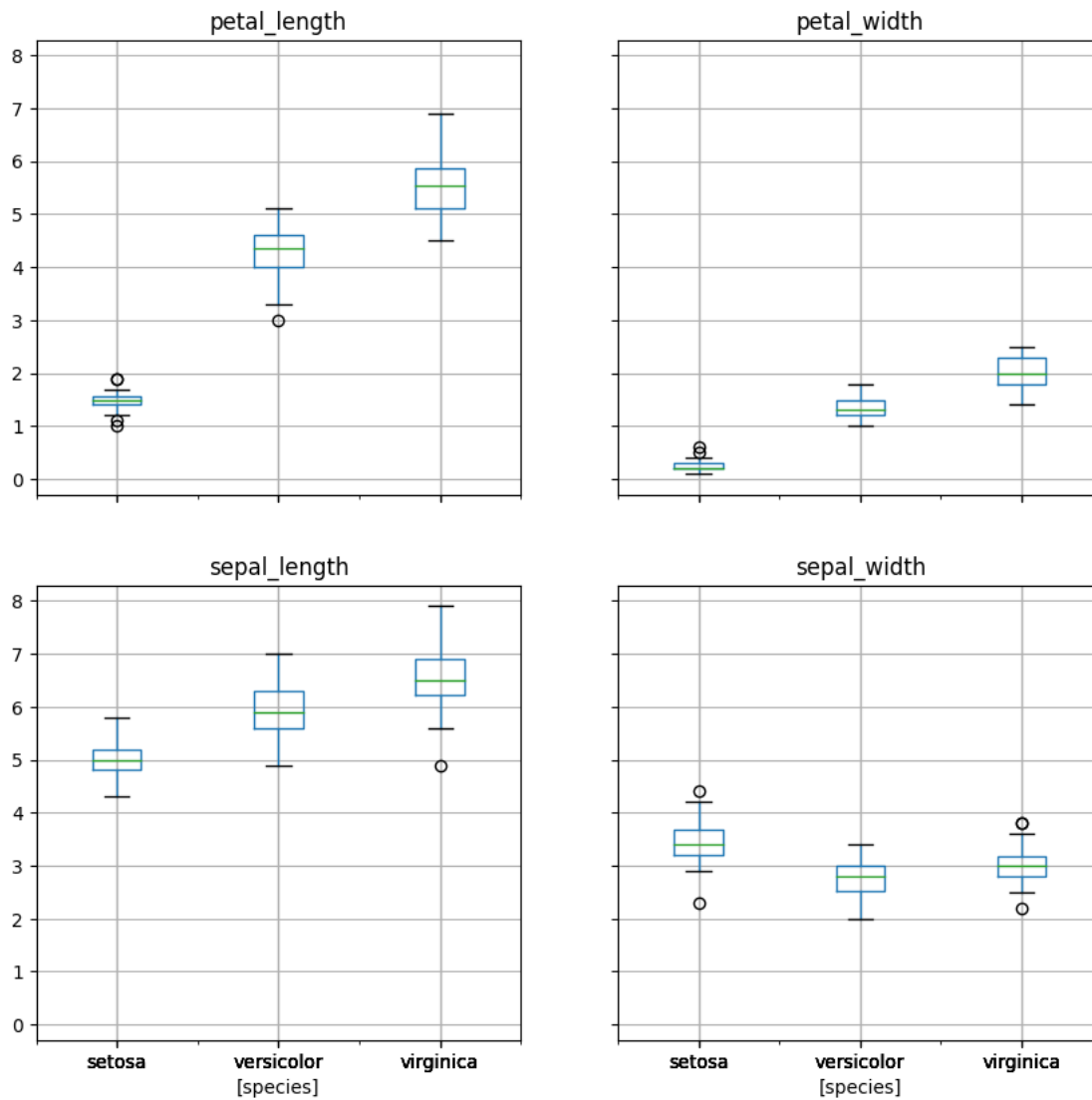
```
<Axes: xlabel='SepalWidthCm'>
```

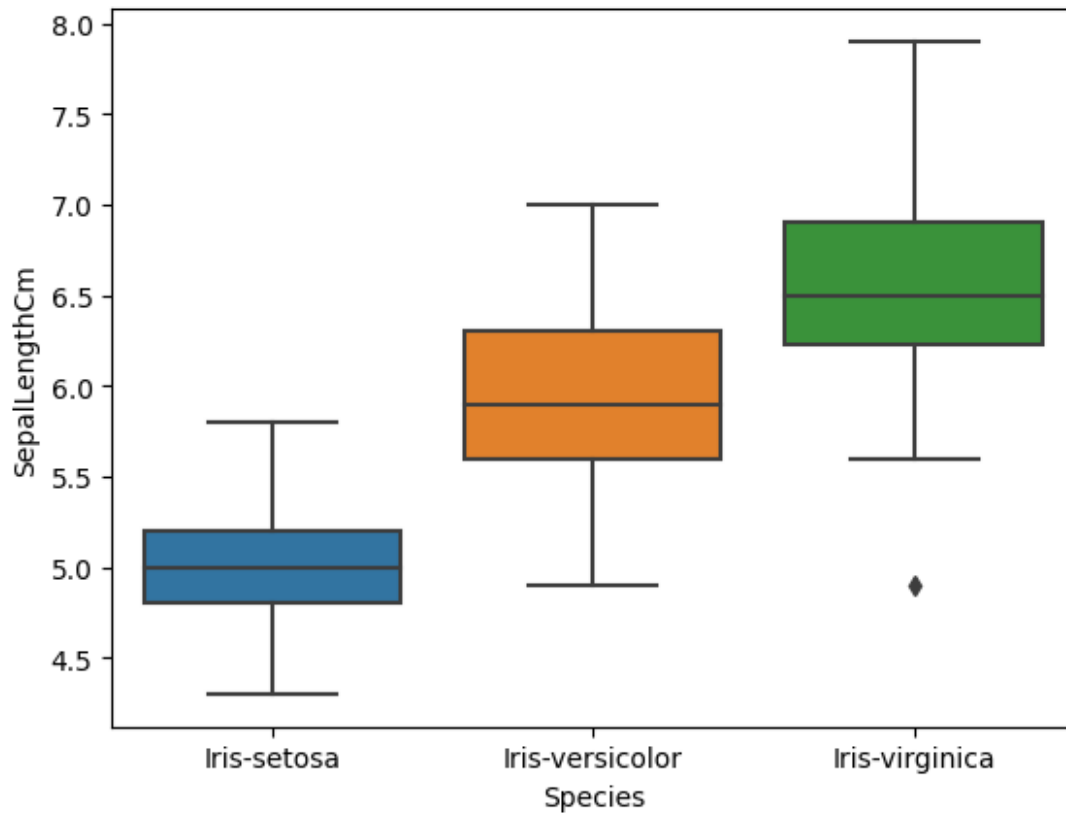
The values above 4 and below 2 are acting as outliers.

```
iris.boxplot(by = 'species', figsize=(10,10))  
