

```
In [ ]: Objective is to classify the flower belong to which category among the all three?
```

```
In [ ]: # Importing Libraries
```

```
In [5]: import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
```

```
In [6]: data = pd.read_csv(r'C:\Users\Hp\Desktop\Pyhton\Python Project_Raw file\IRIS Data\Iris.c
```

```
In [7]: #Understand the data
data.head(2)
```

```
Out[7]:
```

	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

```
In [18]: #To Get an Overview of No of column Present In dataset
data.columns
```

```
Out[18]: Index(['Id', 'sepalength', 'sepalwidth', 'petallength', 'petalwidth',
               'Species'],
              dtype='object')
```

```
In [19]: #rename the column name and Make it readable
data.rename(columns={'SepalLengthCm':'sepalength','SepalWidthCm':'sepalwidth','PetalLen
```

```
In [20]: data.head(2)
```

```
Out[20]:
```

	Id	sepalength	sepalwidth	petallength	petalwidth	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

```
In [21]: #Get the statistical Insights of the data
data.describe()
```

```
Out[21]:
```

	Id	sepalength	sepalwidth	petallength	petalwidth
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
max	150.000000	7.900000	4.400000	6.900000	2.500000

```
In [22]: #We observe from this above line that our dataset has 150 rows and 6 columns
data.shape
```

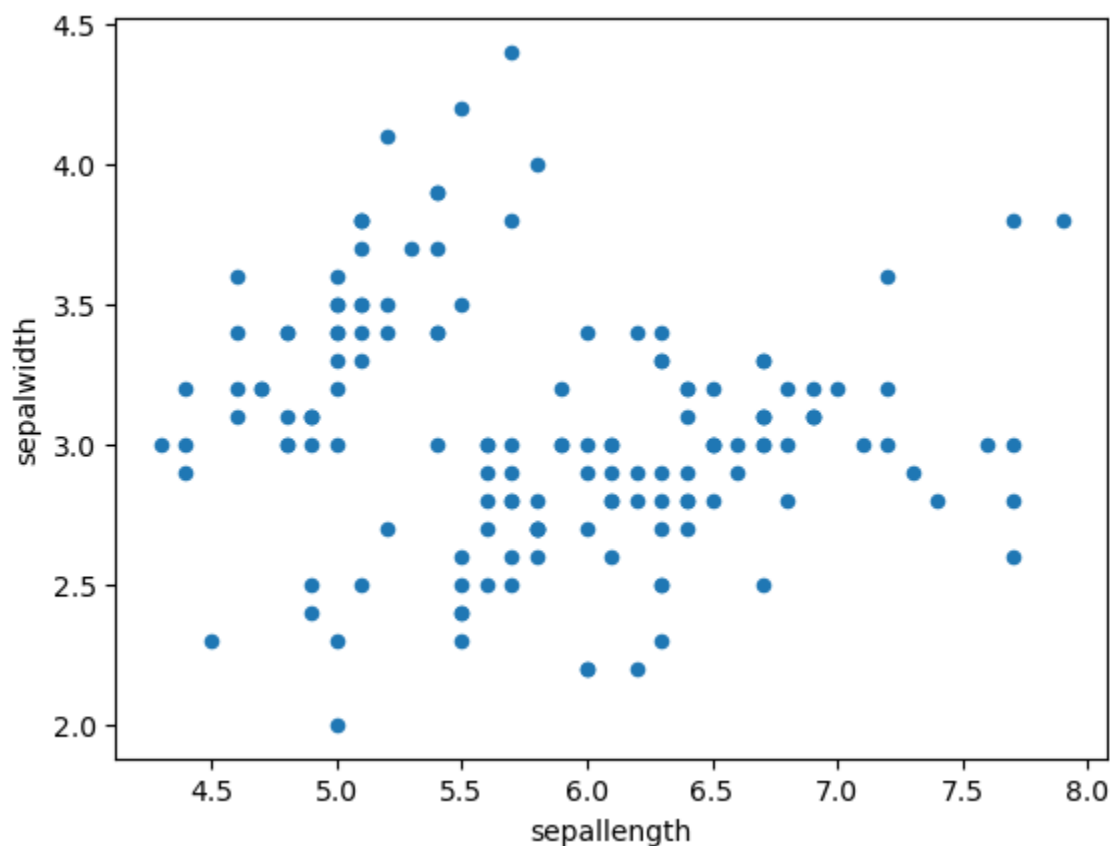
Out[22]: (150, 6)

```
In [23]: #to get more informatio about the dataset  
#iris dataset is a balanced dataset no of datapoints are equal  
data['Species'].value_counts()
```

