

Iris Flower Classification

Importing Libraries

```
In [10]: ▶ import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
%matplotlib inline
import seaborn as sns

## Supress warnings
import warnings
warnings.filterwarnings("ignore")
```

```
In [11]: ▶ iris_df = pd.read_csv("iris.csv")
print("the data has been successfully load.")

the data has been successfully load.
```

```
In [12]: ▶ iris_df
```

```
Out[12]:
```

	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
...
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
149	150	5.9	3.0	5.1	1.8	Iris-virginica

150 rows × 6 columns

Data Exploration

```
In [13]: ▶ #check shape of data
iris_df.shape
```

```
Out[13]: (150, 6)
```

```
In [14]: ▶ # check basic information of data
iris_df.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 [15]: ▶ # check statistical summary of data
iris_df.describe()
```

```
Out[15]:
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
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 [16]: ▶ #Check null values
iris_df.isnull().sum()
```

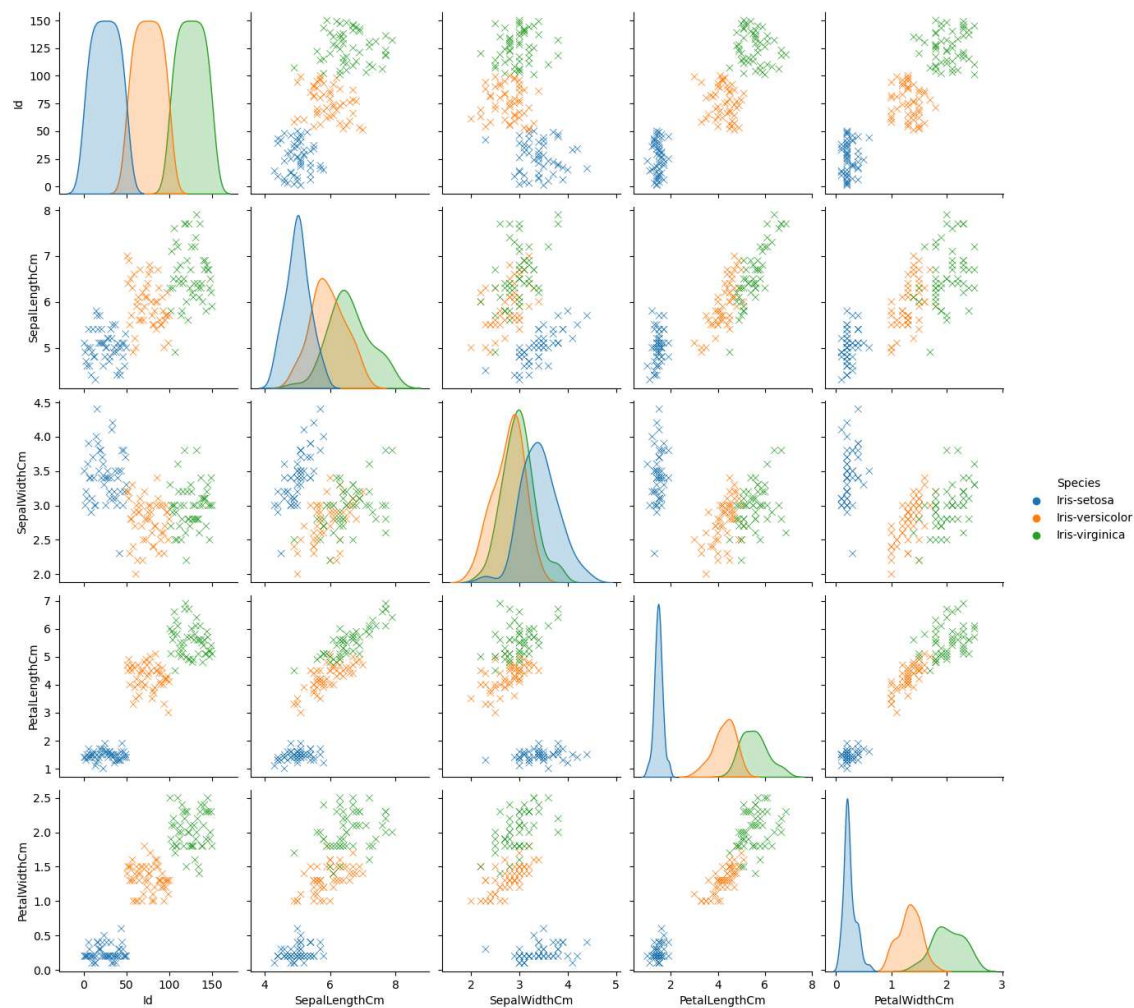
```
Out[16]: Id               0
SepalLengthCm           0
SepalWidthCm            0
PetalLengthCm           0
PetalWidthCm            0
Species                 0
dtype: int64
```

```
In [18]: ▶ print("unique number of values in dataset Species:",iris_df["Species"].nunique())
print("Unique Species in iris dataset:",iris_df["Species"].unique())
```

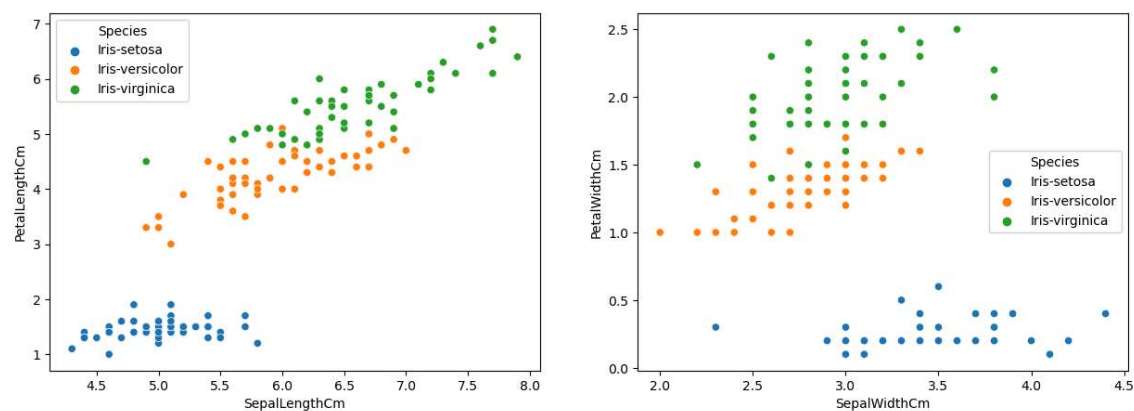
```
unique number of values in dataset Species: 3
Unique Species in iris dataset: ['Iris-setosa' 'Iris-versicolor' 'Iris-virginica']
```