plt.show()
```

Boxplot grouped by species



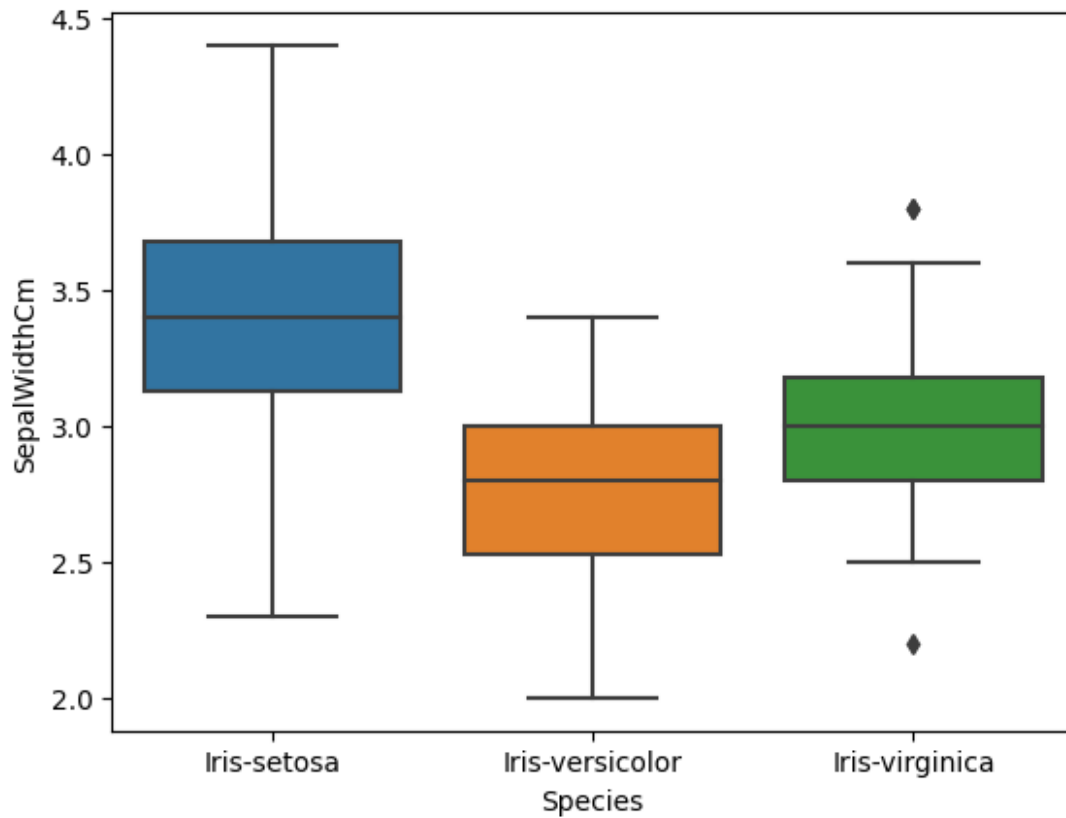
```
def graph(y):  
    sns.boxplot(x="Species", y=y, data=df)  
    plt.figure(figsize=(10,10))  
    graph('SepalLengthCm')  
  
plt.show()
```



<Figure size 1000x1000 with 0 Axes>

```
def graph(y):  
    sns.boxplot(x="Species", y=y, data=df)  
    plt.figure(figsize=(10,10))  
graph('SepalWidthCm')
```

```
plt.show()
```

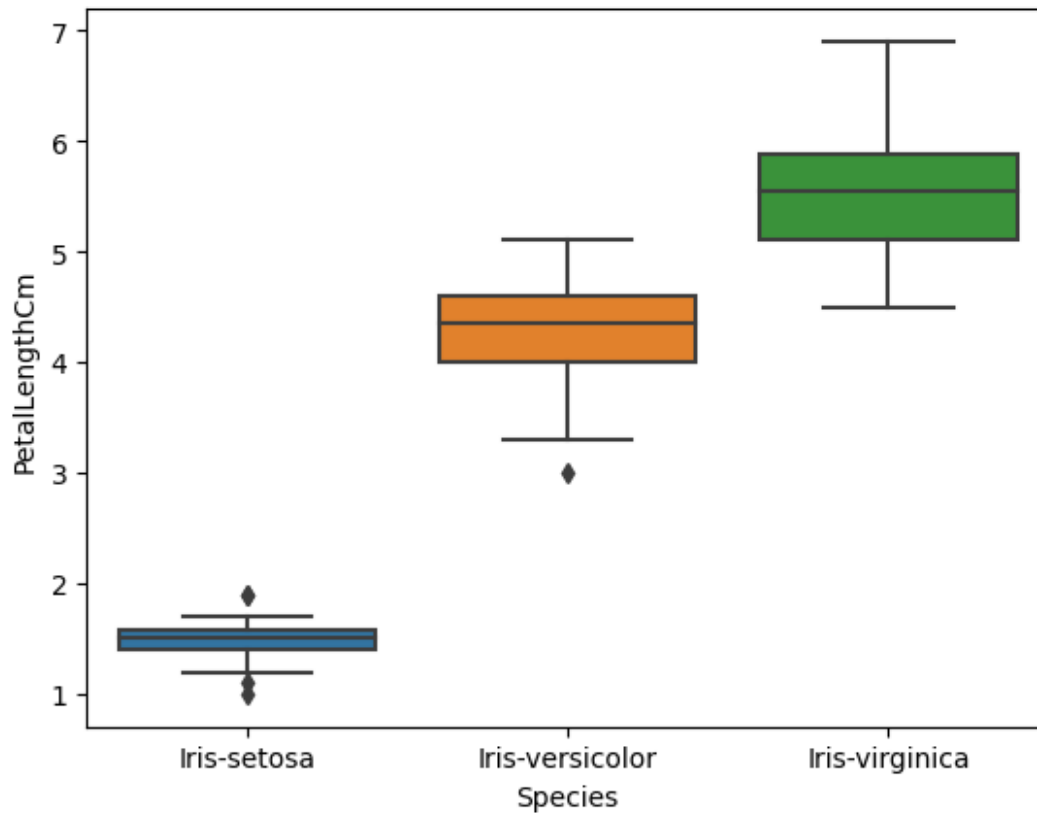


<Figure size 1000x1000 with 0 Axes>

```
def graph(y):  
    sns.boxplot(x="Species", y=y, data=df)  
    plt.figure(figsize=(10,10))
```

```
graph('PetalLengthCm')
```

```
plt.show()
```