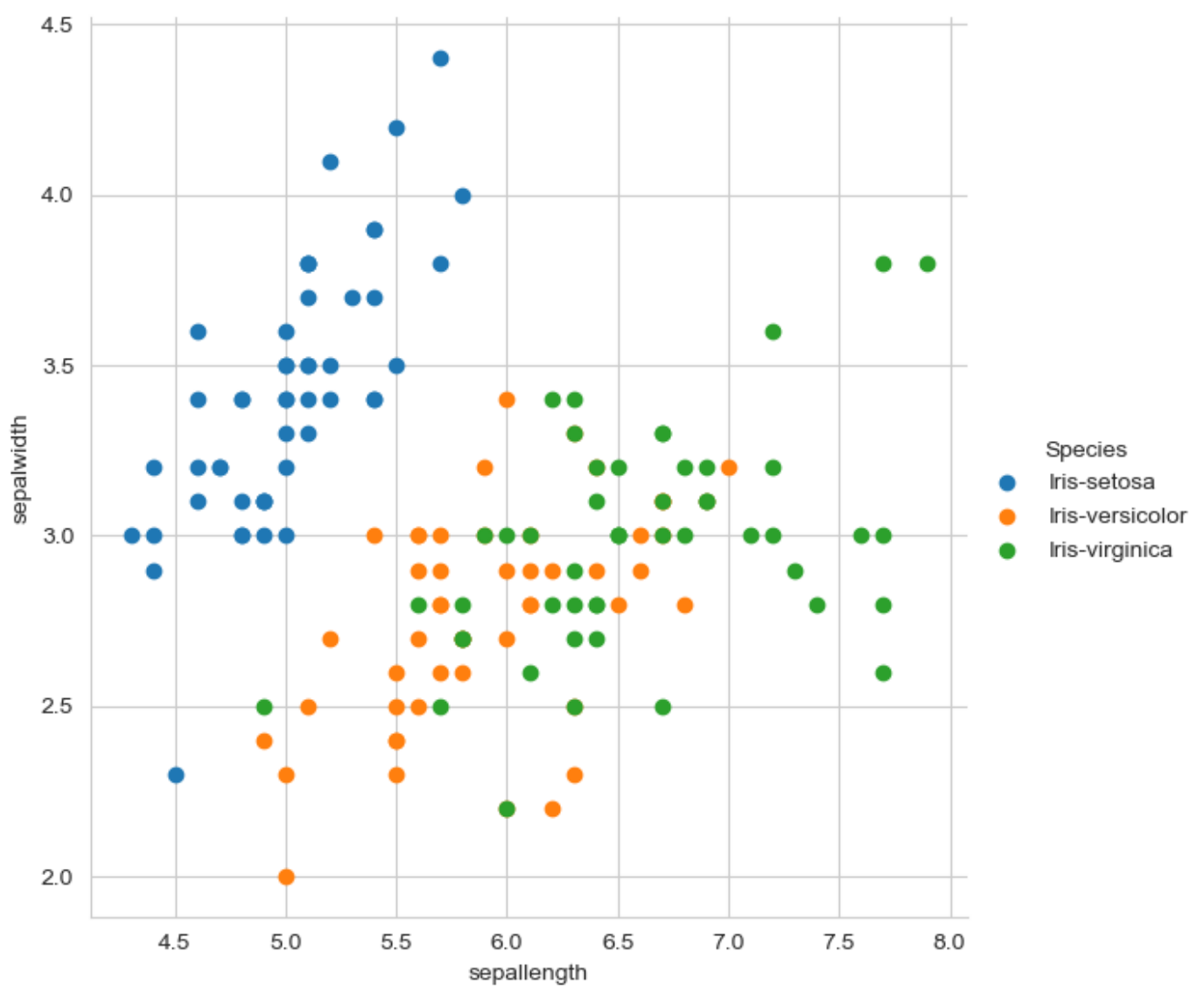
```
Out[23]: Iris-setosa      50  
Iris-versicolor  50  
Iris-virginica   50  
Name: Species, dtype: int64
```

```
In [ ]: #2D Scatter plot  
# to Understand the axis label scale
```

```
In [25]: data.plot(kind='scatter',x='sepallength',y='sepalwidth')  
plt.show()
```



```
In [45]: sns.set_style("whitegrid"),  
sns.FacetGrid(data, hue= "Species", height =6) \  
.map(plt.scatter, "sepallength","sepalwidth") \  
.add_legend();  
plt.show()
```



```
In [31]: data.head(1)
```

```
Out[31]:
```

	sepalength	sepalwidth	petallength	petalwidth	Species
0	5.1	3.5	1.4	0.2	Iris-setosa

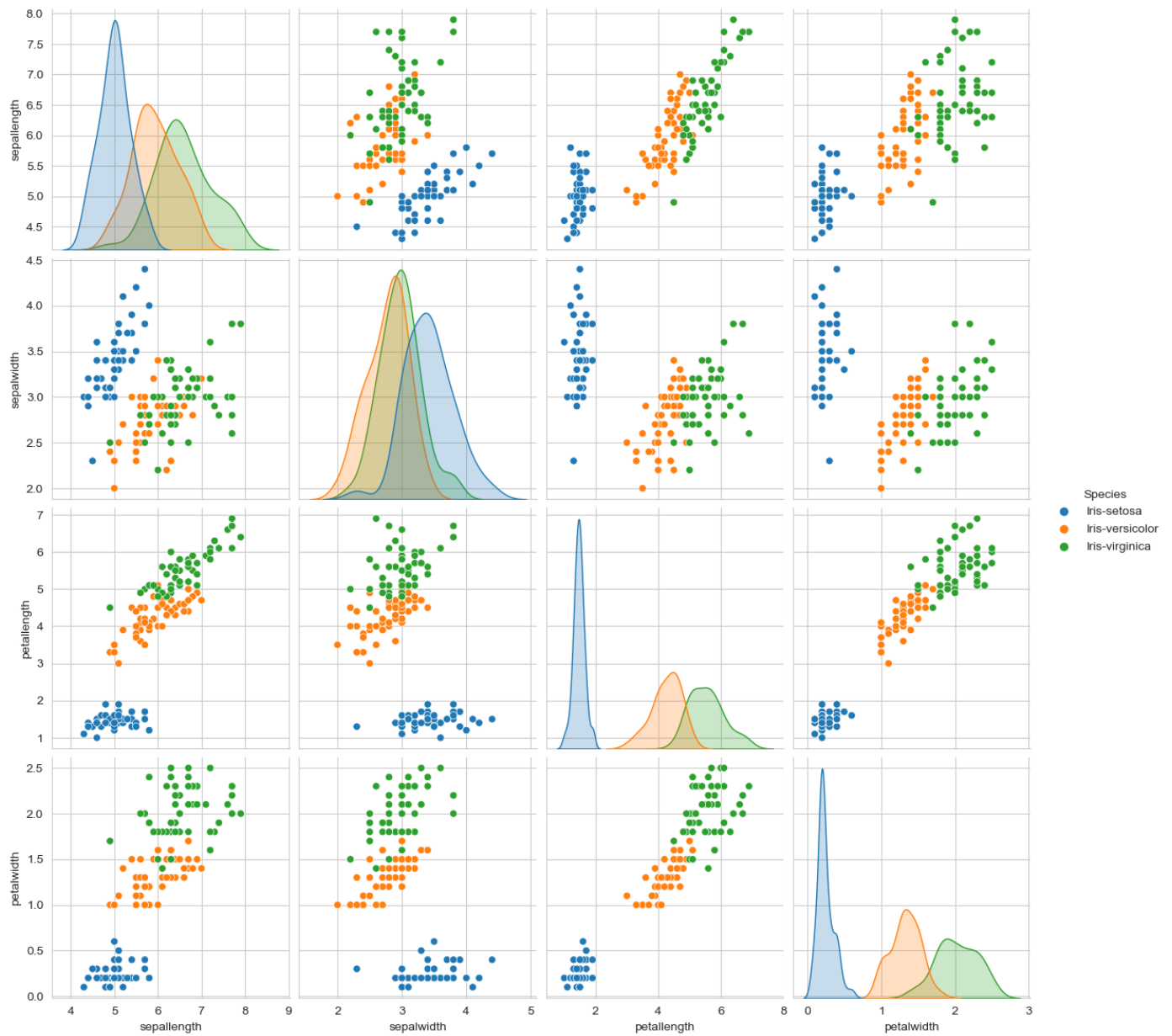
```
In [50]: data.head()
```

```
Out[50]:
```

	sepalength	sepalwidth	petallength	petalwidth	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

```
In [ ]: #Pair plot represents the relationship between our target and the variables
```

```
In [51]: sns.set_style("whitegrid")
sns.pairplot(data, hue= "Species", height =3);
plt.show()
```



