

task1

November 5, 2023

```
[ ]: !pip install matplotlib
```

```
Requirement already satisfied: matplotlib in /usr/local/lib/python3.10/dist-packages (3.7.1)
Requirement already satisfied: contourpy>=1.0.1 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (1.1.1)
Requirement already satisfied: cycler>=0.10 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (0.12.1)
Requirement already satisfied: fonttools>=4.22.0 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (4.43.1)
Requirement already satisfied: kiwisolver>=1.0.1 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (1.4.5)
Requirement already satisfied: numpy>=1.20 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (1.23.5)
Requirement already satisfied: packaging>=20.0 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (23.2)
Requirement already satisfied: pillow>=6.2.0 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (9.4.0)
Requirement already satisfied: pyparsing>=2.3.1 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (3.1.1)
Requirement already satisfied: python-dateutil>=2.7 in /usr/local/lib/python3.10/dist-packages (from matplotlib) (2.8.2)
Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.10/dist-packages (from python-dateutil>=2.7->matplotlib) (1.16.0)
```

```
[ ]: 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")
```

```
[4]: from google.colab import files

uploaded = files.upload()
```

<IPython.core.display.HTML object>

Saving Iris.csv to Iris.csv

```
[ ]: import pandas as pd
```

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

```
[ ]: iris_df = pd.read_csv("Iris.csv")
print("the data has been successfully loaded")
```

the data has been successfully loaded

```
[ ]: iris_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]

```
[ ]: iris_df.shape
```

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

```
[ ]: 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
```

```
[ ]: iris_df.describe()
```

```
[ ]:
count      Id  SepalLengthCm  SepalWidthCm  PetalLengthCm  PetalWidthCm
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
```

```
[ ]: iris_df.isnull().sum()
```

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

```
[ ]: print("unique number of values in dataset species:", iris_df["Species"].
      ↪unique())
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']
```

Exploratory Data Analysis

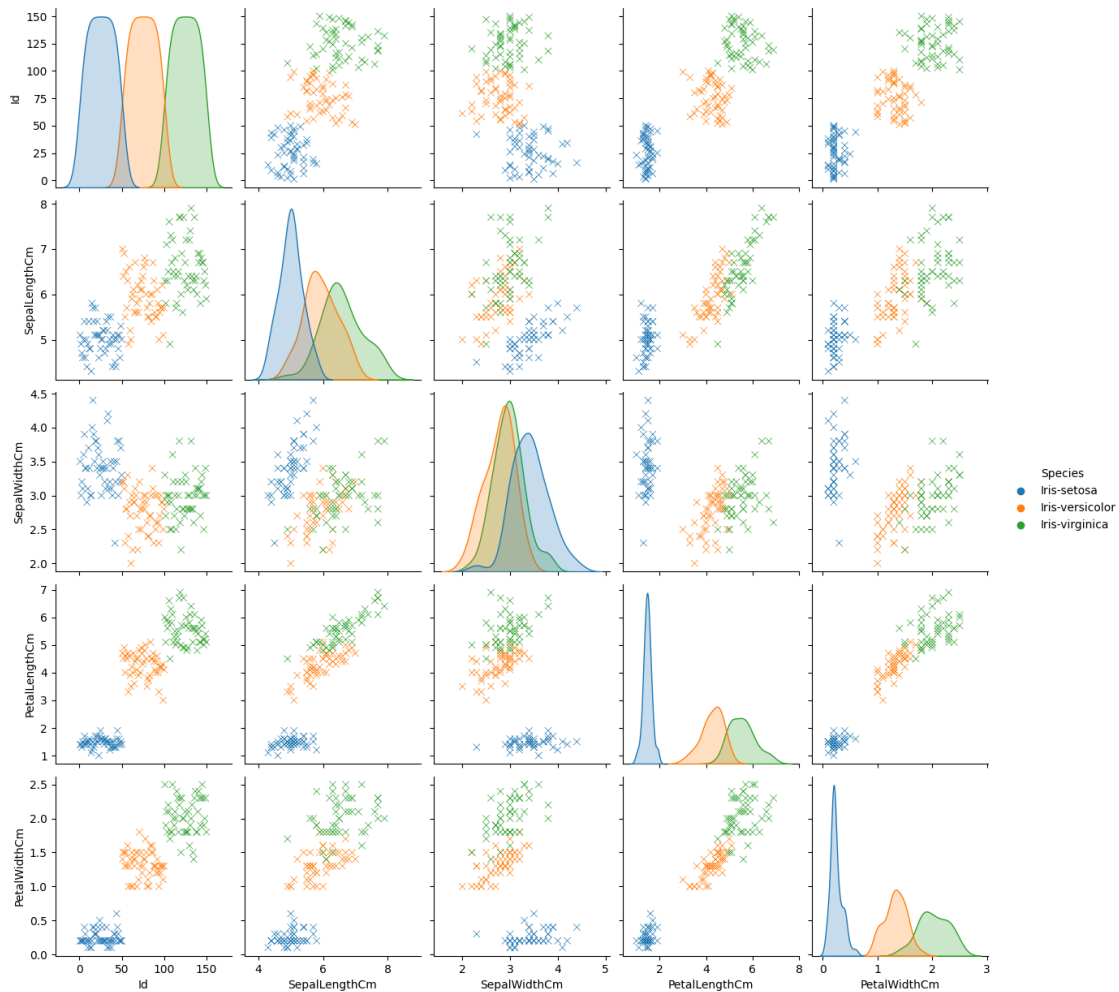
Data Visualisation

```
[2]: import seaborn as sns
```

```
[6]: import pandas as pd
iris_df = pd.read_csv('Iris.csv')
```

```
[7]: import seaborn as sns
import matplotlib.pyplot as plt

sns.pairplot(iris_df, hue="Species", markers="x")
plt.show()
```

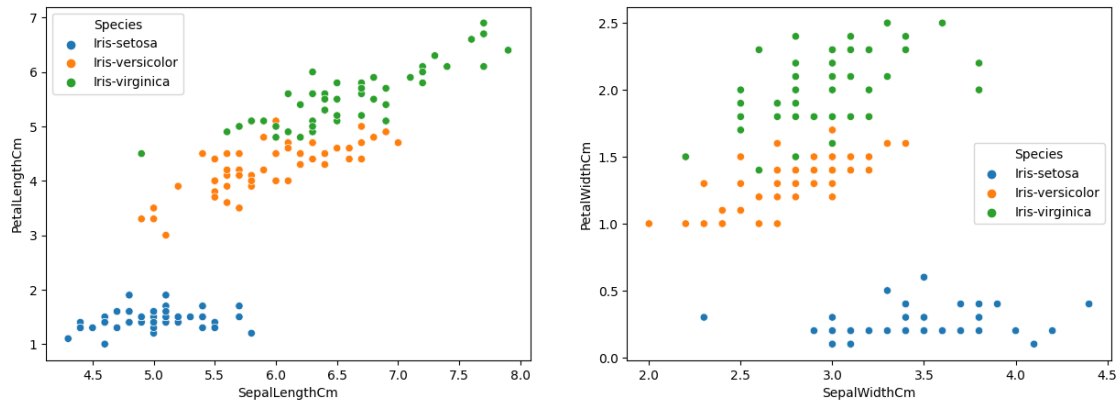


Shows that Iris-setosa is separated from both other species in all features

```
[8]: 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()
```



Check Correction in Dataset

```
[9]: iris_df.corr()
```

<ipython-input-9-1b33314f8075>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_only to silence this warning.

```
iris_df.corr()
```

```
[9]:
```

| | 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 | |

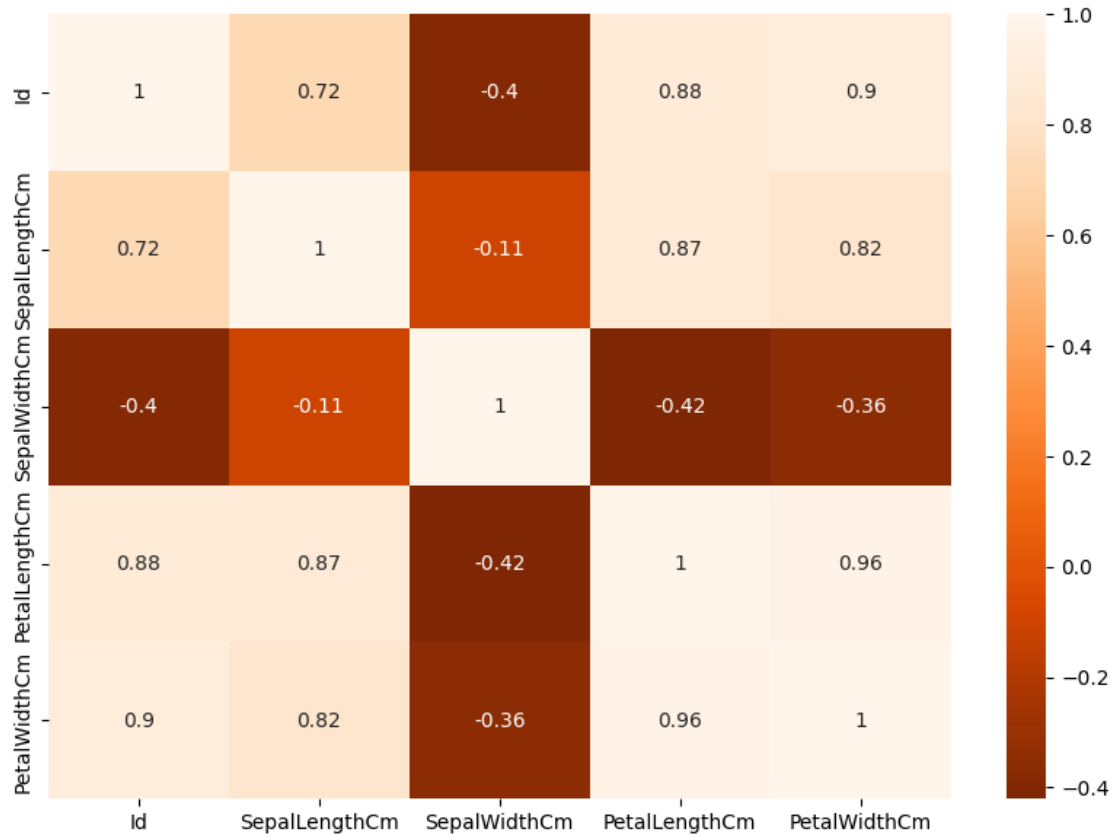
| | PetalWidthCm |
|---------------|--------------|
| Id | 0.899759 |
| SepalLengthCm | 0.817954 |
| SepalWidthCm | -0.356544 |
| PetalLengthCm | 0.962757 |
| PetalWidthCm | 1.000000 |