Data Visualization

```
In [19]: ▶ sns.pairplot(iris_df, hue = "Species", markers = "x")  
plt.show()
```



```
In [20]: ▶ plt.figure(figsize=(15,5))  
plt.subplot(1,2,1)  
sns.scatterplot(x='SepalLengthCm',y='PetalLengthCm',data=iris_df,hue='Species')  
plt.subplot(1,2,2)  
sns.scatterplot(x='SepalWidthCm',y='PetalWidthCm',data=iris_df,hue='Species')  
plt.show()
```



```
In [25]: ► # Select only the numeric columns from the DataFrame
numeric_columns = iris_df.select_dtypes(include=[float, int])

# Calculate the correlation matrix
correlation_matrix = numeric_columns.corr()

# Display the correlation matrix
print(correlation_matrix)
```

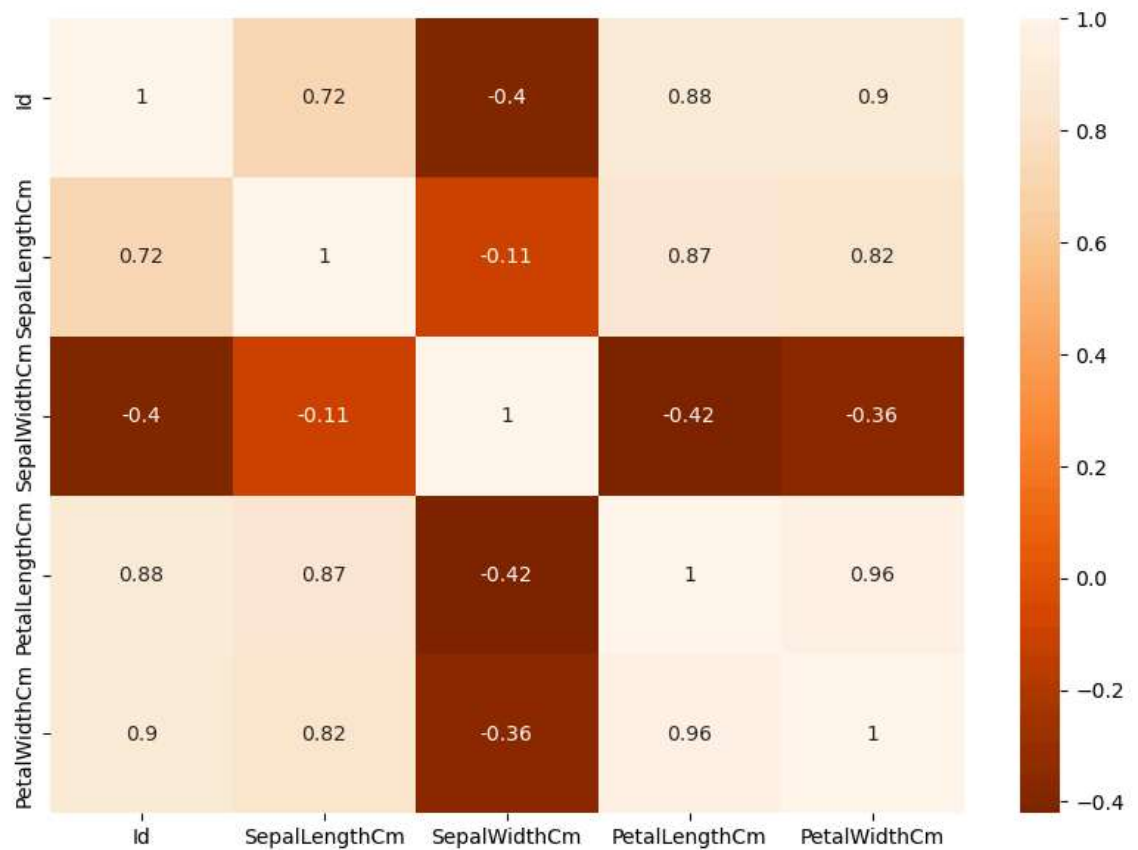
	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	\
Id	1.000000	0.716676	-0.397729	0.882747	
SepalLengthCm	0.716676	1.000000	-0.109369	0.871754	
SepalWidthCm	-0.397729	-0.109369	1.000000	-0.420516	
PetalLengthCm	0.882747	0.871754	-0.420516	1.000000	
PetalWidthCm	0.899759	0.817954	-0.356544	0.962757	

	PetalWidthCm
Id	0.899759
SepalLengthCm	0.817954
SepalWidthCm	-0.356544
PetalLengthCm	0.962757
PetalWidthCm	1.000000

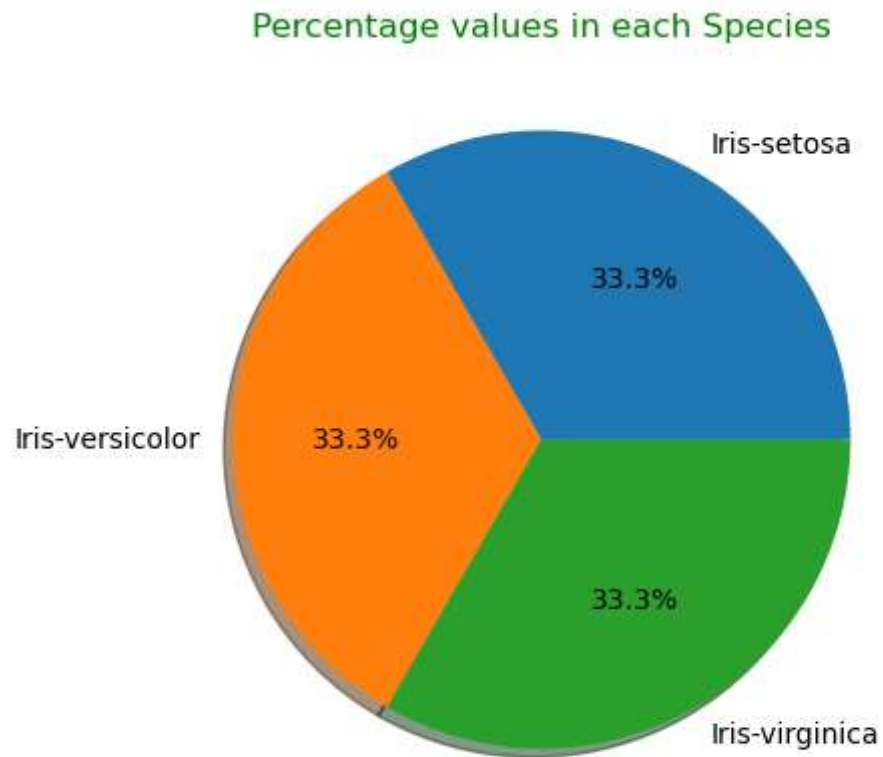
```
In [27]: # Select only the numeric columns from the DataFrame
numeric_columns = iris_df.select_dtypes(include=[float, int])

# Calculate the correlation matrix for the numeric columns
correlation_matrix = numeric_columns.corr()

# Use a heatmap to visualize the correlation matrix
plt.figure(figsize=(10, 7))
sns.heatmap(correlation_matrix, annot=True, cmap="Oranges_r")
plt.show()
```



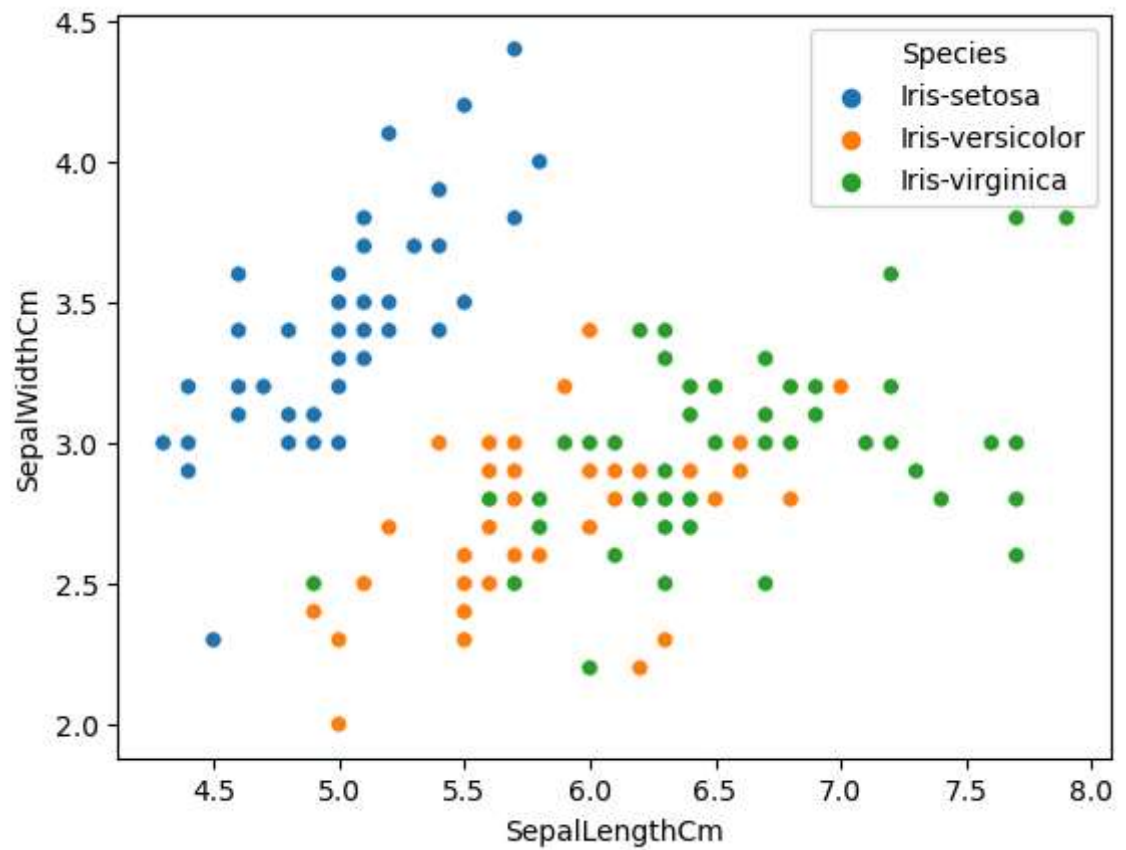
```
In [28]: ▶ # Check value counts
iris_df["Species"].value_counts().plot(kind="pie", autopct = "%1.1f%%", shadow=True)
plt.title("Percentage values in each Species", fontsize = 12, c = "g")
plt.ylabel("", fontsize=10, c="r")
plt.show()
```



Scatterplot for Sepal Length and Sepal Width

```
In [31]: ▶ # Create a scatter plot with SepalLengthCm on the x-axis and SepalWidthCm on the y-axis
sns.scatterplot(x=iris_df["SepalLengthCm"], y=iris_df["SepalWidthCm"], hue=iris_df["Species"])

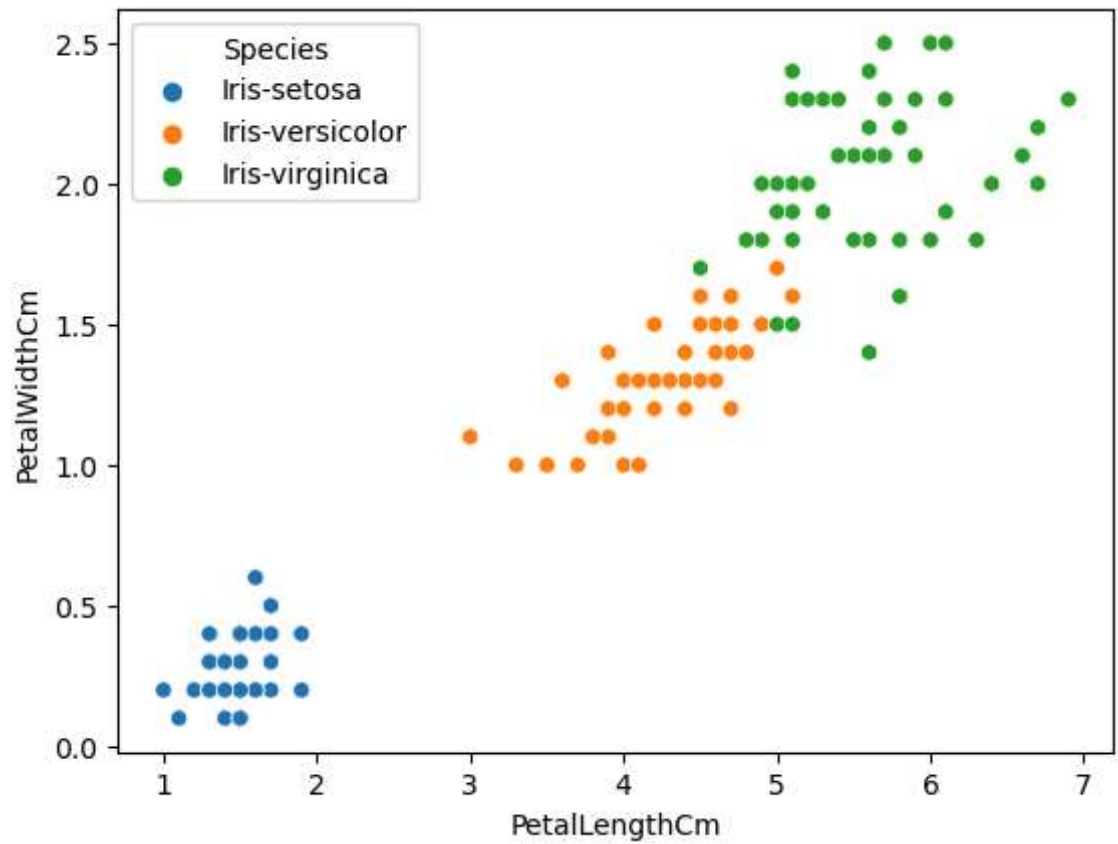
# Show the plot
plt.show()
```



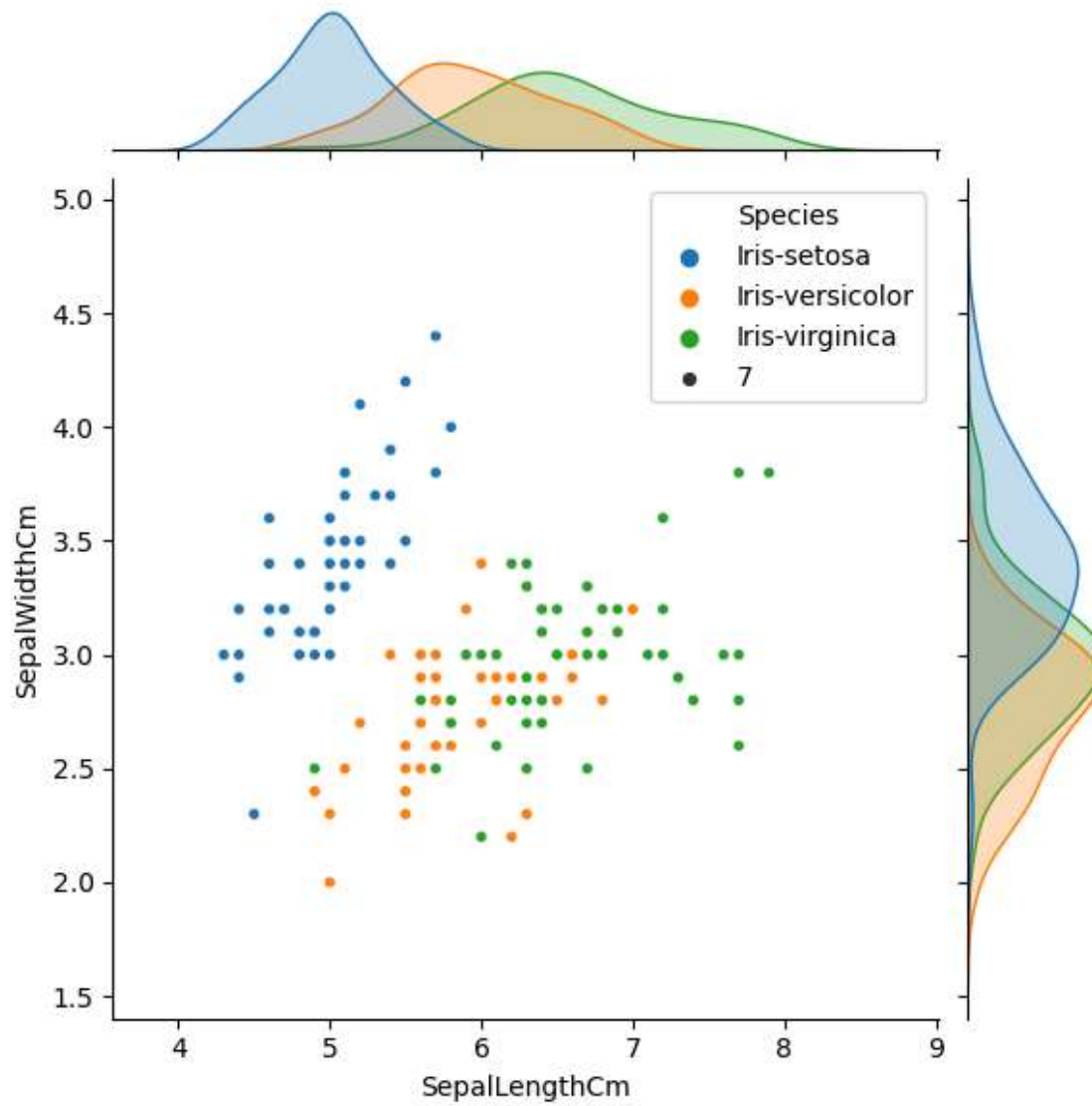
Scatterplot for Petal Length and Petal Width

```
In [32]: ▶ # Create a scatter plot with PetalLengthCm on the x-axis and PetalWidthCm on the y-axis
sns.scatterplot(x=iris_df["PetalLengthCm"], y=iris_df["PetalWidthCm"], hue=iris_df["Species"])

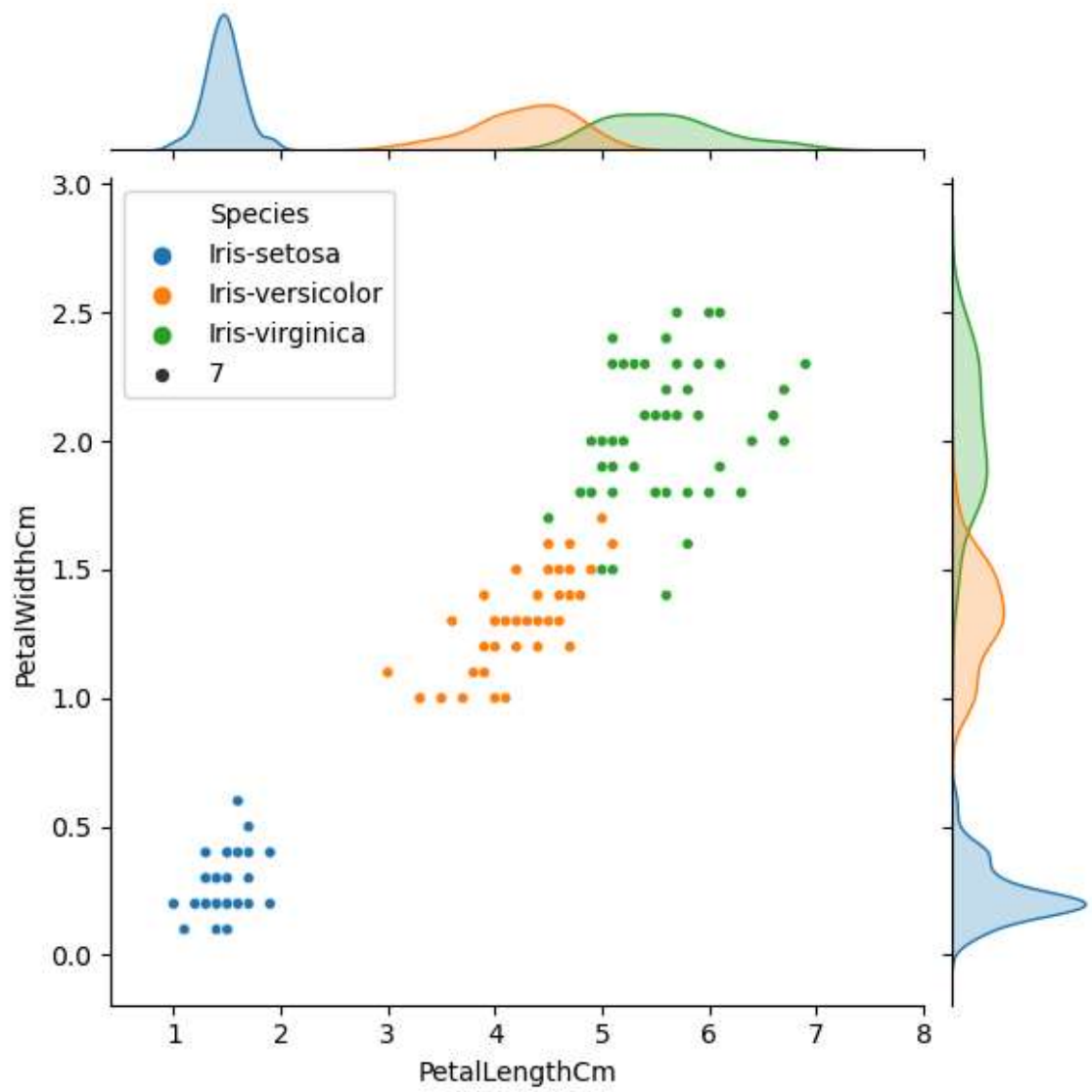
# Show the plot
plt.show()
```




```
In [33]: ▶ sns.jointplot(data = iris_df , x = "SepalLengthCm", y = "SepalWidthCm" , kind = "scatter",
plt.show())
```



```
In [34]: ▶ sns.jointplot(data = iris_df , x = "PetalLengthCm", y = "PetalWidthCm" , size=10,
plt.show())
```



```

In [35]: ▶ plt.figure(figsize = (15,5))
plt.subplot(2,2,1)
sns.barplot(x = "Species",y = "SepallLengthCm", data=iris_df, palette=("Spec
plt.title("Bar plot SepallLengthCm Vs Species")

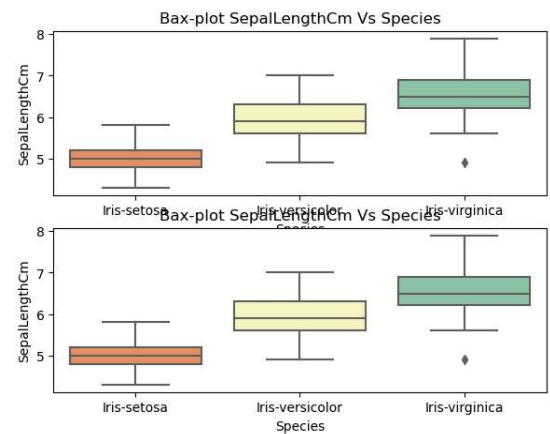
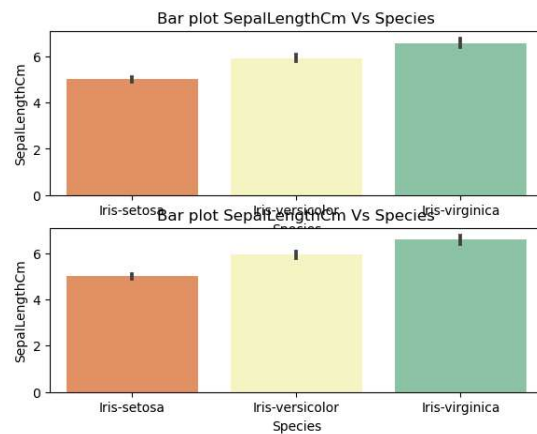
plt.subplot(2,2,2)
sns.boxplot(x = "Species",y = "SepallLengthCm", data=iris_df, palette=("Spec
plt.title("Bax-plot SepallLengthCm Vs Species")

plt.subplot(2,2,3)
sns.barplot(x = "Species",y = "SepallLengthCm", data=iris_df, palette=("Spec
plt.title("Bar plot SepallLengthCm Vs Species")

plt.subplot(2,2,4)
sns.boxplot(x = "Species",y = "SepallLengthCm", data=iris_df, palette=("Spec
plt.title("Bax-plot SepallLengthCm Vs Species")

plt.show()

```



```

In [37]: ▶ plt.figure(figsize = (15,5))
plt.subplot(2,2,1)
sns.barplot(x = "Species",y = "SepallLengthCm", data=iris_df, palette=("cool")
plt.title("Bar plot SepallLengthCm Vs Species")

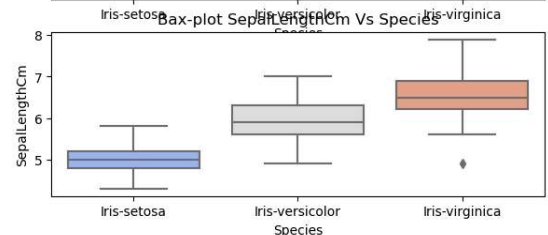
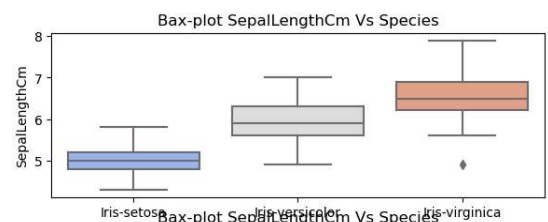
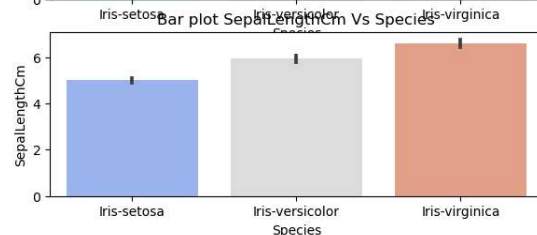
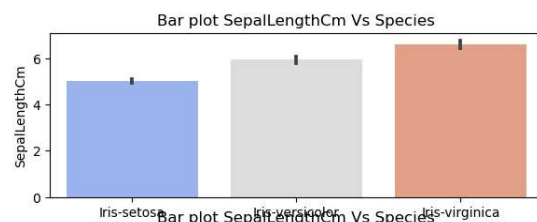
plt.subplot(2,2,2)
sns.boxplot(x = "Species",y = "SepallLengthCm", data=iris_df, palette=("cool")
plt.title("Bax-plot SepallLengthCm Vs Species")

plt.subplot(2,2,3)
sns.barplot(x = "Species",y = "SepallLengthCm", data=iris_df, palette=("cool")
plt.title("Bar plot SepallLengthCm Vs Species")

plt.subplot(2,2,4)
sns.boxplot(x = "Species",y = "SepallLengthCm", data=iris_df, palette=("cool")
plt.title("Bax-plot SepallLengthCm Vs Species")

plt.show()

```



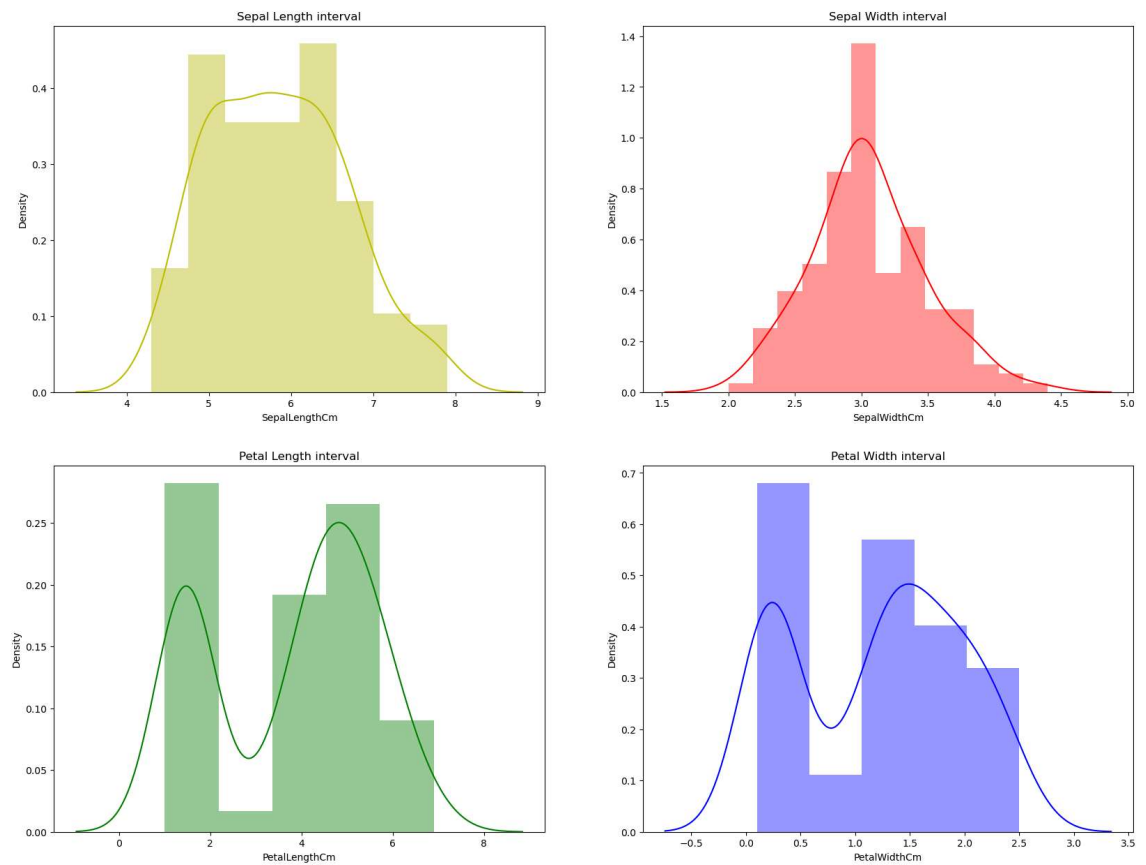
```
In [38]: ▶ plt.figure(figsize=(20,15))
plt.subplot(2,2,1)
sns.distplot(iris_df["SepalLengthCm"],color="y").set_title("Sepal Length interval")

plt.subplot(2,2,2)
sns.distplot(iris_df["SepalWidthCm"],color="r").set_title("Sepal Width interval")

plt.subplot(2,2,3)
sns.distplot(iris_df["PetalLengthCm"],color="g").set_title("Petal Length interval")

plt.subplot(2,2,4)
sns.distplot(iris_df["PetalWidthCm"],color="b").set_title("Petal Width interval")

plt.show()
```



Data Cleaning

```
In [40]: # Change Categorical Data into numerical value
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
iris_df["Species"] = le.fit_transform(iris_df["Species"])
iris_df.head()
```

```
Out[40]:
```

	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 [41]: iris_df['Species'].unique()
```

```
Out[41]: array([0, 1, 2])
```

```
In [42]: X = iris_df.iloc[:, [0,1,2,3]]
X.head()
```

```
Out[42]:
```

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

```
In [43]: y = iris_df.iloc[:, - 1]
y.head()
```

```
Out[43]: 0    0
1    0
2    0
3    0
4    0
Name: Species, dtype: int32
```

```
In [44]: print(X.shape)
print(y.shape)
```

```
(150, 4)
(150,)
```

Model Building

Split data into Training and Testing Set

```
In [45]: 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,rand
```

```
In [46]: print(X_train.shape)
print(X_test.shape)
print(y_train.shape)
print(y_test.shape)
```

```
(120, 4)
(30, 4)
(120,)
(30,)
```

Logistic Regression

```
In [47]: from sklearn.linear_model import LogisticRegression
lr= LogisticRegression()
lr.fit(X_train, y_train)
print("Logistic regression successfully implemented")
y_pred = lr.predict(X_test)
#Confusion Matrix
from sklearn.metrics import confusion_matrix, accuracy_score, classification_report
cm = confusion_matrix(y_test, y_pred)
print("Confusion Matrix:-")
print(cm)
accuracy = accuracy_score(y_test, y_pred)
print("accuracy is:-",accuracy*100)
print("Classification Report:-")
print(classification_report(y_test,y_pred))
```

Logistic regression successfully implemented

Confusion Matrix:-

```
[[11  0  0]
 [ 0 13  0]
 [ 0  0  6]]
```

accuracy is:- 100.0

Classification Report:-

	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	1.00	1.00	1.00	13
2	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

Random Forest Classifier

```
In [48]: ▶ from sklearn.ensemble import RandomForestClassifier
rfc = RandomForestClassifier()

rfc.fit(X_train, y_train)
print("Random Forest Classifier successfully Implimented")

y_pred = rfc.predict(X_test)

#confusion matrix
cm = confusion_matrix(y_test, y_pred)
print("Confusion Matrix:- ")
print(cm)

#accuracy test
accuracy = accuracy_score(y_test,y_pred)

print("accuracy:- ", accuracy*100)

print("Classification Report:-")
print( classification_report(y_test, y_pred))
```

Rndom Forest Classifier successfully Implimented

Confusion Matrix:-

```
[[11  0  0]
 [ 0 13  0]
 [ 0  0  6]]
```

accuracy:- 100.0

Classification Report:-

	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	1.00	1.00	1.00	13
2	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

Decision Tree

In [49]: `from sklearn.tree import DecisionTreeClassifier`

```
dtree = DecisionTreeClassifier()

dtree.fit(X_train, y_train)
print("Decision Tree Algorithm is successfully implimented.")

y_pred = dtree.predict(X_test)

#confusion matrix
cm = confusion_matrix(y_test, y_pred)
print("Confusion Matrix:- ")
print(cm)

#accuracy test
accuracy = accuracy_score(y_test,y_pred)
print("accuracy:- ", accuracy*100)

print("Classification Report:-")
print( classification_report(y_test, y_pred))
```

Decision Tree Algorithm is successfully implimented.

Confusion Matrix:-

```
[[11  0  0]
 [ 0 13  0]
 [ 0  1  5]]
```

accuracy:- 96.66666666666667

Classification Report:-

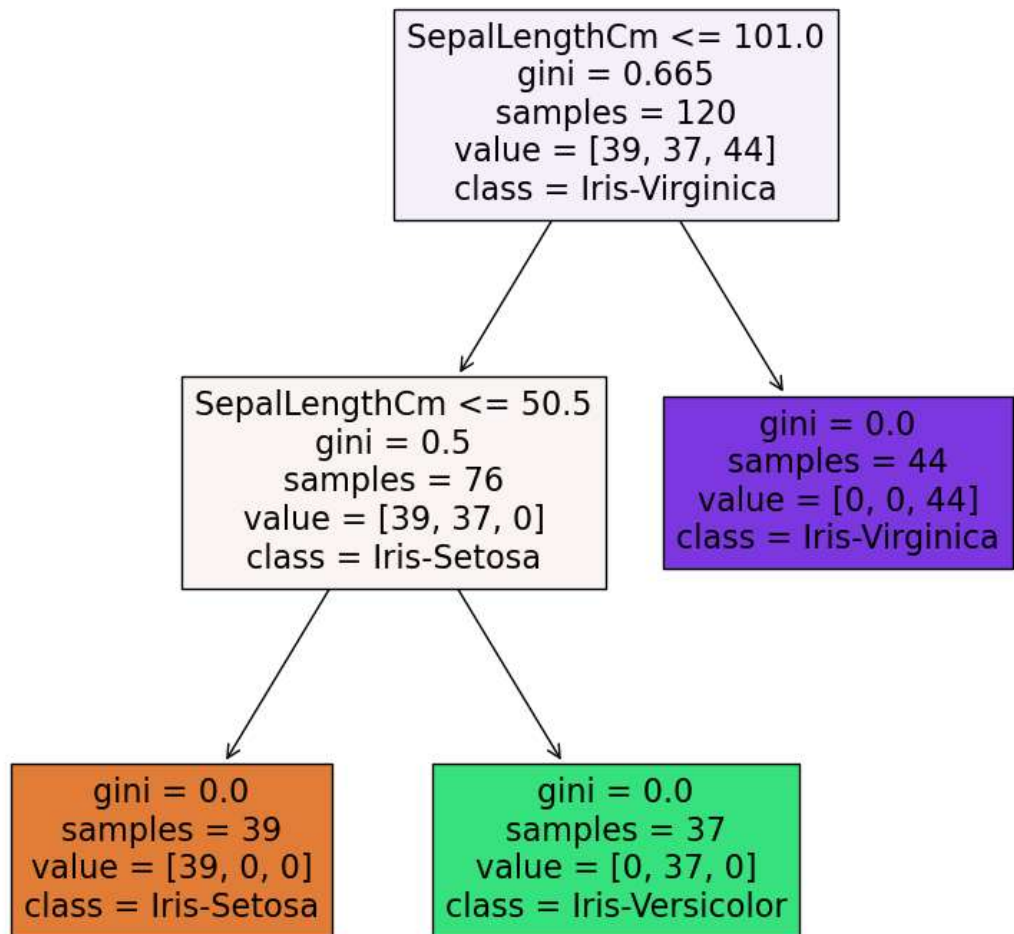
	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	0.93	1.00	0.96	13
2	1.00	0.83	0.91	6
accuracy			0.97	30
macro avg	0.98	0.94	0.96	30
weighted avg	0.97	0.97	0.97	30

In [50]: `from sklearn.tree import plot_tree`

In [51]: `# for visualziing the Decision Tree`

```
feature = ['SepalLengthCm','SepalWidthCm','PetalLengthCm','PetalWidthCm']
classes = ['Iris-Setosa','Iris-Versicolor','Iris-Virginica']
```

```
In [52]: ▶ plt.figure(figsize=(10,10))  
plot_tree(dtree, feature_names = feature, class_names = classes, filled = 1
```



Support Vector Machine

```
In [53]:  from sklearn.svm import SVC
          svc= SVC()

          svc.fit(X_train, y_train)
          print("Support vector classifier is successfully implemented")

          y_pred = svc.predict(X_test)

          #confusion matrix
          cm = confusion_matrix(y_test, y_pred)
          print("Confusion Matrix:- ")
          print(cm)

          #accuracy test
          accuracy = accuracy_score(y_test,y_pred)
          print("accuracy:- ", accuracy*100)

          print("Classification Report:-")

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

```
Support vector classifier is successfully implemented
Confusion Matrix:-
[[11  0  0]
 [ 0 13  0]
 [ 0  0  6]]
accuracy:- 100.0
Classification Report:-
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	1.00	1.00	1.00	13
2	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

K - NN Classifier

```
In [54]: ► from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n_neighbors= 7)
knn.fit(X_train, y_train)

print("K-Nearest Neighbors classifier is successfully implemented")

y_pred = knn.predict(X_test)

#confusion matrix
cm = confusion_matrix(y_test, y_pred)
print("Confusion Matrix:- ")
print(cm)

#accuracy test
accuracy = accuracy_score(y_test,y_pred)

print("accuracy:- ", accuracy*100)

print("Classification Report:-")
print( classification_report(y_test, y_pred))
```

K-Nearest Neighbors classifier is successfully implemented

Confusion Matrix:-

```
[[11  0  0]
 [ 0 13  0]
 [ 0  0  6]]
```

accuracy:- 100.0

Classification Report:-

	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	1.00	1.00	1.00	13
2	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

Naive Bayes

```
In [55]: from sklearn.naive_bayes import GaussianNB
gnb = GaussianNB()
gnb.fit(X_train, y_train)
print("Naive Bayes is successfully implemented")

y_pred = gnb.predict(X_test)

cm = confusion_matrix(y_pred, y_test)
print("Confusion Matrix:- ")
print(cm)

# Accuracy test
accuracy = accuracy_score(y_test, y_pred)
print("accuracy:- ", accuracy*100)

print("Classification Report:-")
print(classification_report(y_test, y_pred))
```


```
Naive Bayes is successfully implemented
Confusion Matrix:-
[[11  0  0]
 [ 0 13  0]
 [ 0  0  6]]
accuracy:- 100.0
Classification Report:-
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	1.00	1.00	1.00	13
2	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

Result

1. Accuracy of Logistic Regression :- 100%
2. Accuracy of Random Forest Classifier:-100%
3. Accuracy of Decision Tree :- 96.66%
4. Accuracy of Support Vector Machine :- 100%
5. Accuracy of K-NN Classifier :- 100%
6. Accuracy of Naive Bayes :- 100%

Test Model

In [58]:  input_data=(4.9,3.0,1.4,0.2)

```
#changing the input data to a numpy array
```

```
input_data_as_ndarray = np.asarray(input_data)
```

```
#reshape the data as we are predicting the label for only the instance
```

```
input_data_reshaped = input_data_as_ndarray.reshape(1,-1)
```

```
prediction = dtree.predict(input_data_reshaped)
```

```
print("The category is",prediction)
```

The category is [0]