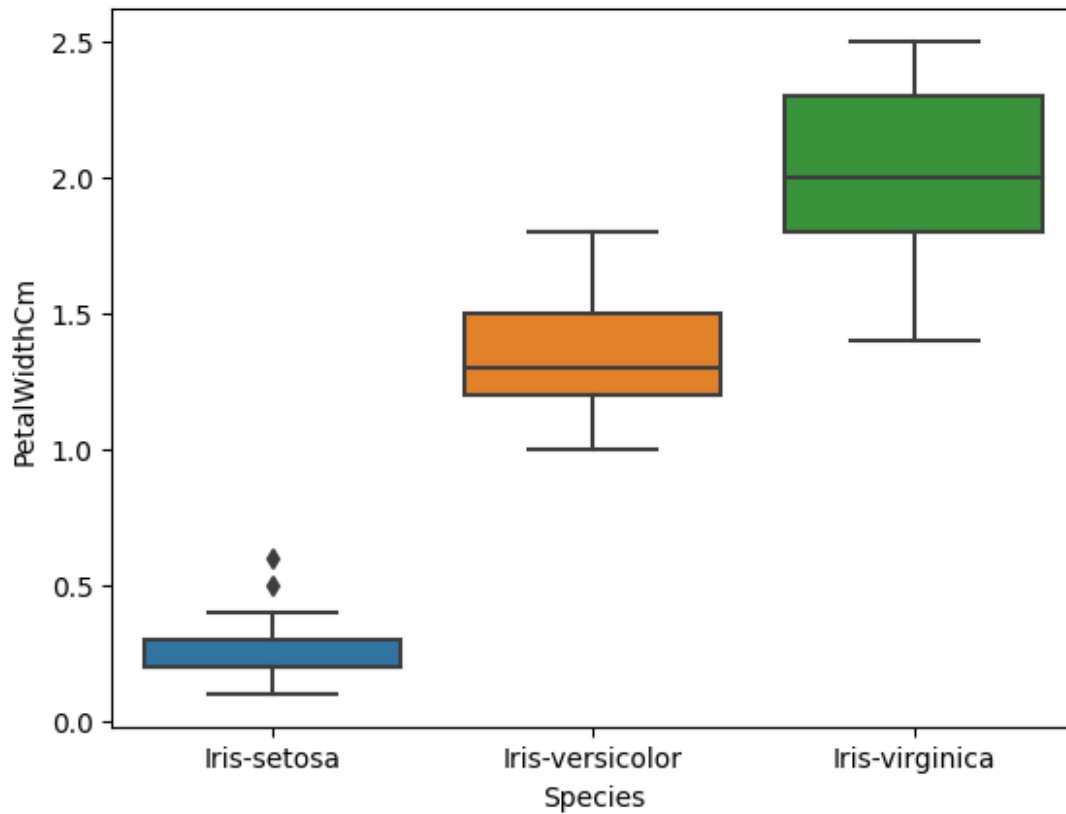


<Figure size 1000x1000 with 0 Axes>

```
def graph(y):  
    sns.boxplot(x="Species", y=y, data=df)  
    plt.figure(figsize=(10,10))
```

```
graph('PetalWidthCm')
```

```
plt.show()
```



<Figure size 1000x1000 with 0 Axes>

Dropping the Id Column

```
df = df.drop('Id', axis=1)
```

df

	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	
Species					
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 x 5 columns]

X contains Independent Variables and Y contains Dependent Variables

```
x = df.iloc[:, :4]
```

x

	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
...
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

[150 rows x 4 columns]

This line selects all rows and the first four columns of the DataFrame df which are **SepalLengthCm, SepalWidthCm, PetalLengthCm, PetalWidthCm** using the iloc function. df.iloc[:, :4] selects all rows (:) and the columns indexed from 0 to 3 (:4). **The resulting x will contain the input features for classification.**

```
y = df.iloc[:, 4]
```

y

0	Iris-setosa
1	Iris-setosa
2	Iris-setosa
3	Iris-setosa
4	Iris-setosa
...	
145	Iris-virginica
146	Iris-virginica
147	Iris-virginica
148	Iris-virginica
149	Iris-virginica

Name: Species, Length: 150, dtype: object

This line selects all rows and the fifth column of the DataFrame df which is (**Species**), which corresponds to the target labels for classification. Again, df.iloc[:, 4] selects all rows (:) and the column at index 4. **The resulting y will contain the target labels**

Import train_test_split to split the data into train and test datasets.

```
from sklearn.model_selection import train_test_split
x_train, x_test, y_train, y_test = train_test_split(x, y,
test_size=0.2, random_state=45)

x_train.shape

(120, 4)
```

x_train has a shape of (112, 4), it means that there are 112 training samples, and each sample has 4 features or input variables.

```
x_test.shape

(30, 4)
```

x_test has a shape of (38, 4), it means that there are 38 test samples, and each sample has 4 features or input variables.

```
y_train.shape

(120,)

y_test.shape

(30,)
```

Creating Model - Classification -- Classifying the Iris Flower Dataset using the Logistic Regression

```
from sklearn.linear_model import LogisticRegression
import warnings
warnings.filterwarnings('ignore')

from sklearn.metrics import accuracy_score, confusion_matrix,
classification_report

from sklearn import preprocessing
from sklearn import utils

lab = preprocessing.LabelEncoder()
encoded = lab.fit_transform(y_train)

print(utils.multiclass.type_of_target(y_train))

multiclass

print(utils.multiclass.type_of_target(encoded))
```



```
multiclass
```

```
model = LogisticRegression()
```

Training the model using the fit method -- Passing the x_train and y_train in the fit function

```
threshold = 0.5
```

```
model.fit(x_train,y_train)
```

```
LogisticRegression()
```

Predicting the results using Predict Method

```
y_pred=model.predict(x_test)
```

```
print(classification_report(y_test, y_pred))
```

	precision	recall	f1-score	support
Iris-setosa	1.00	1.00	1.00	11
Iris-versicolor	0.88	1.00	0.93	7
Iris-virginica	1.00	0.92	0.96	12
accuracy			0.97	30
macro avg	0.96	0.97	0.96	30
weighted avg	0.97	0.97	0.97	30

```
print("Accuracy Score:" , accuracy_score(y_test, y_pred))
```

Accuracy Score: 0.9666666666666667

```
print("Accuracy Score:" , accuracy_score(y_test, y_pred) *100)
```

Accuracy Score: 96.66666666666667

Accuracy is 96.66

Predicting the Model

```
y_pred = model.predict([[5.0, 3.6, 1.4, 0.2]])
```

```
print(*y_pred)
```

Iris-setosa

Predicted the Species of Iris Flower which is IRIS- SETOSA

KNN alogrithm

```
from sklearn.neighbors import KNeighborsClassifier
```

```
model = KNeighborsClassifier()
model.fit(x_test, y_test)
KNeighborsClassifier()
print("Accuracy:" ,model.score(x_test, y_test) )
Accuracy: 1.0
print("Accuracy:" ,model.score(x_test, y_test) * 100)
Accuracy: 100.0
y_pred = model.predict([[5.6, 2.5, 3.9, 1.1]])
print(*y_pred)
Iris-versicolor
```

The predicted flower is IRIS-VERSICOLOR

Decision Tree

```
from sklearn.tree import DecisionTreeClassifier
model = DecisionTreeClassifier()
model.fit(x_train, y_train)
DecisionTreeClassifier()
print("Accuracy:" ,model.score(x_test, y_test))
Accuracy: 1.0
print("Accuracy:", model.score(x_test, y_test) *100)
Accuracy: 100.0
```

Accuracy of the model is 100

Predict the Model

```
y_pred = model.predict([[6.2, 2.8, 4.8, 1.8]])
print(*y_pred)
Iris-virginica
```

Predicted flower is IRIS-VIRGINICA

Random Forest

```
from sklearn.ensemble import RandomForestClassifier
model = RandomForestClassifier()
```

```
model.fit(x_train, y_train)
RandomForestClassifier()
RF_predictions = model.predict(x_test)
print("Accuracy:", model.score(x_test, y_test) *100)
Accuracy: 96.66666666666667
```

Accuracy of the model is 93.33

```
y_pred = model.predict([[6.9, 1.8, 4.4, 1.6]])
print(*y_pred)
```

Iris-versicolor

Predicted flower is IRIS-Versicolor

Support Vector Machine Algorithm

```
from sklearn.svm import SVC
model = SVC()
model.fit(x_train, y_train)
SVC()
SVM_predictions = model.predict(x_test)
print("Accuracy:", model.score(x_test, y_test) *100)
Accuracy: 96.66666666666667
```

Accuracy of the model is 96.66

```
y_pred = model.predict([[5.1, 3.5, 1.4, 0.2]])
print(*y_pred)
```

Iris-setosa

Predicted flower is IRIS-SETOSA

End of the code