



Iris flower prediction

importing libraries

In [1]:

```
import pandas as pd
import numpy as np
from matplotlib import pyplot as plt
%matplotlib inline
import seaborn as sns
import warnings
warnings.filterwarnings('ignore')
pd.options.display.max_rows=5000
```

In [2]:

```
aggtype = ['mean', 'std']
```

In [3]:

```
df=pd.read_csv("C:/Users/ayith/Downloads/Iris.csv") # reading dataset
```

In [4]:

```
df.head()
```

Out[4]:

	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

checking unique values

In [5]:

```
df['Species'].unique()
```

Out[5]:

```
array(['Iris-setosa', 'Iris-versicolor', 'Iris-virginica'], dtype=object)
```

In [6]:

```
df['PetalLengthCm'].unique()
```

Out[6]:

```
array([1.4, 1.3, 1.5, 1.7, 1.6, 1.1, 1.2, 1. , 1.9, 4.7, 4.5, 4.9, 4. ,
       4.6, 3.3, 3.9, 3.5, 4.2, 3.6, 4.4, 4.1, 4.8, 4.3, 5. , 3.8, 3.7,
       5.1, 3. , 6. , 5.9, 5.6, 5.8, 6.6, 6.3, 6.1, 5.3, 5.5, 6.7, 6.9,
       5.7, 6.4, 5.4, 5.2])
```

In [7]:

```
df[df.duplicated()]      #checking duplicates
```

Out[7]:

<u>Id</u>	<u>SepalLengthCm</u>	<u>SepalWidthCm</u>	<u>PetalLengthCm</u>	<u>PetalWidthCm</u>	<u>Species</u>
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In [8]:

```
df.isnull().sum()      #finding null values
```

Out[8]:

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

In [9]:

```
df.info
```

Out[9]:

```
<bound method DataFrame.info 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
5      6           5.4           3.9           1.7           0.4
6      7           4.6           3.4           1.4           0.3
7      8           5.0           3.4           1.5           0.2
8      9           4.4           2.9           1.4           0.2
9     10           4.9           3.1           1.5           0.1
10    11           5.4           3.7           1.5           0.2
11    12           4.8           3.4           1.6           0.2
12    13           4.8           3.0           1.4           0.1
13    14           4.3           3.0           1.1           0.1
14    15           5.8           4.0           1.2           0.2
15    16           5.7           4.4           1.5           0.4
```

In [10]:

```
df.describe().T
```

Out[10]:

	count	mean	std	min	25%	50%	75%	max
Id	150.0	75.500000	43.445368	1.0	38.25	75.50	112.75	150.0
SepalLengthCm	150.0	5.843333	0.828066	4.3	5.10	5.80	6.40	7.9
SepalWidthCm	150.0	3.054000	0.433594	2.0	2.80	3.00	3.30	4.4
PetalLengthCm	150.0	3.758667	1.764420	1.0	1.60	4.35	5.10	6.9
PetalWidthCm	150.0	1.198667	0.763161	0.1	0.30	1.30	1.80	2.5

In [11]:

```
df.groupby('Species').agg(aggtype).T
```

Out[11]:

Species		Iris-setosa	Iris-versicolor	Iris-virginica
Id	mean	25.500000	75.500000	125.500000
	std	14.577380	14.577380	14.577380
SepalLengthCm	mean	5.006000	5.936000	6.588000
	std	0.352490	0.516171	0.635880
SepalWidthCm	mean	3.418000	2.770000	2.974000
	std	0.381024	0.313798	0.322497
PetalLengthCm	mean	1.464000	4.260000	5.552000
	std	0.173511	0.469911	0.551895
PetalWidthCm	mean	0.244000	1.326000	2.026000
	std	0.107210	0.197753	0.274650

In [12]:

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

Out[12]:

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

In [13]:

```
pd.crosstab(df['Species'],df['SepalLengthCm']).sum()
```

Out[13]:

SepalLengthCm

4.3	1
4.4	3
4.5	1
4.6	4
4.7	2
4.8	5
4.9	6
5.0	10
5.1	9
5.2	4
5.3	1
5.4	6
5.5	7
5.6	6
5.7	8
5.8	7
5.9	3
6.0	6
6.1	6
6.2	4
6.3	9
6.4	7
6.5	5
6.6	2
6.7	8
6.8	3
6.9	4
7.0	1
7.1	1
7.2	3
7.3	1
7.4	1
7.6	1
7.7	4
7.9	1

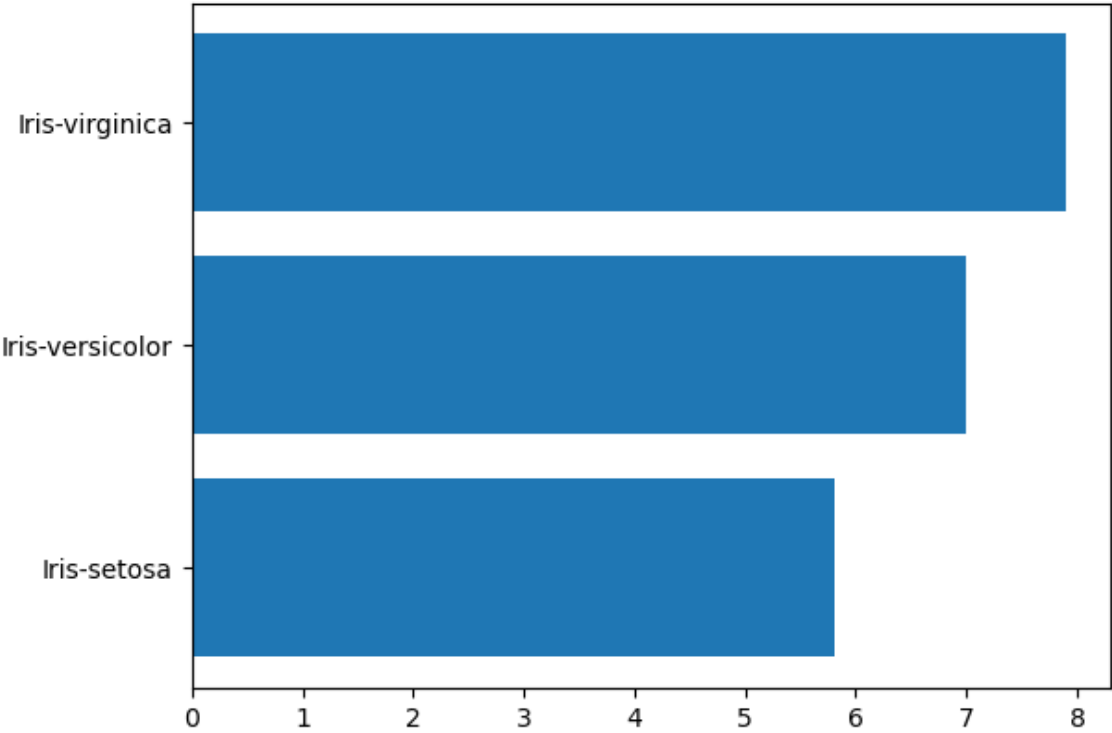
dtype: int64

Data graphs

In [14]:

```
fig,ax=plt.subplots()
y=ax.barh('Species','SepalLengthCm',data=df)

plt.show()
```



In [15]:

```
df
```

Out[15]:

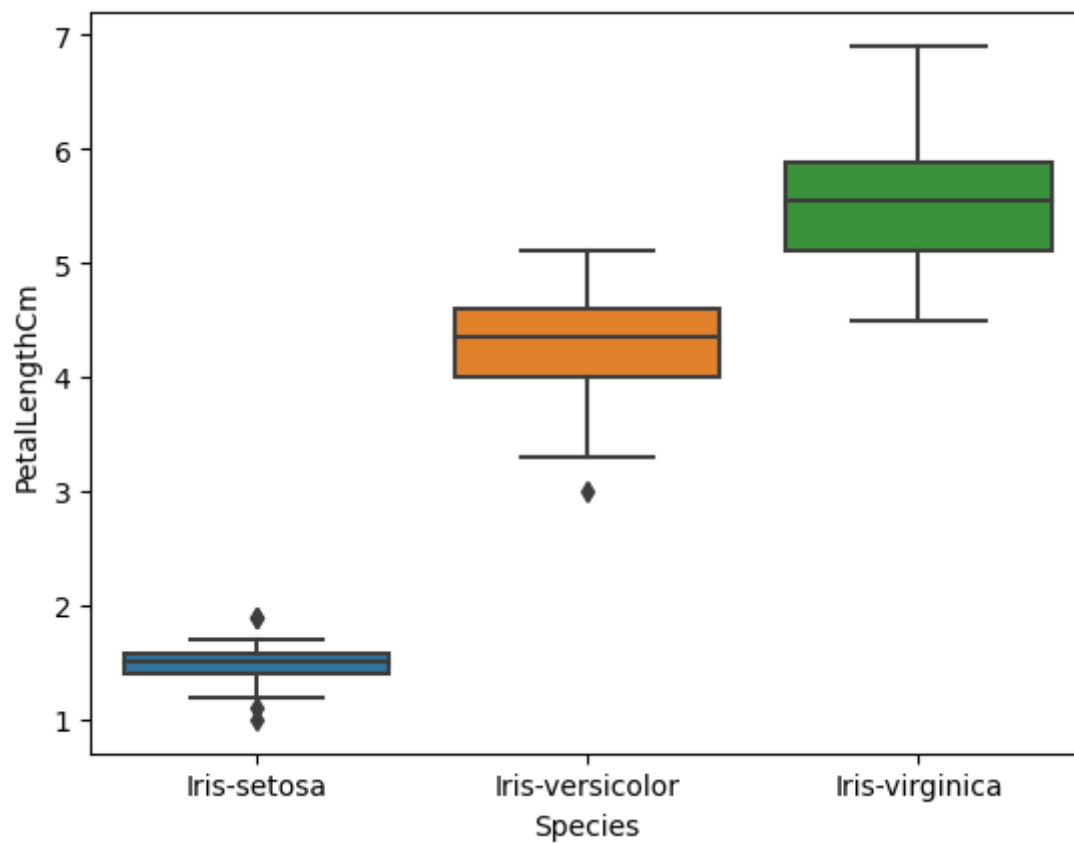
	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa
5	6	5.4	3.9	1.7	0.4	Iris-setosa
6	7	4.6	3.4	1.4	0.3	Iris-setosa
7	8	5.0	3.4	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

In [16]:

```
sns.boxplot(x = 'Species', y = 'PetalLengthCm', data=df)
```

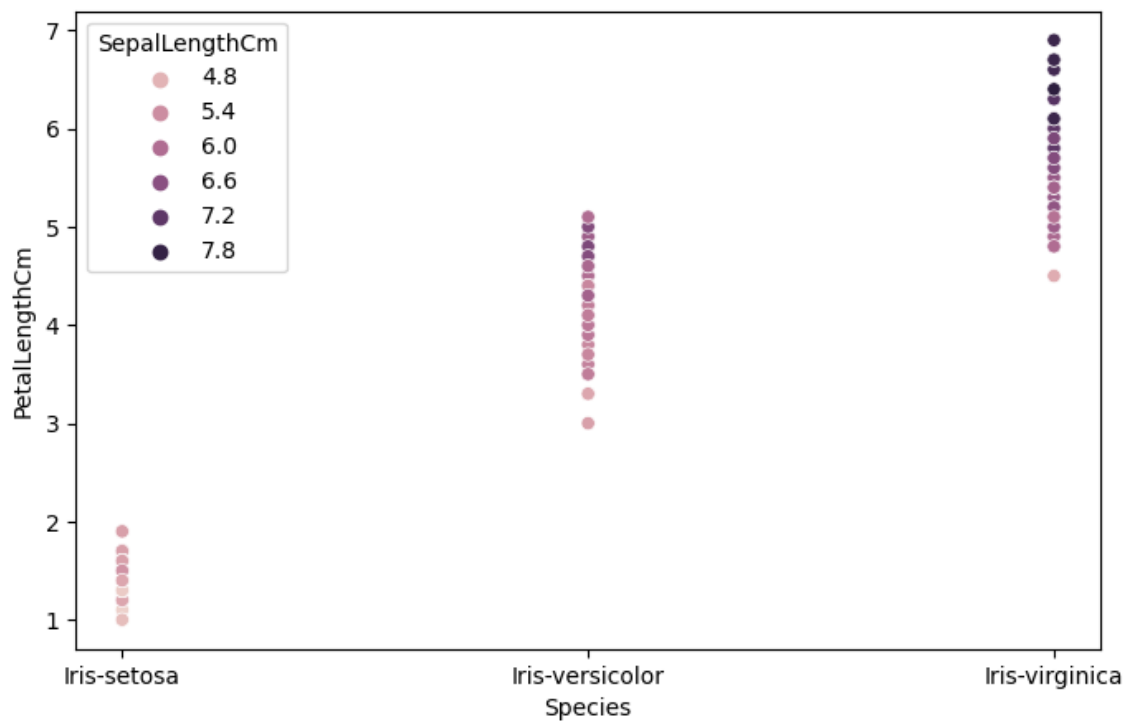
Out[16]:

<AxesSubplot:xlabel='Species', ylabel='PetalLengthCm'>



In [17]:

```
plt.figure(figsize=(8,5))  
sns.scatterplot(x = 'Species', y = 'PetalLengthCm', hue = 'SepalLengthCm', data = df)  
plt.show()
```



In [18]:

```
plt.figure(figsize=(15,10))
plt.suptitle('Bivariate Analysis of Categorical Features & numerical features', fontsize

plt.subplot(2,2,1)
sns.boxplot(x = df['Species'], y = df['PetalWidthCm'])
plt.title("Boxplot")

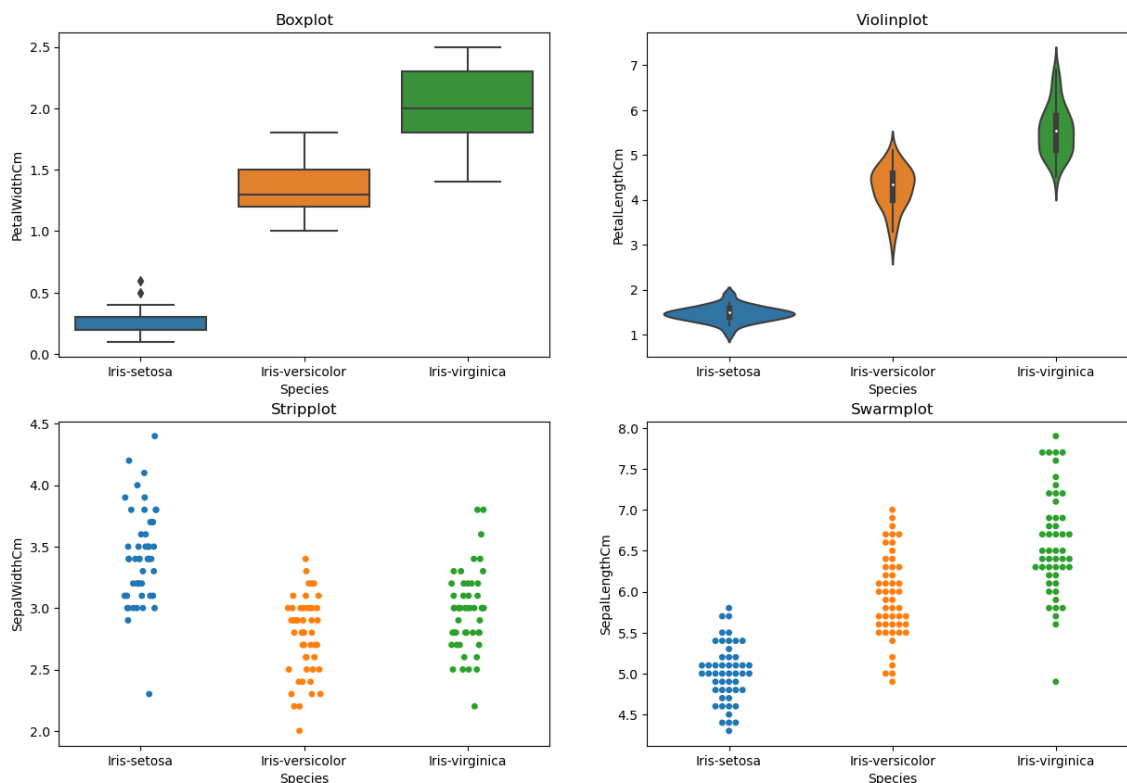
plt.subplot(2,2,2)
sns.violinplot(x = df['Species'], y = df['PetalLengthCm'])
plt.title("Violinplot")

plt.subplot(2,2,3)
sns.stripplot(x = df['Species'], y = df['SepalWidthCm'])
plt.title("Stripplot")

plt.subplot(2,2,4)
sns.swarmplot(x = df['Species'], y = df['SepalLengthCm'])
plt.title("Swarmplot")

plt.show()
```

Bivariate Analysis of Categorical Features & numerical features



In [19]:

```
num_features=[feature for feature in df.columns if df[feature].dtype != 'O']
```

In [20]:

```
num_features
```

Out[20]:

```
['Id', 'SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm']
```

In [21]:

```
cat_features=[feature for feature in df.columns if df[feature].dtype == 'O']
```

In [22]:

```
cat_features
```

Out[22]:

```
['Species']
```

In [23]:

```
for col in num_features:  
    print(f"{col}:{df[col].value_counts(normalize=True)*100}")  
    print("=====")
```

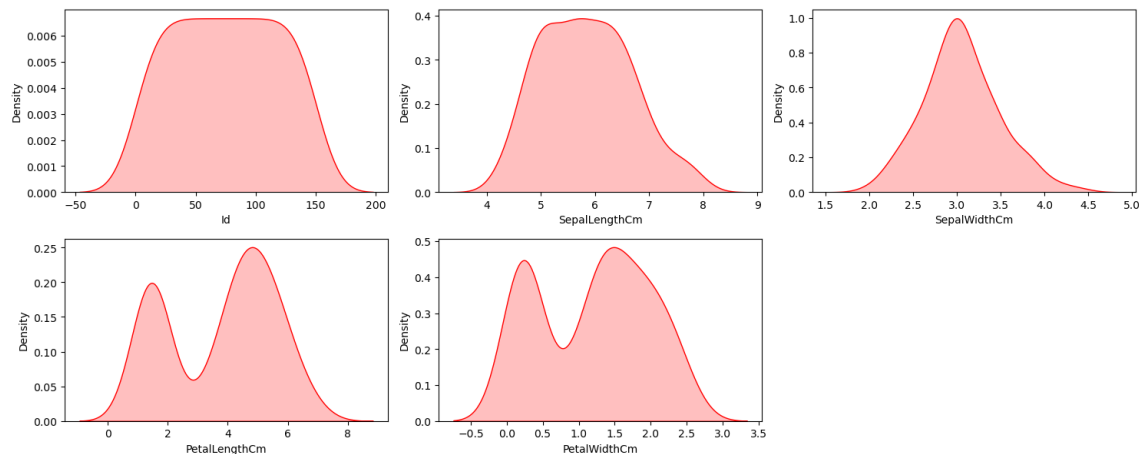
```
Id:1      0.666667  
95      0.666667  
97      0.666667  
98      0.666667  
99      0.666667  
100     0.666667  
101     0.666667  
102     0.666667  
103     0.666667  
104     0.666667  
105     0.666667  
106     0.666667  
107     0.666667  
108     0.666667  
109     0.666667  
110     0.666667  
111     0.666667  
96      0.666667  
94      0.666667  
112     0.666667
```

In [24]:

```
plt.figure(figsize=(15, 15))
plt.suptitle('Univariate Analysis of Numerical Features', fontsize=20, fontweight='bold')

for i in range(0, len(num_features)):
    plt.subplot(5, 3, i+1)
    sns.kdeplot(x=df[num_features[i]], shade=True, color='r')
    plt.xlabel(num_features[i])
    plt.tight_layout()
```

Univariate Analysis of Numerical Features

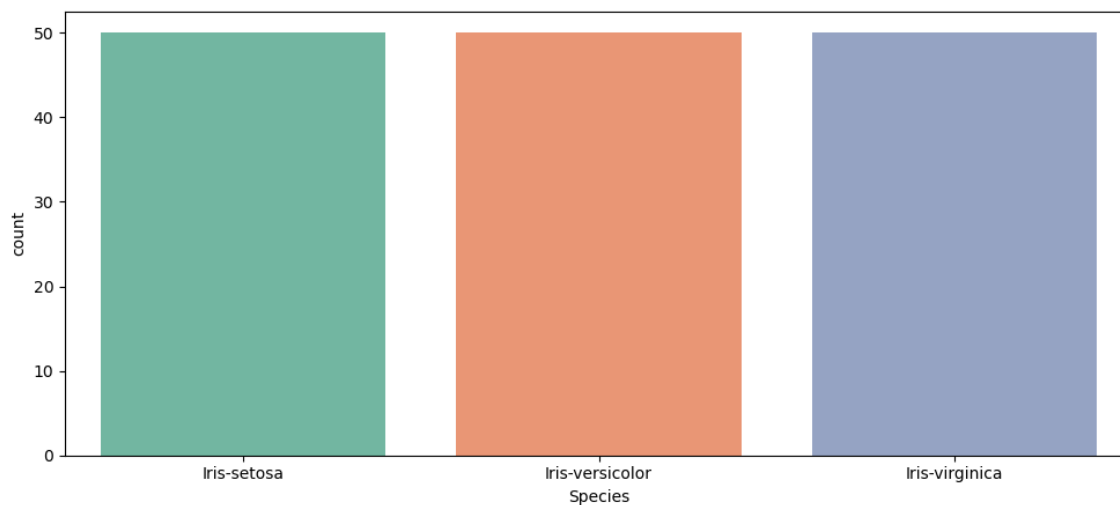


In [25]:

```
plt.figure(figsize=(10,5))
plt.suptitle('Univariate Analysis of Categorical Features', fontsize=20, fontweight='bold')
category = [ 'Species' ]
for i in range(0, len(category)):

    sns.countplot(x=df[category[i]], palette="Set2")
    plt.xlabel(category[i])
    plt.xticks(rotation=0)
    plt.tight_layout()
```

Univariate Analysis of Categorical Features



Lables encodeing

In [26]:

```
# Import Label encoder
from sklearn import preprocessing

# Label_encoder object knows how to understand word Labels.
label_encoder = preprocessing.LabelEncoder()

# Encode labels in column 'species'.
df['Species'] = label_encoder.fit_transform(df['Species'])

df['Species'].unique()
```

Out[26]:

```
array([0, 1, 2])
```

In [27]:

```
from sklearn import preprocessing
```

In [28]:

```
labe_encodeing= preprocessing.LabelEncoder()
df['Species']=labe_encodeing.fit_transform(df['Species'])
df['Species'].unique()
```

Out[28]:

```
array([0, 1, 2], dtype=int64)
```

In [29]:

```
df.sample(5)
```

Out[29]:

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
37	38	4.9	3.1	1.5	0.1	0
132	133	6.4	2.8	5.6	2.2	2
102	103	7.1	3.0	5.9	2.1	2
2	3	4.7	3.2	1.3	0.2	0
33	34	5.5	4.2	1.4	0.2	0

In [30]:

df

Out[30]:

	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
5	6	5.4	3.9	1.7	0.4	0
6	7	4.6	3.4	1.4	0.3	0
7	8	5.0	3.4	1.5	0.2	0
8	9	4.4	2.9	1.4	0.2	0
9	10	4.9	3.1	1.5	0.1	0
10	11	5.4	3.7	1.5	0.2	0

Data splitting

In [31]:

```
x = df.iloc[:, [1,2,3, 4]].values  
y=df.iloc[:,[5]].values
```

In [32]:

```
from sklearn.model_selection import train_test_split  
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.25, random_state
```

In [33]:

```
X_train.shape,X_test.shape,y_train.shape,y_test.shape
```

Out[33]:

```
((112, 4), (38, 4), (112, 1), (38, 1))
```

In [34]:

```
type(y_test)
```

Out[34]:

```
numpy.ndarray
```

scaling data

In [35]:

```
from sklearn.preprocessing import StandardScaler
```

In [36]:

```
sc_X=StandardScaler()
```

In [37]:

```
xtrain = sc_X.fit_transform(X_train)
```

In [38]:

```
xtest = sc_X.fit_transform(X_test)
```

creating model

In [39]:

```
from sklearn.linear_model import LogisticRegression  
classifier = LogisticRegression(random_state = 0)  
classifier.fit(X_train, y_train)
```

Out[39]:

```
LogisticRegression(random_state=0)
```

In [40]:

```
y_pred = classifier.predict(X_test)
```

In [41]:

```
from sklearn.metrics import confusion_matrix  
cm = confusion_matrix(y_test, y_pred)
```

In [42]:

```
print('accuracy of model: ', classifier.score(X_test, y_test))
```

```
accuracy of model: 1.0
```

In [43]:

```
from sklearn.model_selection import cross_val_score  
cvs = cross_val_score(classifier, X_train, y_train, cv=10)  
print('accuracy of model after CVS: ', np.mean(cvs))
```

```
accuracy of model after CVS: 0.9469696969696969
```

checking the predict values with original values

In [44]:

```
y_test=y_test.flatten()
predict = classifier.predict(X_test)
compar = pd.DataFrame({'actual':y_test, 'predicted': predict})
compar = compar.reset_index(drop = True)
compar[:10]
```

Out[44]:

	actual	predicted
0	0	0
1	0	0
2	0	0
3	0	0
4	1	1
5	1	1
6	1	1
7	0	0
8	1	1
9	2	2

In [45]:

```
from sklearn.metrics import accuracy_score
print ("Accuracy : ", accuracy_score(y_test, y_pred))
```

Accuracy : 1.0

In [46]:

```
from sklearn.metrics import classification_report
```

In [47]:

```
print(classification_report(y_test, predict))
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	13
1	1.00	1.00	1.00	13
2	1.00	1.00	1.00	12
accuracy			1.00	38
macro avg	1.00	1.00	1.00	38
weighted avg	1.00	1.00	1.00	38

In []:

In []: