

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import accuracy_score
```

```
In [2]: df=pd.read_csv('Iris.csv')
```

```
In [3]: df.head()
```

```
Out[3]:
```

	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

```
In [4]: df.tail()
```

```
Out[4]:
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
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

```
In [5]: df.shape
```

```
Out[5]: (150, 6)
```

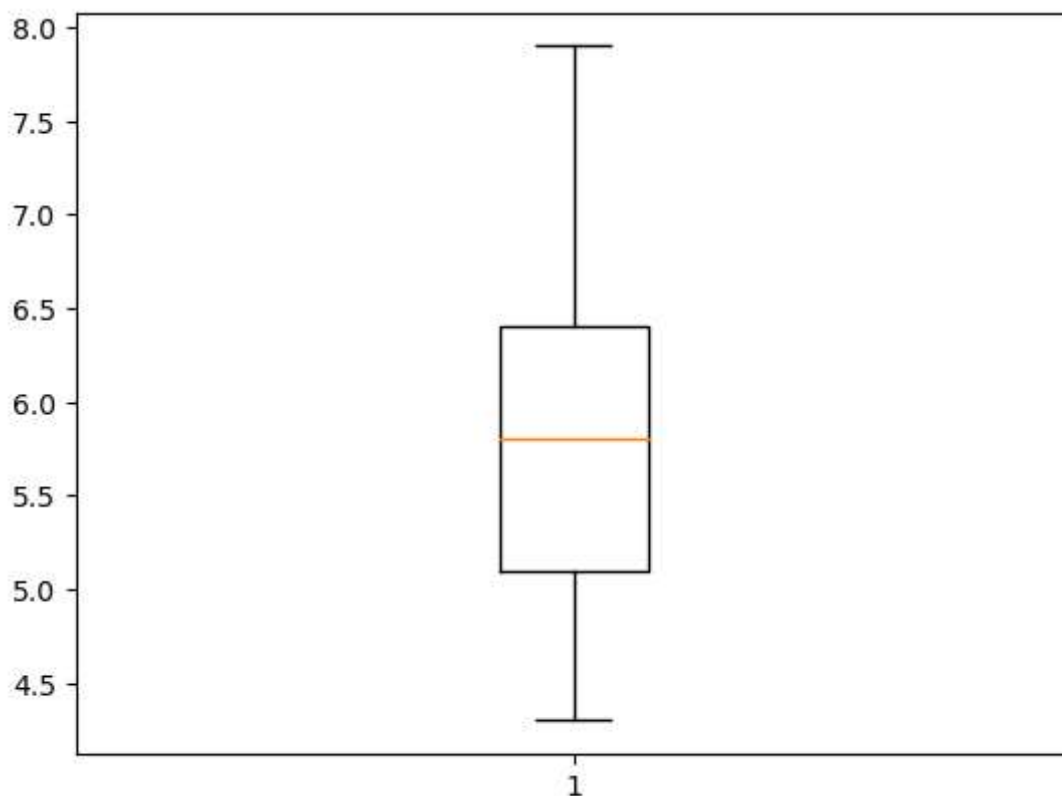
```
In [6]: data=df.groupby('Species')
```

```
In [7]: df['Species'].unique()
```

```
Out[7]: array(['Iris-setosa', 'Iris-versicolor', 'Iris-virginica'], dtype=object)
```

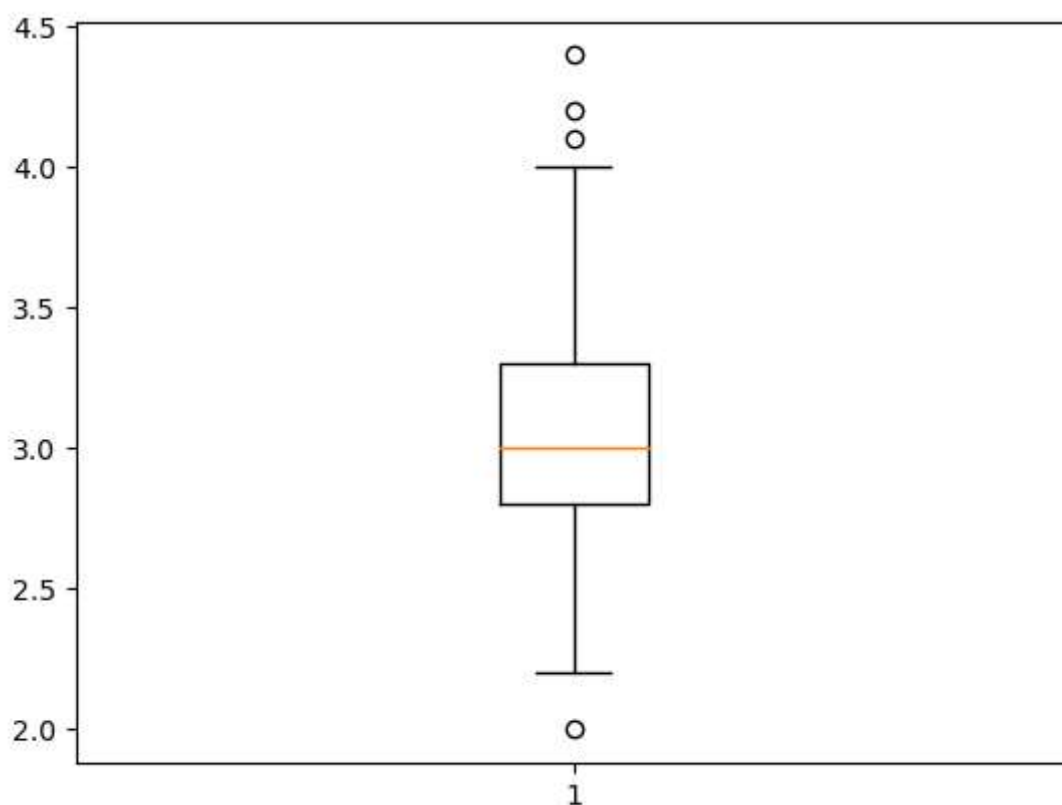
```
In [8]: #2. visualizing the dataset
plt.boxplot(df['SepalLengthCm'])
```

```
Out[8]: {'whiskers': [<matplotlib.lines.Line2D at 0x2389ad93210>,
<matplotlib.lines.Line2D at 0x2389ba11d10>],
'caps': [<matplotlib.lines.Line2D at 0x2389b414d50>,
<matplotlib.lines.Line2D at 0x2389ba2d790>],
'boxes': [<matplotlib.lines.Line2D at 0x2389ba118d0>],
'medians': [<matplotlib.lines.Line2D at 0x2389ba2e390>],
'fliers': [<matplotlib.lines.Line2D at 0x2389b2714d0>],
'means': []}
```



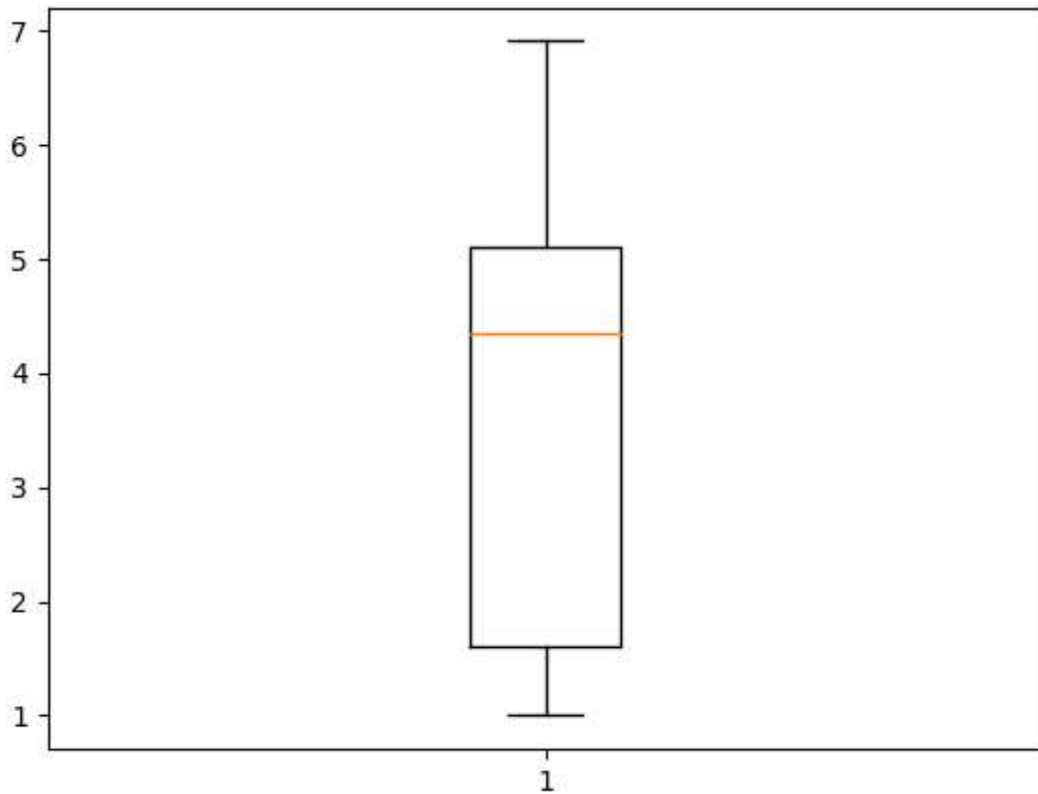
```
In [9]: plt.boxplot(df['SepalWidthCm'])
```

```
Out[9]: {'whiskers': [<matplotlib.lines.Line2D at 0x2389baf8e50>,
<matplotlib.lines.Line2D at 0x2389baf9a90>],
'caps': [<matplotlib.lines.Line2D at 0x2389bafa650>,
<matplotlib.lines.Line2D at 0x2389bafb0d0>],
'boxes': [<matplotlib.lines.Line2D at 0x2389baf8290>],
'medians': [<matplotlib.lines.Line2D at 0x2389bafbc90>],
'fliers': [<matplotlib.lines.Line2D at 0x2389bac20d0>],
'means': []}
```



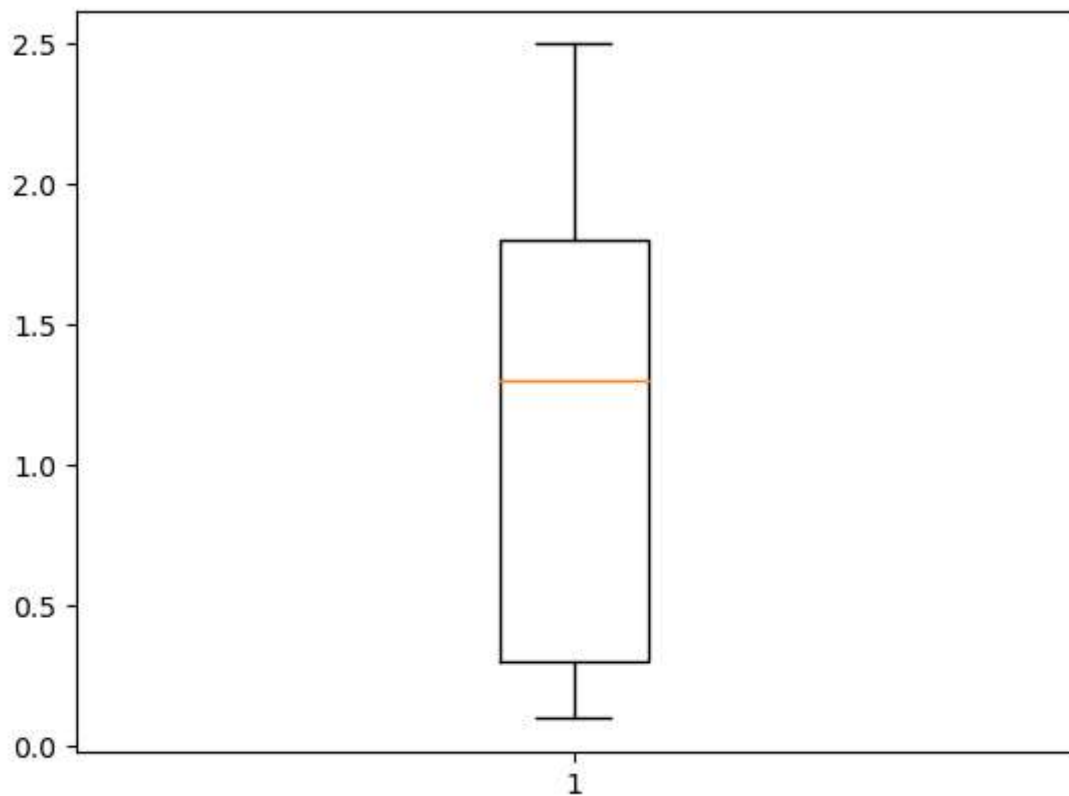
```
In [10]: plt.boxplot(df['PetalLengthCm'])
```

```
Out[10]: {'whiskers': [<matplotlib.lines.Line2D at 0x2389bb66290>,  
<matplotlib.lines.Line2D at 0x2389bb67010>],  
'caps': [<matplotlib.lines.Line2D at 0x2389bb67b50>,  
<matplotlib.lines.Line2D at 0x2389bb6c710>],  
'boxes': [<matplotlib.lines.Line2D at 0x2389bb65610>],  
'medians': [<matplotlib.lines.Line2D at 0x2389bb6d190>],  
'fliers': [<matplotlib.lines.Line2D at 0x2389bb1dbd0>],  
'means': []}
```



```
In [11]: plt.boxplot(df['PetalWidthCm'])
```

```
Out[11]: {'whiskers': [<matplotlib.lines.Line2D at 0x2389bbcc5d0>,  
<matplotlib.lines.Line2D at 0x2389bbcd0d0>],  
'caps': [<matplotlib.lines.Line2D at 0x2389bbcdc10>,  
<matplotlib.lines.Line2D at 0x2389bbcd710>],  
'boxes': [<matplotlib.lines.Line2D at 0x2389bbc38d0>],  
'medians': [<matplotlib.lines.Line2D at 0x2389bbcf110>],  
'fliers': [<matplotlib.lines.Line2D at 0x2389bbc3710>],  
'means': []}
```

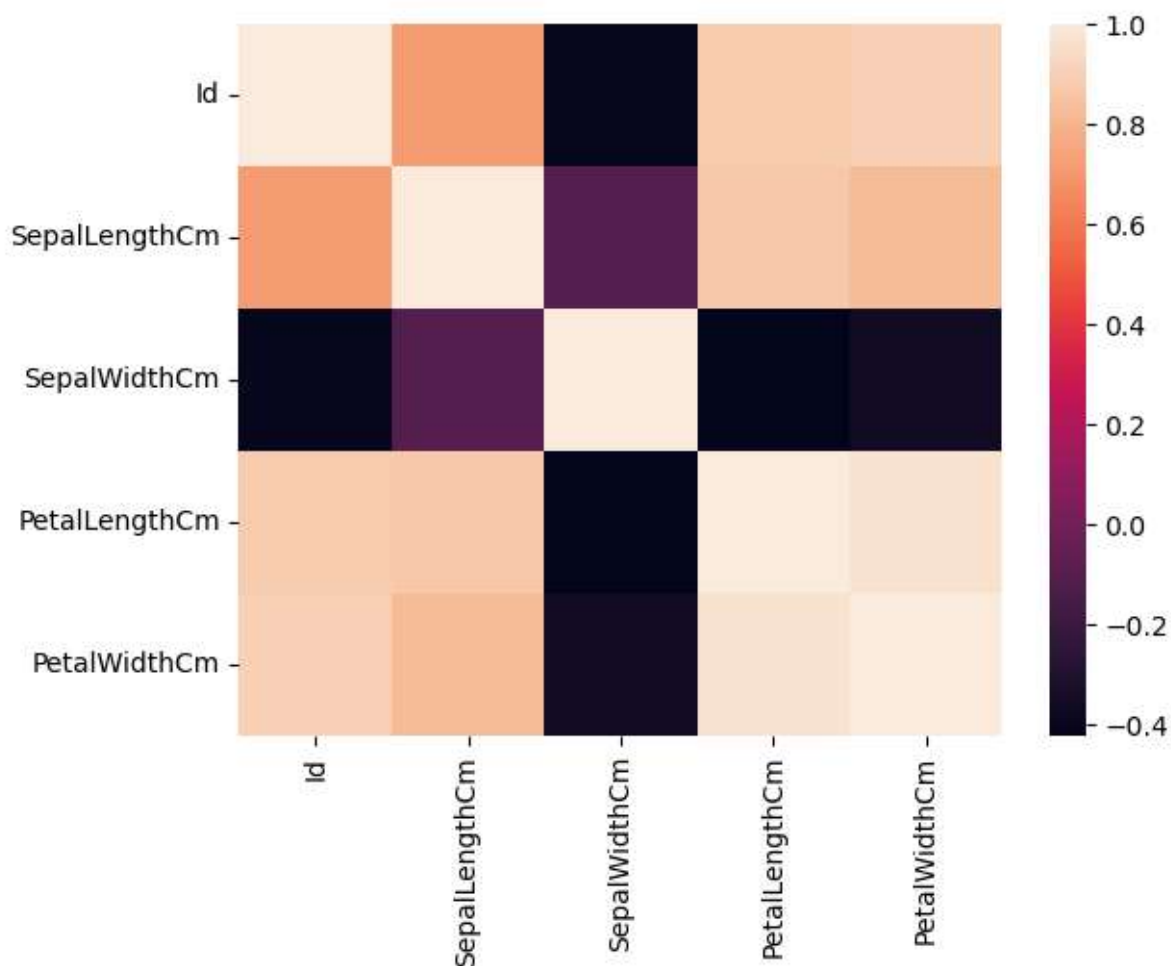


```
In [12]: sns.heatmap(df.corr())
```

C:\Users\91772\AppData\Local\Temp\ipykernel_22852\58359773.py: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.

```
sns.heatmap(df.corr())
```

```
Out[12]: <Axes: >
```



```
In [13]: #3. Data Preparation  
df.drop('Id',axis=1,inplace=True)
```

```
In [14]: sp={'Iris-setosa':1,'Iris-versicolor':2,'Iris-virginica':3}
```

```
In [15]: df.Species=[sp[i] for i in df.Species]
```

```
In [16]: df
```

Out[16]:

	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	5.1	3.5	1.4	0.2	1
1	4.9	3.0	1.4	0.2	1
2	4.7	3.2	1.3	0.2	1
3	4.6	3.1	1.5	0.2	1
4	5.0	3.6	1.4	0.2	1
...
145	6.7	3.0	5.2	2.3	3
146	6.3	2.5	5.0	1.9	3
147	6.5	3.0	5.2	2.0	3
148	6.2	3.4	5.4	2.3	3
149	5.9	3.0	5.1	1.8	3

150 rows × 5 columns

In [17]:

X=df.iloc[:,0:4]

In [19]:

X

Out[19]:

	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
...
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

150 rows × 4 columns

In [20]:

y=df.iloc[:,4]

In [21]:

y

```
Out[21]: 0      1
          1      1
          2      1
          3      1
          4      1
          ..
        145     3
        146     3
        147     3
        148     3
        149     3
        Name: Species, Length: 150, dtype: int64
```

```
In [22]: X_train,X_test,y_train,y_test=train_test_split(X,y,test_size=0.33,random_state=42)
```

```
In [23]: #Traning Model
        model=LinearRegression()
```

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

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

```
In [25]: model.score(X,y) #coef of prediction
```

```
Out[25]: 0.9304223675331595
```

```
In [26]: model.coef_
```

```
Out[26]: array([-0.10974146, -0.04424045,  0.22700138,  0.60989412])
```

```
In [27]: model.intercept_
```

```
Out[27]: 1.192083994828144
```

```
In [28]: #Making Prediction
        y_pred=model.predict(X_test)
```

```
In [29]: #Model Evolution
        print("Mean squared error: %.2f" % np.mean((y_pred - y_test) ** 2))
```

```
Mean squared error: 0.04
```

```
In [ ]:
```