In []: Histogram PDF

In [62]: data

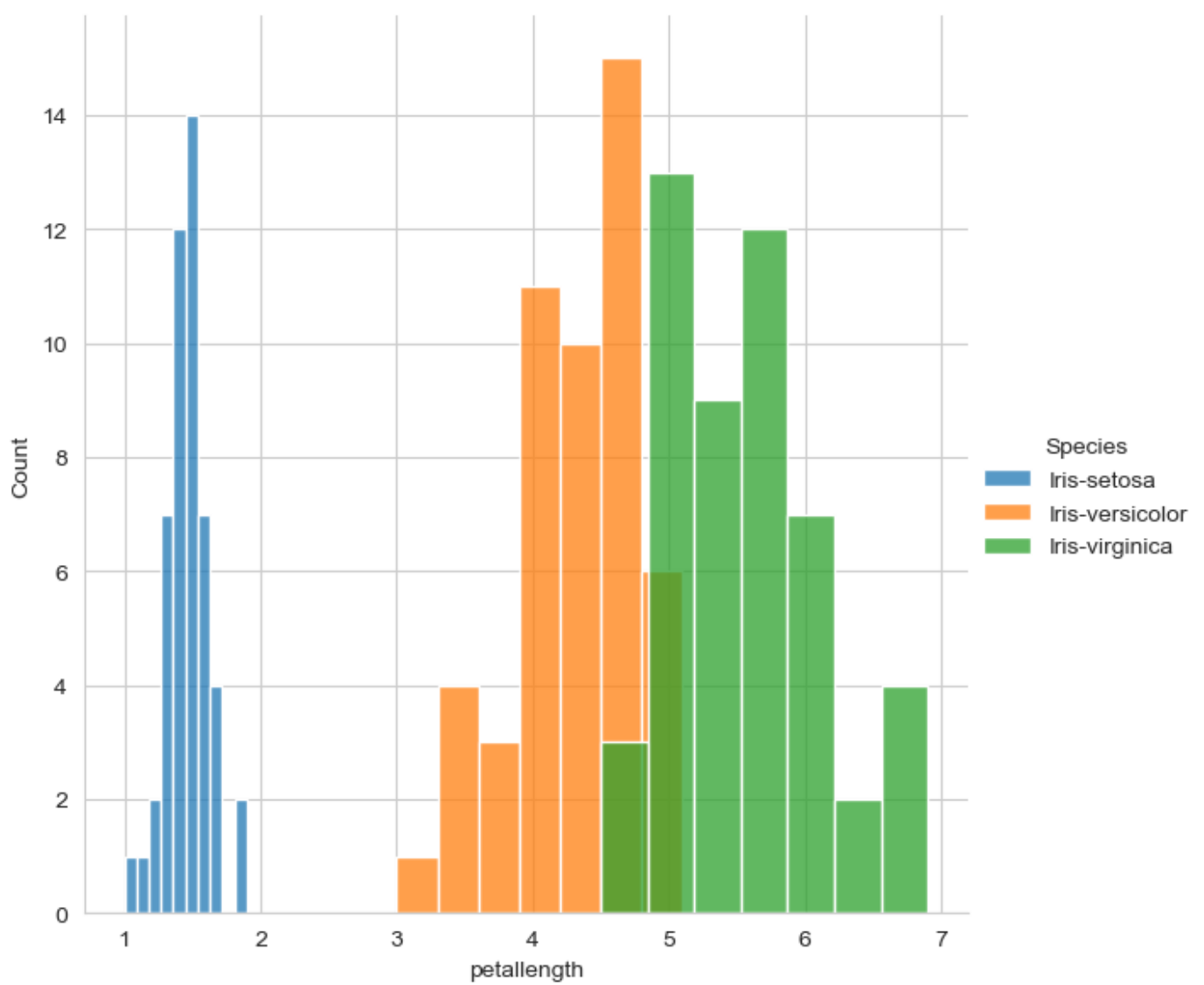
Out[62]:

	sepalength	sepalwidth	petallength	petalwidth	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

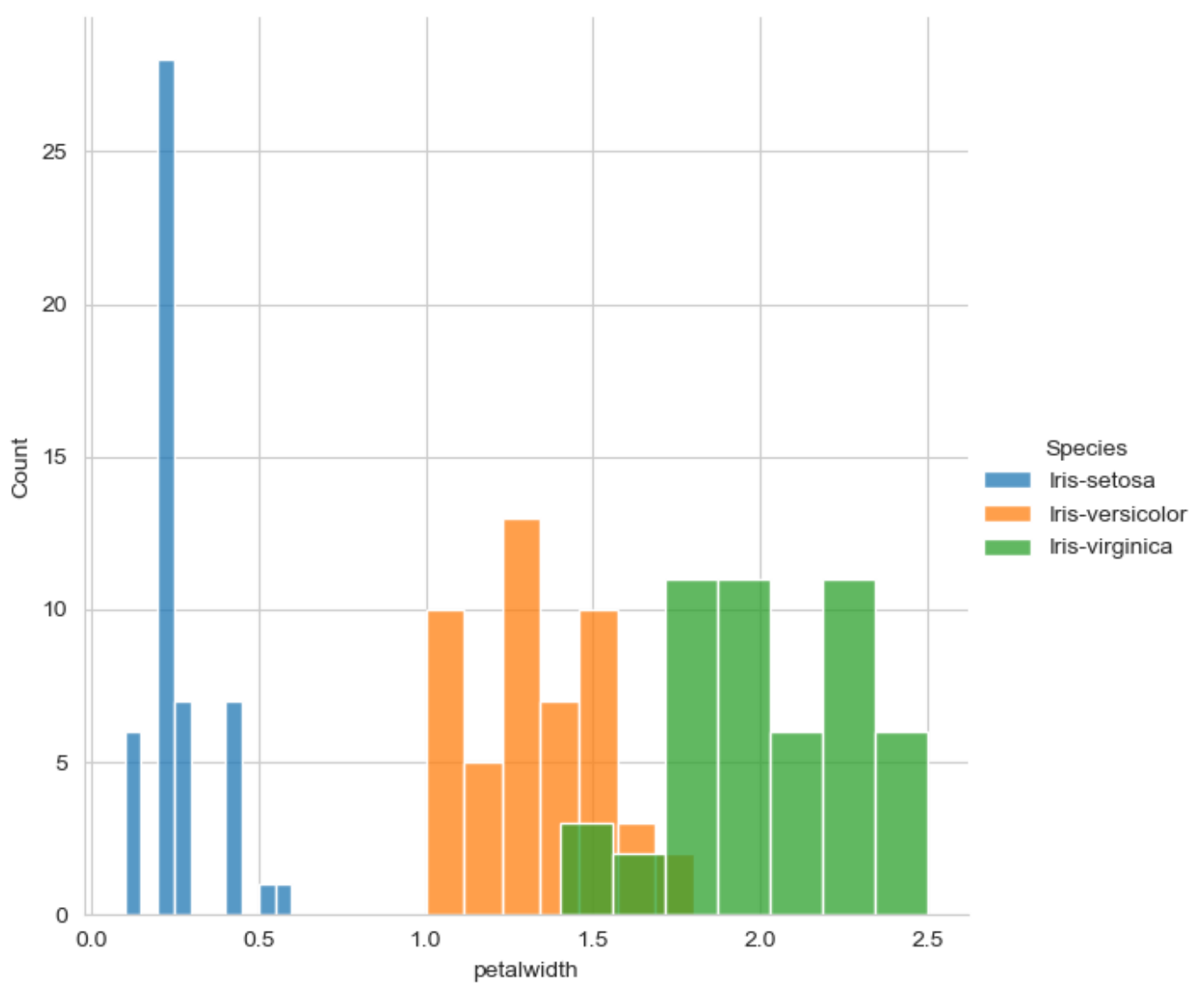
150 rows × 5 columns

```
In [105... import numpy as np
iris_setosa = data.loc[data["Species"]=="iris-setosa"]
iris_virginica = data.loc[data["Species"]=="Iris-virginica"];
iris_versicolor = data.loc[data["Species"]=="Iris-versicolor"]
```

