

```
In [1]: import pandas as pd
import matplotlib.pyplot as plt
%matplotlib inline
import seaborn as sns
import warnings
warnings.filterwarnings('ignore')

from sklearn.linear_model import LogisticRegression as lr
from sklearn import metrics
from sklearn.metrics import classification_report
from sklearn.metrics import accuracy_score
```

```
In [2]: irisdf = pd.read_csv("C:/Users/HP/Downloads/iris.csv")
#first 5 dataset
irisdf.head()
```

```
Out[2]:
```

	<b>Id</b>	<b>SepalLengthCm</b>	<b>SepalWidthCm</b>	<b>PetalLengthCm</b>	<b>PetalWidthCm</b>	<b>Species</b>
<b>0</b>	1	5.1	3.5	1.4	0.2	Iris-setosa
<b>1</b>	2	4.9	3.0	1.4	0.2	Iris-setosa
<b>2</b>	3	4.7	3.2	1.3	0.2	Iris-setosa
<b>3</b>	4	4.6	3.1	1.5	0.2	Iris-setosa
<b>4</b>	5	5.0	3.6	1.4	0.2	Iris-setosa

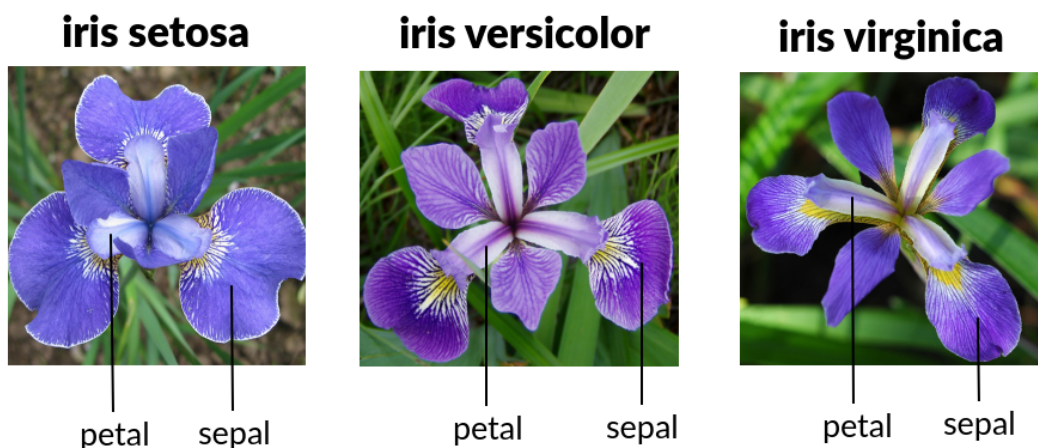
```
In [3]: #last 5 dataset
irisdf.tail()
```

```
Out[3]:
```

	<b>Id</b>	<b>SepalLengthCm</b>	<b>SepalWidthCm</b>	<b>PetalLengthCm</b>	<b>PetalWidthCm</b>	<b>Species</b>
<b>145</b>	146	6.7	3.0	5.2	2.3	Iris-virginica
<b>146</b>	147	6.3	2.5	5.0	1.9	Iris-virginica
<b>147</b>	148	6.5	3.0	5.2	2.0	Iris-virginica
<b>148</b>	149	6.2	3.4	5.4	2.3	Iris-virginica
<b>149</b>	150	5.9	3.0	5.1	1.8	Iris-virginica

```
In [4]: from IPython import display
display.Image("C:/Users/HP/Downloads/iris.png")
```

```
Out[4]:
```



```
In [5]: irisdf.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 6 columns):
 #   Column          Non-Null Count  Dtype  
---  -
 0   Id              150 non-null   int64  
 1   SepalLengthCm   150 non-null   float64
 2   SepalWidthCm    150 non-null   float64
 3   PetalLengthCm   150 non-null   float64
 4   PetalWidthCm    150 non-null   float64
 5   Species         150 non-null   object  
dtypes: float64(4), int64(1), object(1)
memory usage: 7.2+ KB
```

```
In [6]: irisdf.shape
```

```
Out[6]: (150, 6)
```

```
In [7]: irisdf.size
```