```
[10]: plt.figure(figsize=(10,7))
sns.heatmap(iris_df.corr(),annot = True,cmap = "Oranges_r")
plt.show()
```

<ipython-input-10-fe35941b4079>:2: FutureWarning: The default value of

numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_only to silence this warning.

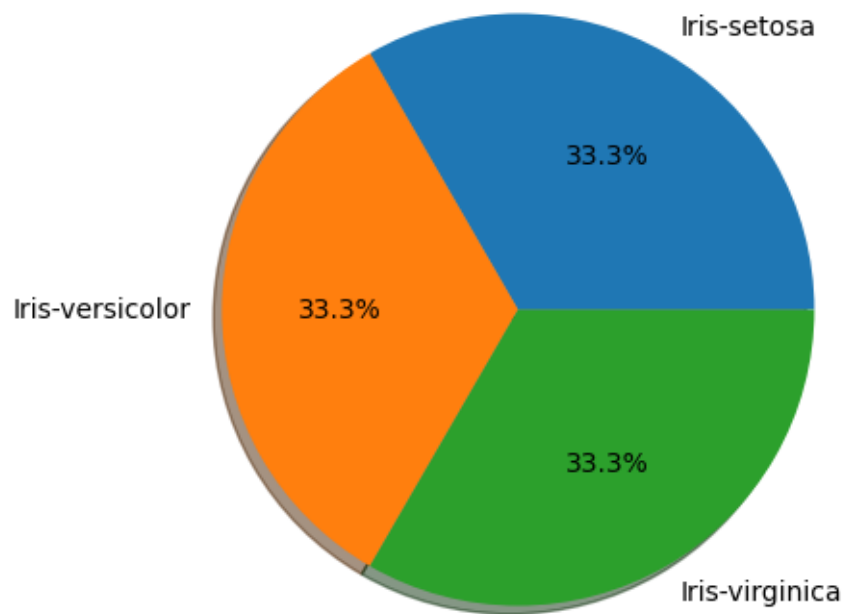
```
sns.heatmap(iris_df.corr(),annot = True,cmap = "Oranges_r")
```



We see petal length and petal width is highly correlated in above heatmap

```
[11]: iris_df["Species"].value_counts().plot(kind="pie",autopct = "%1.
    ↪1f%",shadow=True, figsize=(5,5))
plt.title("Percentage values in each species", fontsize = 12, c='g')
plt.ylabel("",fontsize=10,c="r")
plt.show()
```

Percentage values in each species

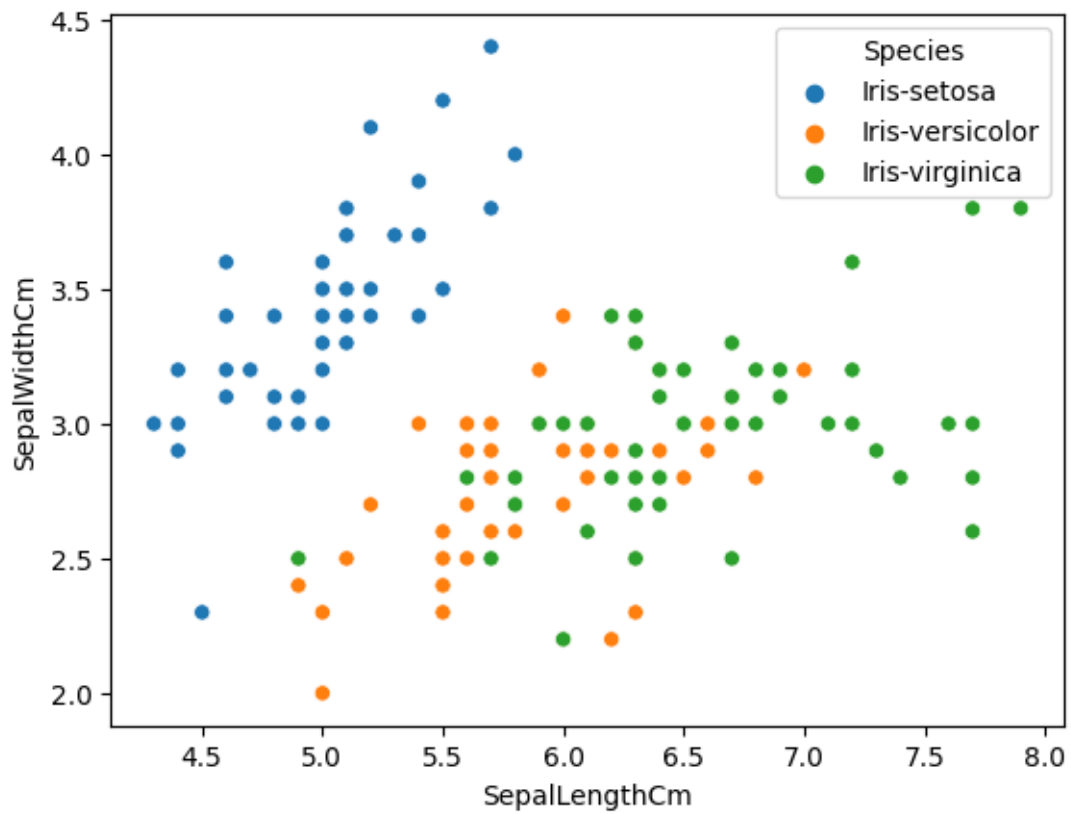


- All species have equal values in dataset
- Iris-Sentosa:50
- Iris-Versicolor:50
- Iris-Virginica:50

Scatterplot for Sepal Lenth and Sepal Width

```
[13]: import seaborn as sns
import matplotlib.pyplot as plt

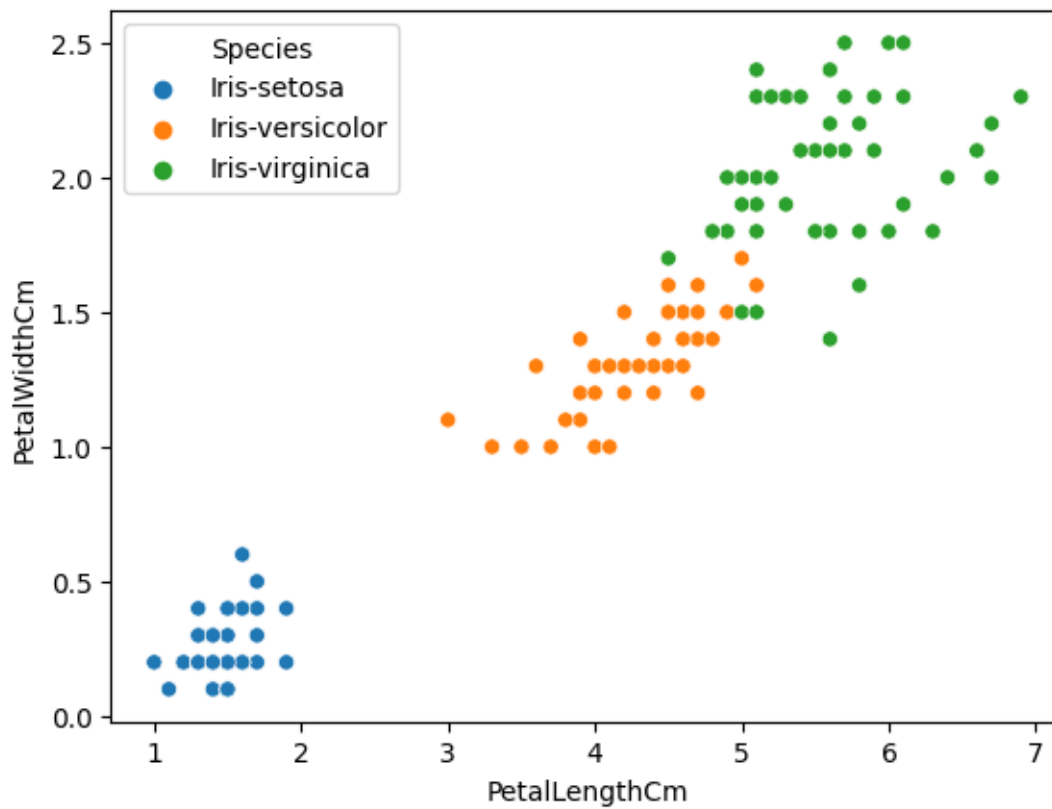
sns.scatterplot(x=iris_df["SepalLengthCm"], y=iris_df["SepalWidthCm"],
               ↪hue=iris_df["Species"])
plt.show()
```



Scatterplot for Petal Length and Petal Width

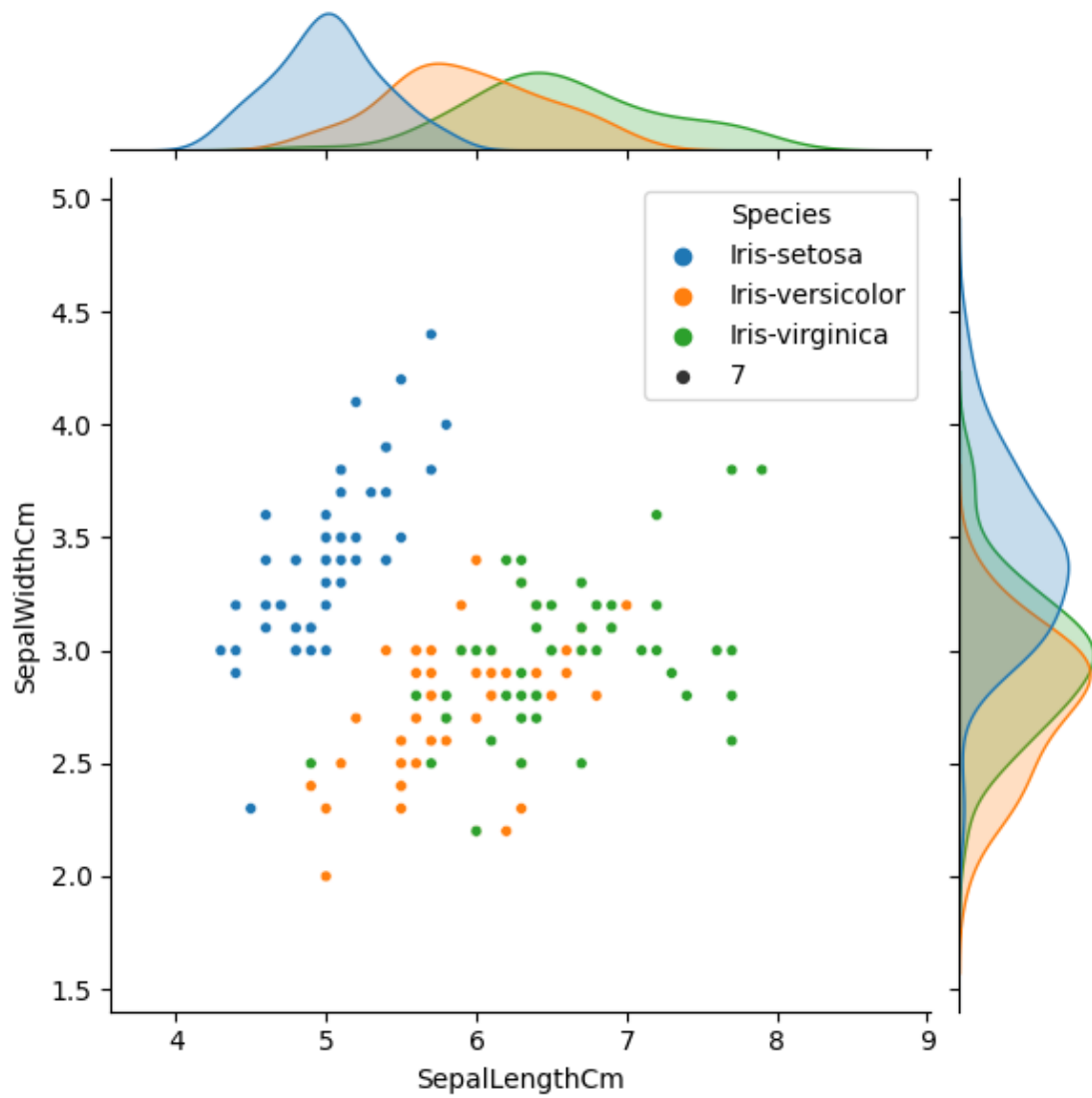
```
[15]: import seaborn as sns
import matplotlib.pyplot as plt

sns.scatterplot(x=iris_df["PetalLengthCm"], y=iris_df["PetalWidthCm"],
               hue=iris_df["Species"])
plt.show()
```

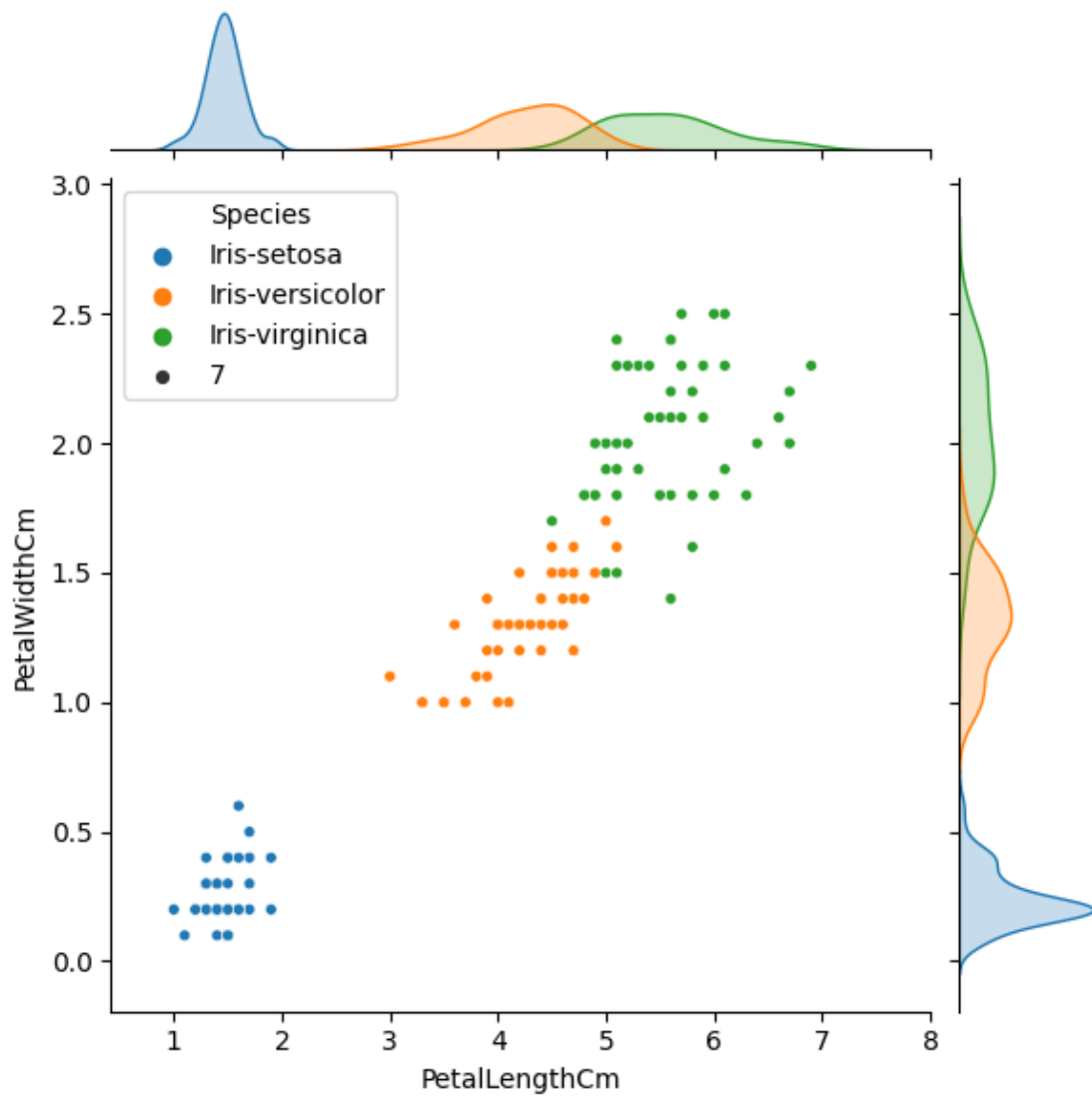



```
[16]: sns.jointplot(data = iris_df, x="SepalLengthCm", y="SepalWidthCm", size = 7,
hue = "Species")

plt.show()
```



```
[17]: sns.jointplot(data = iris_df, x = "PetalLengthCm", y = "PetalWidthCm", size = 7,
hue = "Species")
plt.show()
```



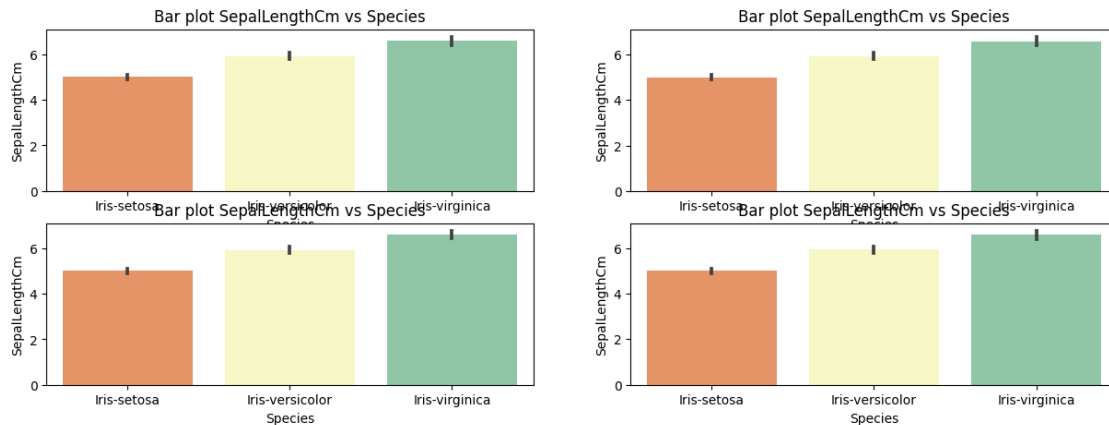
```
[18]: plt.figure(figsize = (15,5))
plt.subplot(2,2,1)
sns.barplot(x = "Species", y="SepalLengthCm",data=iris_df, palette="Spectral"))
plt.title("Bar plot SepalLengthCm vs Species")

plt.subplot(2,2,2)
sns.barplot(x = "Species", y="SepalLengthCm",data=iris_df, palette="Spectral"))
plt.title("Bar plot SepalLengthCm vs Species")

plt.subplot(2,2,3)
sns.barplot(x = "Species", y="SepalLengthCm",data=iris_df, palette="Spectral"))
plt.title("Bar plot SepalLengthCm vs Species")
```

```
plt.subplot(2,2,4)
sns.barplot(x = "Species", y="SepallLengthCm",data=iris_df, palette=("Spectral"))
plt.title("Bar plot SepallLengthCm vs Species")

plt.show()
```



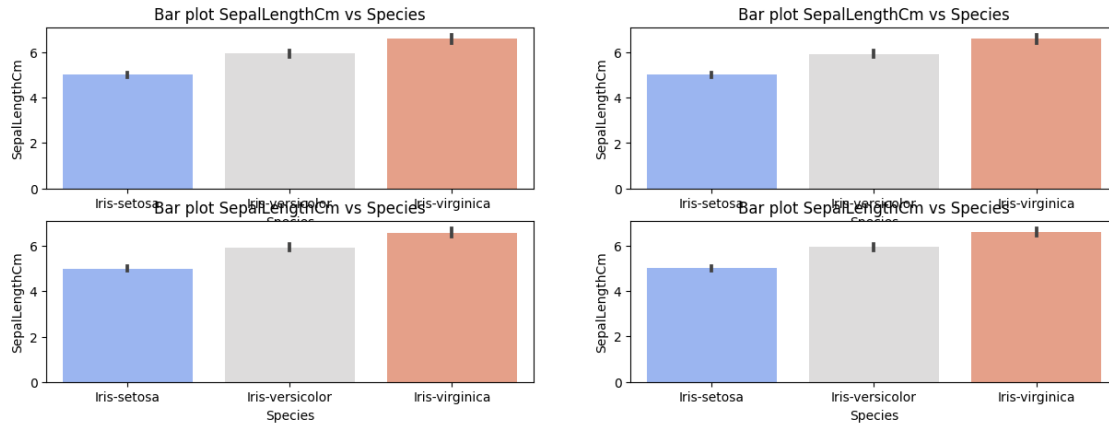
```
[19]: plt.figure(figsize = (15,5))
plt.subplot(2,2,1)
sns.barplot(x = "Species", y="SepallLengthCm",data=iris_df, palette=("coolwarm"))
plt.title("Bar plot SepallLengthCm vs Species")

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

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

plt.subplot(2,2,4)
sns.barplot(x = "Species", y="SepallLengthCm",data=iris_df, palette=("coolwarm"))
plt.title("Bar plot SepallLengthCm vs Species")

plt.show()
```



```
[22]: 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()
```

<ipython-input-22-d489ffb1486c>:3: UserWarning:

`distplot` is a deprecated function and will be removed in seaborn v0.14.0.

Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see <https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751>

```
sns.distplot(iris_df["SepalLengthCm"],color="y").set_title("Sepal Length
interval")
```

<ipython-input-22-d489ffb1486c>:6: UserWarning:

``distplot`` is a deprecated function and will be removed in seaborn v0.14.0.

Please adapt your code to use either ``displot`` (a figure-level function with similar flexibility) or ``histplot`` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see <https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751>

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

<ipython-input-22-d489ffb1486c>:9: UserWarning:

``distplot`` is a deprecated function and will be removed in seaborn v0.14.0.

Please adapt your code to use either ``displot`` (a figure-level function with similar flexibility) or ``histplot`` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see <https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751>

```
sns.distplot(iris_df["PetalLengthCm"],color="g").set_title("Petal Length interval")
```

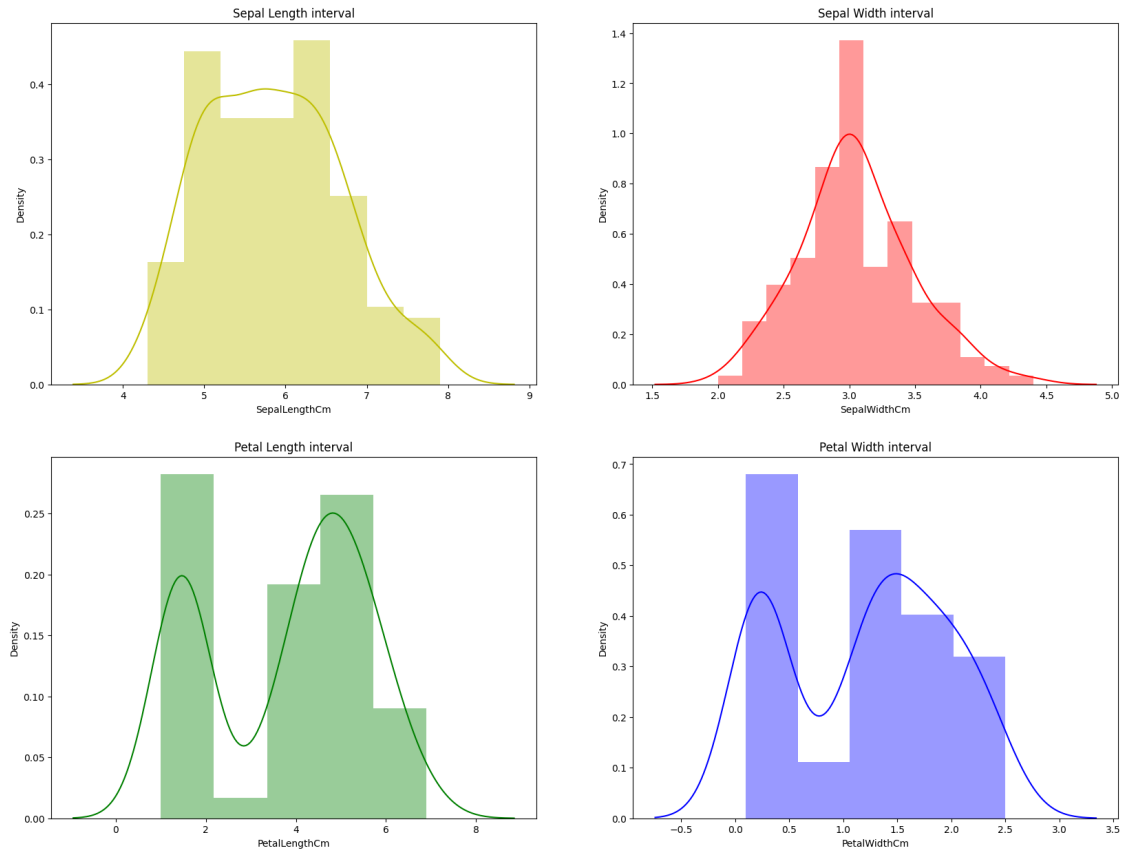
<ipython-input-22-d489ffb1486c>:12: UserWarning:

``distplot`` is a deprecated function and will be removed in seaborn v0.14.0.

Please adapt your code to use either ``displot`` (a figure-level function with similar flexibility) or ``histplot`` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see <https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751>

```
sns.distplot(iris_df["PetalWidthCm"],color="b").set_title("Petal Width interval")
```



Data Cleaning

```
[24]: from sklearn.preprocessing import LabelEncoder
```

```
[25]: from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()

iris_df["Species"] = le.fit_transform(iris_df["Species"])
iris_df.head()
```

```
[25]:
```

| | Id | SepalLengthCm | SepalWidthCm | Petal.LengthCm | 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 |

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

```
[26]:
```

| | 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 |

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

```
[27]: 0    0
1    0
2    0
3    0
4    0
Name: Species, dtype: int64
```

```
[28]: print(X.shape)
print(y.shape)
```

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

Model Building

```
[29]: 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=0)
```

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

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

Logistic Regression

```
[33]: 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)
```



```

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:-")
from sklearn.metrics import 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 |

/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_logistic.py:458:

ConvergenceWarning: lbfgs failed to converge (status=1):

STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max_iter) or scale the data as shown in:

<https://scikit-learn.org/stable/modules/preprocessing.html>

Please also refer to the documentation for alternative solver options:

https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression

```
n_iter_i = _check_optimize_result(
```

Random Forest Classifier

```
[35]: from sklearn.ensemble import RandomForestClassifier
rfc= RandomForestClassifier()

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

y_pred = rfc.predict(X_test)

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:-")
from sklearn.metrics import classification_report

# ...

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

Random Forest Classifier 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 |

Decision Tree

```
[36]: from sklearn.tree import DecisionTreeClassifier

dtc = DecisionTreeClassifier()
```

```

dtc.fit(X_train, y_train)

print("Decision Tree Classifier successfully implemented")

y_pred = dtc.predict(X_test)

cm = confusion_matrix(y_test, y_pred)

print("Confusion Matrix:")
print(cm)
accuracy = accuracy_score(y_test, y_pred)

print("Accuracy:", accuracy*100, "%")

from sklearn.metrics import classification_report
print("Classification Report:")
print(classification_report(y_test, y_pred))

```

Decision Tree Classifier successfully implemented

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 |

```
[37]: from sklearn.tree import plot_tree
```

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

Support Vector Machine

```
[22]: from google.colab import files
```

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

<IPython.core.display.HTML object>

Saving Iris.csv to Iris.csv

```
[23]: import pandas as pd

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

```
[27]: import seaborn as sns

# Load the Iris dataset
iris_df = sns.load_dataset('iris')

# Now you can proceed with your machine learning tasks using `iris_df`
```

```
[28]: import pandas as pd

# Load your dataset
your_dataset_df = pd.read_csv('Iris.csv')

# Now you can proceed with your machine learning tasks using `your_dataset_df`
```

```
[33]: X = iris_df.drop(columns=["species"]) # Features
y = iris_df["species"] # Labels

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=42)
from sklearn.svm import SVC

svc = SVC()
svc.fit(X_train, y_train)

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

y_pred = svc.predict(X_test)

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

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:-

```
[[10  0  0]
 [ 0  9  0]
 [ 0  0 11]]
```

accuracy:- 100.0

Classification Report:-

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| setosa | 1.00 | 1.00 | 1.00 | 10 |
| versicolor | 1.00 | 1.00 | 1.00 | 9 |
| virginica | 1.00 | 1.00 | 1.00 | 11 |
| accuracy | | | 1.00 | 30 |
| macro avg | 1.00 | 1.00 | 1.00 | 30 |
| weighted avg | 1.00 | 1.00 | 1.00 | 30 |

```
[32]: print(iris_df.columns)
```

```
Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
      'species'],
      dtype='object')
```

K-NN Classifier

```
[34]: from sklearn.neighbors import KNeighborsClassifier
      from sklearn.metrics import accuracy_score, classification_report, \
      ↪confusion_matrix

      knn = KNeighborsClassifier(n_neighbors=5) # You can change the number of \
      ↪neighbors if needed
      knn.fit(X_train, y_train)

      y_pred = knn.predict(X_test)

      cm = confusion_matrix(y_test, y_pred)

      print("Confusion Matrix:")
      print(cm)

      accuracy = accuracy_score(y_test, y_pred)

      print("Accuracy:", accuracy*100, "%")

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

Confusion Matrix:

```
[[10  0  0]
 [ 0  9  0]
 [ 0  0 11]]
Accuracy: 100.0 %
Classification Report:
              precision    recall  f1-score   support

   setosa         1.00        1.00        1.00        10
  versicolor      1.00        1.00        1.00         9
   virginica      1.00        1.00        1.00        11

 accuracy         1.00        1.00        1.00        30
 macro avg        1.00        1.00        1.00        30
weighted avg        1.00        1.00        1.00        30
```

Naive Bayes

```
[35]: from sklearn.naive_bayes import GaussianNB
      from sklearn.metrics import accuracy_score, classification_report, \
      ↪confusion_matrix

naive_bayes = GaussianNB()

naive_bayes.fit(X_train, y_train)

y_pred = naive_bayes.predict(X_test)

cm = confusion_matrix(y_test, y_pred)

print("Confusion Matrix:")
print(cm)

accuracy = accuracy_score(y_test, y_pred)

print("Accuracy:", accuracy*100, "%")

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

```
Confusion Matrix:
[[10  0  0]
 [ 0  9  0]
 [ 0  0 11]]
Accuracy: 100.0 %
Classification Report:
              precision    recall  f1-score   support

   setosa         1.00        1.00        1.00        10
  versicolor      1.00        1.00        1.00         9
   virginica      1.00        1.00        1.00        11

 accuracy         1.00        1.00        1.00        30
 macro avg        1.00        1.00        1.00        30
weighted avg        1.00        1.00        1.00        30
```

| | | | | |
|--------------|------|------|------|----|
| setosa | 1.00 | 1.00 | 1.00 | 10 |
| versicolor | 1.00 | 1.00 | 1.00 | 9 |
| virginica | 1.00 | 1.00 | 1.00 | 11 |
| accuracy | | | 1.00 | 30 |
| macro avg | 1.00 | 1.00 | 1.00 | 30 |
| weighted avg | 1.00 | 1.00 | 1.00 | 30 |

Test Model

```
[38]: import numpy as np
```

```
[46]: from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split

X = iris_df.drop(columns=["species"])
y = iris_df["species"]

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
↳random_state=42)

dtc = DecisionTreeClassifier()

dtc.fit(X_train, y_train)
input_data = [[4.9, 3.0, 1.4, 0.2]]

input_data_as_ndarray = np.asarray(input_data)
input_data_reshaped = input_data_as_ndarray.reshape(1,-1)

prediction = dtc.predict(input_data_reshaped)
print("The category is", prediction)
```

The category is ['setosa']

/usr/local/lib/python3.10/dist-packages/sklearn/base.py:439: UserWarning: X does not have valid feature names, but DecisionTreeClassifier was fitted with feature names

```
warnings.warn(
```

```
[ ]:
```