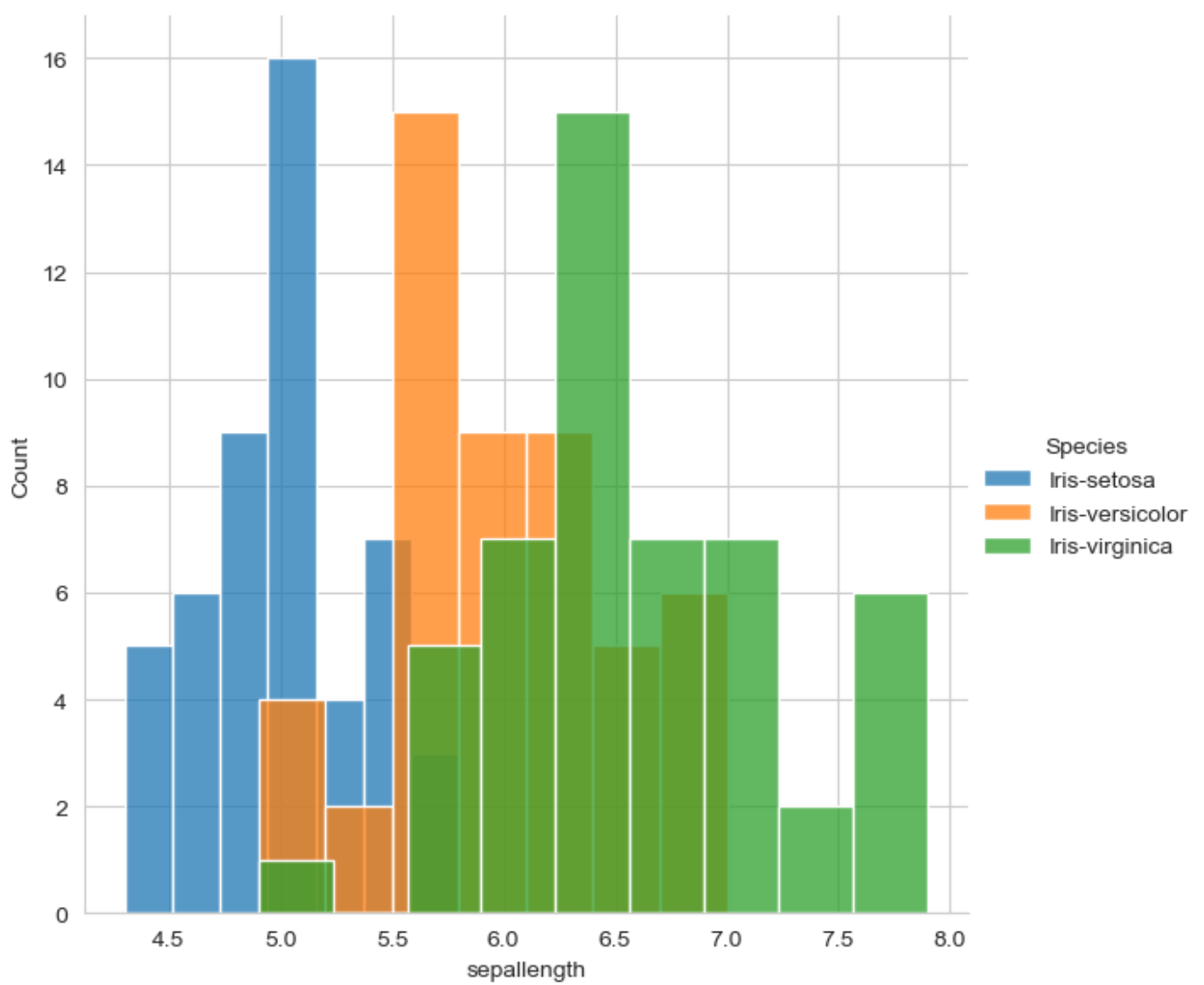
```
In [113... #Univariate Analysis
sns.FacetGrid(data, hue="Species", height=6)\
.map(sns.histplot, "petallength")\
.add_legend()
plt.show()
```



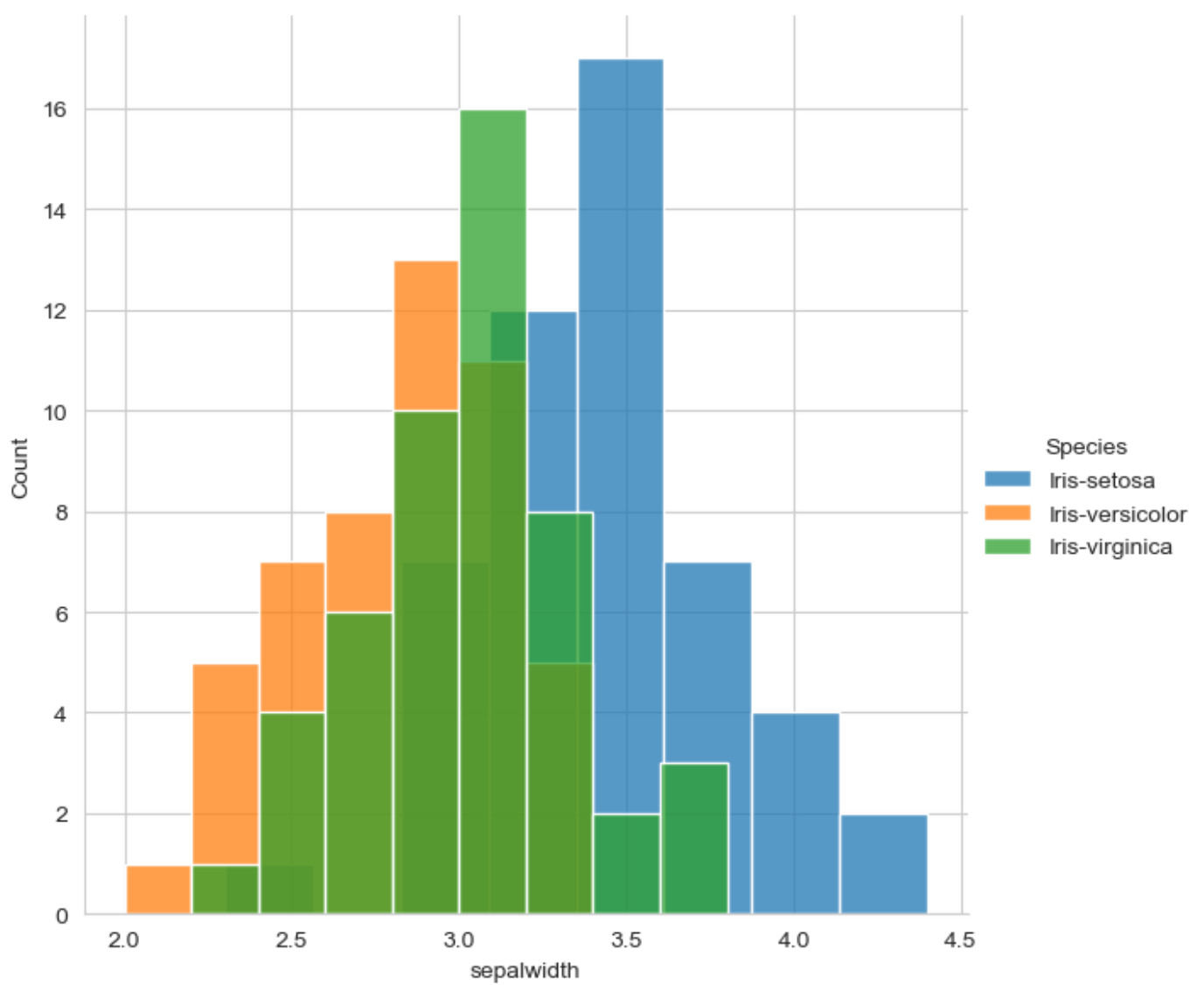
```
In [111... sns.FacetGrid(data, hue="Species", height=6)\
.map(sns.histplot, "petalwidth")\
.add_legend()\
plt.show()
```



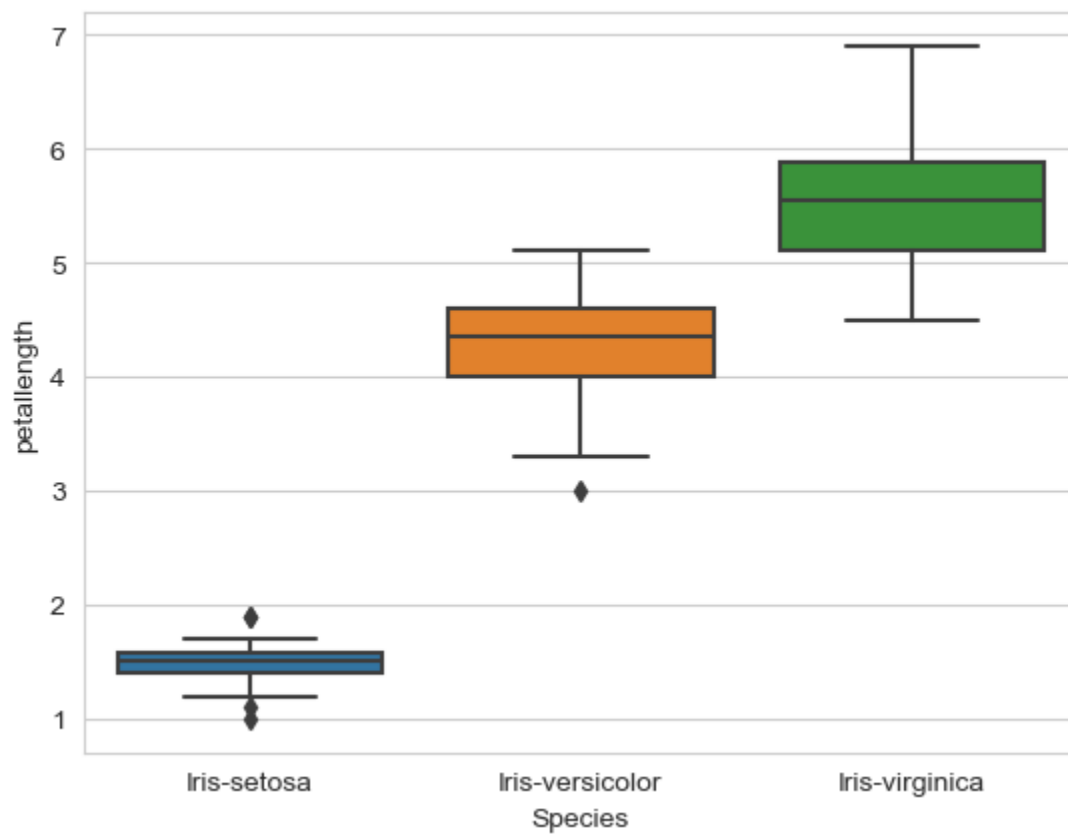
```
In [109... sns.FacetGrid(data, hue="Species", height=6)\  
            .map(sns.histplot, "sepalength")\  
            .add_legend()  
            plt.show()
```