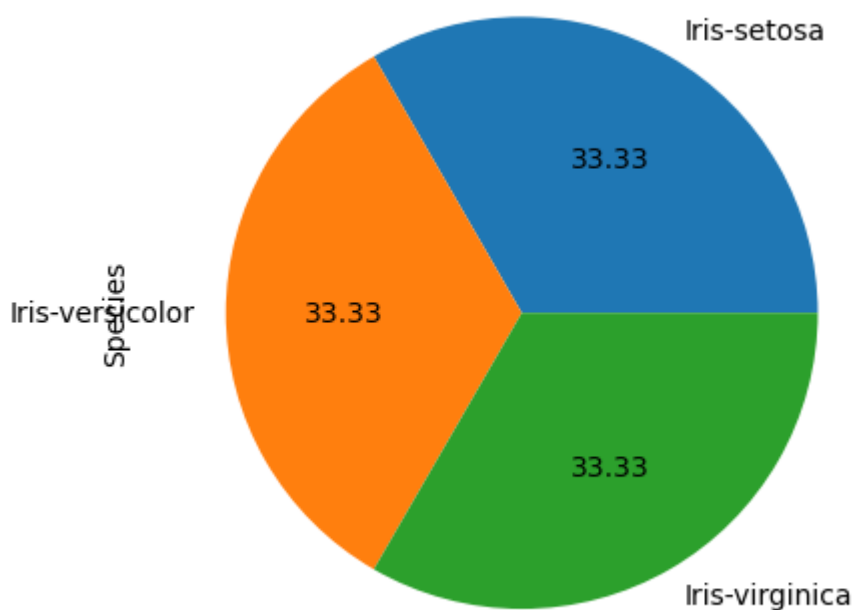
```
Out[7]: 900
```

```
In [8]: #check for values of each species whether its balanced or imbalanced
irisdf['Species'].value_counts()
```

```
Out[8]: Iris-setosa      50
Iris-versicolor    50
Iris-virginica     50
Name: Species, dtype: int64
```

```
In [9]: # Visualization in form of pie chart
irisdf['Species'].value_counts().plot(kind='pie', autopct='%.2f')
```

```
Out[9]: <AxesSubplot:ylabel='Species'>
```



```
In [10]: #Check if any null values is present in give dataset
irisdf.isnull().sum()
```

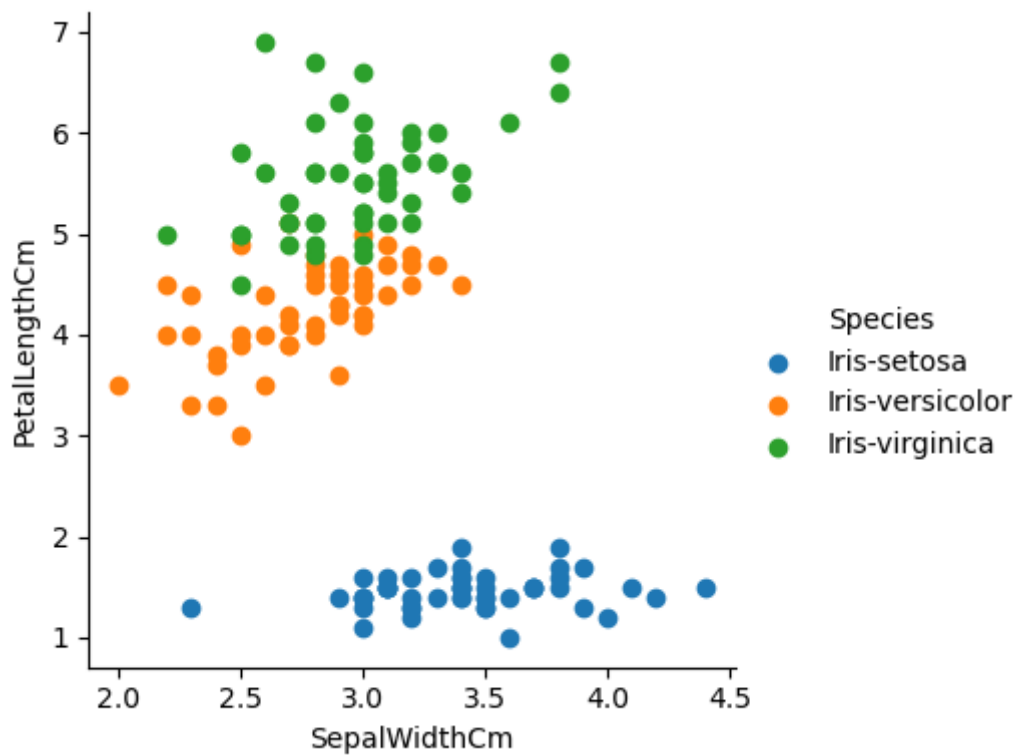
```
Out[10]: Id                0
SepalLengthCm            0
SepalWidthCm             0
PetalLengthCm            0
PetalWidthCm            0
Species                 0
dtype: int64
```

```
In [11]: #Check for duplicate values
duplicate_count = irisdf.duplicated().sum()
print(duplicate_count)
```

```
0
```

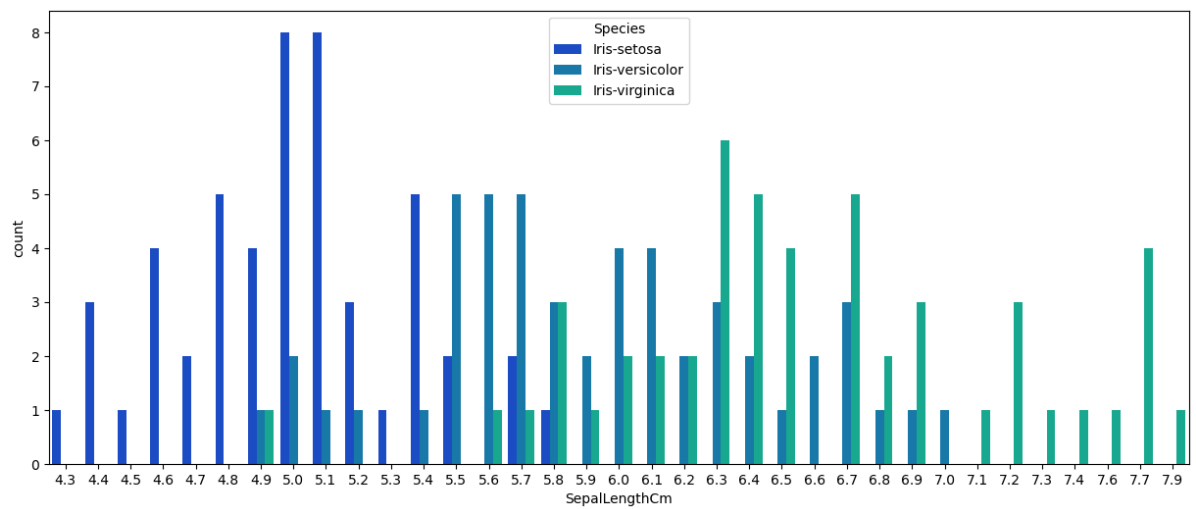
```
In [12]: sns.FacetGrid(irisdf,hue='Species',height=4).map(plt.scatter,"SepalWidthCm" , "Peta
```

```
Out[12]: <seaborn.axisgrid.FacetGrid at 0x23c093c2e20>
```

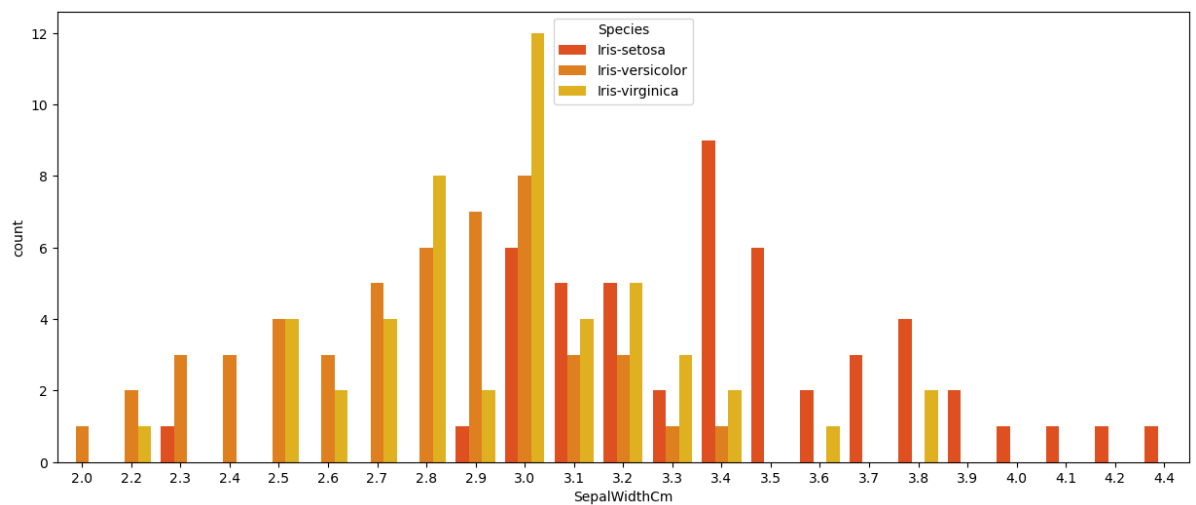


```
In [13]: plt.figure(figsize =(15,6))
sns.countplot(x='SepalLengthCm', data=irisdf, hue= irisdf['Species'], palette= 'w
```

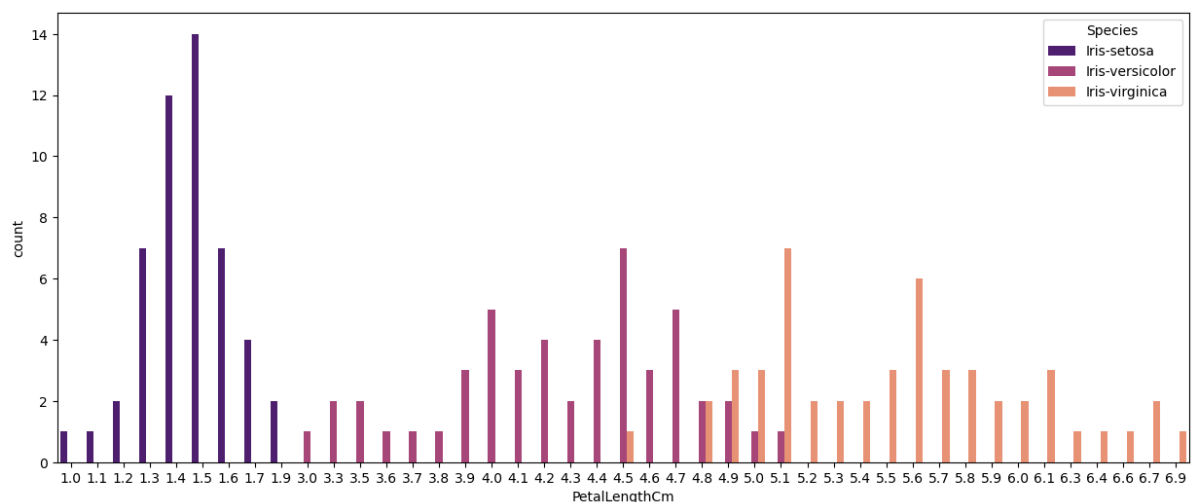
```
plt.show()
```



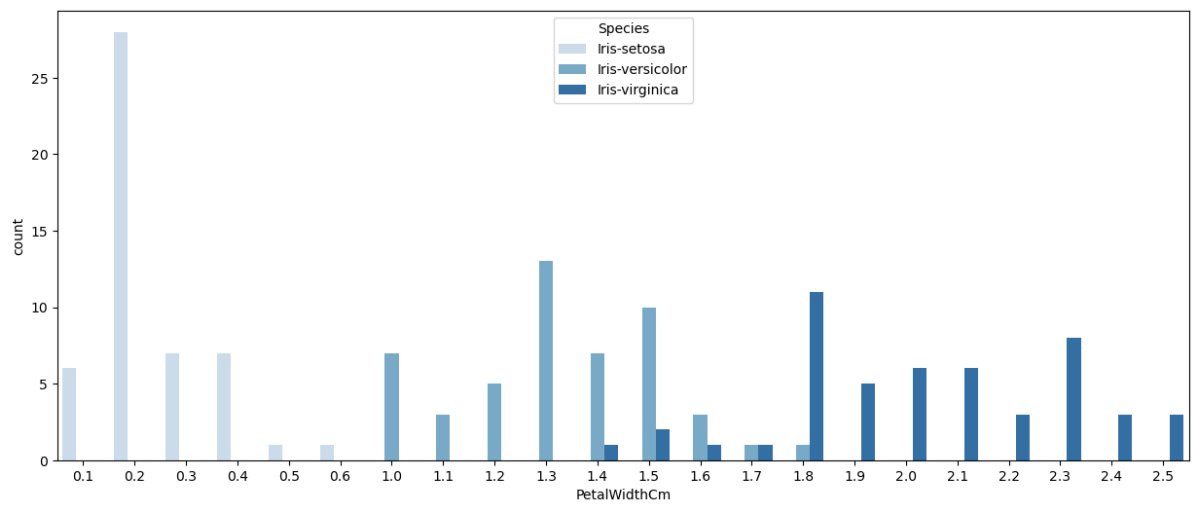
```
In [14]: plt.figure(figsize =(15,6))
sns.countplot(x='SepalWidthCm', data=irisdf, hue= irisdf['Species'], palette= 'autu
plt.show()
```



```
In [15]: plt.figure(figsize =(15,6))
sns.countplot(x='PetalLengthCm', data=irisdf, hue= irisdf['Species'], palette= 'mag
plt.show()
```



```
In [16]: plt.figure(figsize =(15,6))
sns.countplot(x='PetalWidthCm', data=irisdf, hue= irisdf['Species'], palette= 'Blue
plt.show()
```



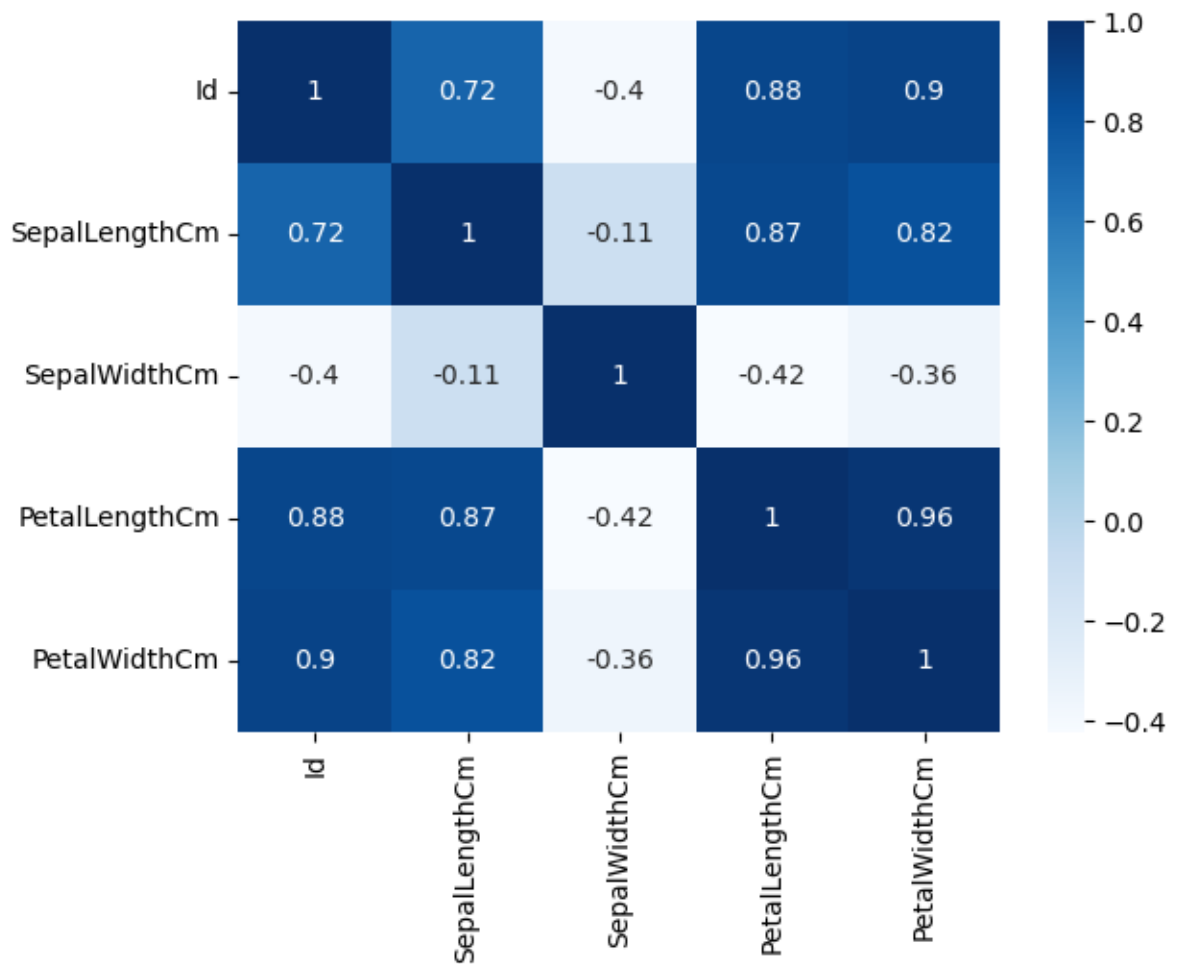
```
In [17]: irisdf.corr()
```

```
Out[17]:
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
Id	1.000000	0.716676	-0.397729	0.882747	0.899759
SepalLengthCm	0.716676	1.000000	-0.109369	0.871754	0.817954
SepalWidthCm	-0.397729	-0.109369	1.000000	-0.420516	-0.356544
PetalLengthCm	0.882747	0.871754	-0.420516	1.000000	0.962757
PetalWidthCm	0.899759	0.817954	-0.356544	0.962757	1.000000

```
In [18]: corr = irisdf.corr()
sns.heatmap(corr,annot=True,cmap='Blues')
```

```
Out[18]: <AxesSubplot:>
```



```
In [19]: flower_mapping= {'Iris-setosa': 0, 'Iris-versicolor' : 1, 'Iris-virginica' :2}
irisdf['Species'] = irisdf ['Species'].map (flower_mapping)
```

```
In [20]: irisdf.head()
```

```
Out[20]:
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	0
1	2	4.9	3.0	1.4	0.2	0
2	3	4.7	3.2	1.3	0.2	0
3	4	4.6	3.1	1.5	0.2	0
4	5	5.0	3.6	1.4	0.2	0

```
In [21]: irisdf.tail()
```

```
Out[21]:
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
145	146	6.7	3.0	5.2	2.3	2
146	147	6.3	2.5	5.0	1.9	2
147	148	6.5	3.0	5.2	2.0	2
148	149	6.2	3.4	5.4	2.3	2
149	150	5.9	3.0	5.1	1.8	2

```
In [22]: x=irisdf[['SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm','PetalWidthCm']].values  
y=irisdf[['Species']].values
```

```
In [23]: from sklearn.linear_model import LinearRegression  
model= LinearRegression()
```

```
In [24]: model.fit(x,y)
```

```
Out[24]: LinearRegression()
```