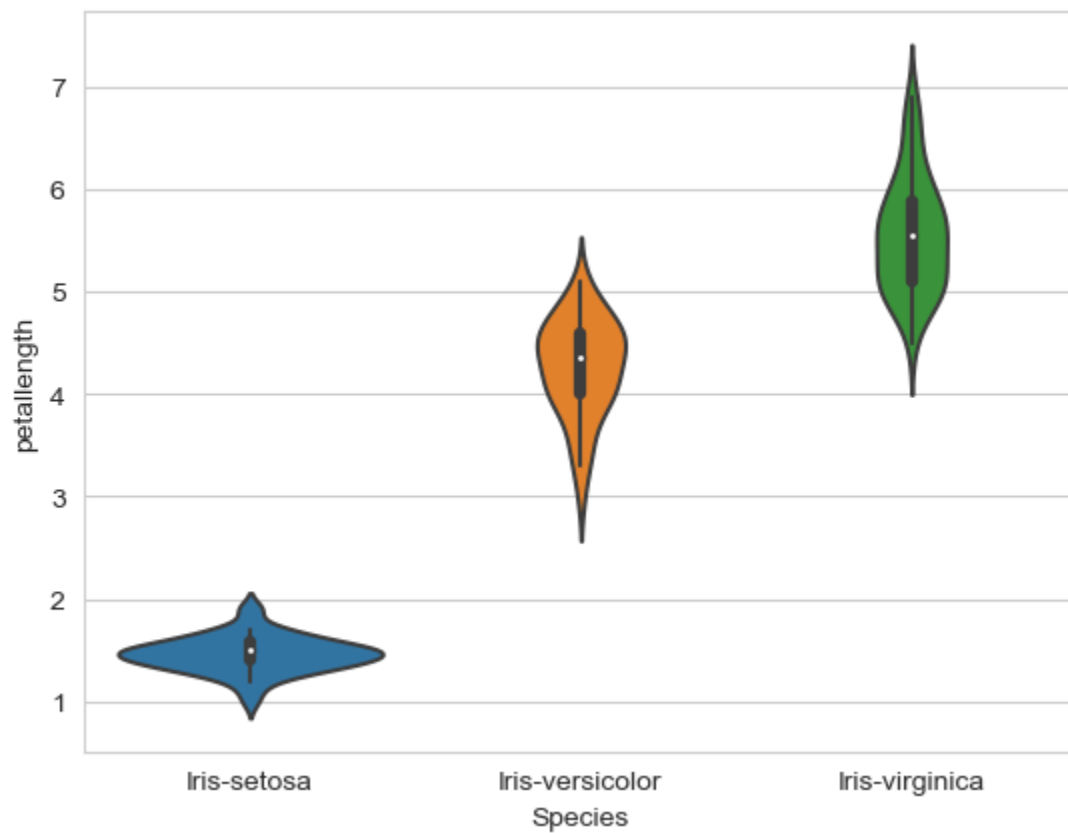
```
In [112... sns.FacetGrid(data, hue="Species", height=6)\  
            .map(sns.histplot, "sepalwidth")\  
            .add_legend()  
            plt.show()
```

```
In [92]: #Box Plot
#Box-plot can be visualized as a PDF on the side-ways
sns.boxplot(x= 'Species', y= 'petallength', data = data)
plt.show()
```



```
In [101... #Violen plot
sns.violinplot(x= "Species", y= "petal length", data = data, height=8)
plt.show()
```



In []: