In [47]: import pandas as pd
import numpy as np
import pickle
import warnings
warnings.filterwarnings('ignore')

Out[48]:

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa
145	6.7	3.0	5.2	2.3	virginica
146	6.3	2.5	5.0	1.9	virginica
147	6.5	3.0	5.2	2.0	virginica
148	6.2	3.4	5.4	2.3	virginica
149	5.9	3.0	5.1	1.8	virginica

150 rows × 5 columns

In [49]: data.describe()

Out[49]:

	sepal_length	sepal_width	petal_length	petal_width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.054000	3.758667	1.198667
std	0.828066	0.433594	1.764420	0.763161
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.300000
50%	5.800000	3.000000	4.350000	1.300000
75%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	2.500000

```
In [50]:
          data.info()
          <class 'pandas.core.frame.DataFrame'>
          RangeIndex: 150 entries, 0 to 149
          Data columns (total 5 columns):
                Column
                                Non-Null Count Dtype
                ____
                                -----
                                                  ----
                sepal_length 150 non-null
                                                  float64
           0
            1
                sepal_width
                                150 non-null
                                                  float64
                                                  float64
            2
                petal_length 150 non-null
            3
                petal_width
                                150 non-null
                                                  float64
           4
                species
                                150 non-null
                                                  object
          dtypes: float64(4), object(1)
          memory usage: 6.0+ KB
In [51]:
          data.head()
Out[51]:
              sepal length sepal width petal length petal width species
           0
                      5.1
                                  3.5
                                              1.4
                                                          0.2
                                                               setosa
           1
                      4.9
                                  3.0
                                              1.4
                                                          0.2
                                                               setosa
           2
                      4.7
                                  3.2
                                              1.3
                                                          0.2
                                                               setosa
           3
                                                          0.2
                      4.6
                                  3.1
                                              1.5
                                                               setosa
           4
                      5.0
                                  3.6
                                              1.4
                                                          0.2
                                                               setosa
In [52]:
          data.tail()
Out[52]:
                sepal_length sepal_width petal_length petal_width species
           145
                        6.7
                                    3.0
                                                5.2
                                                            2.3 virginica
                                    2.5
           146
                        6.3
                                                5.0
                                                            1.9
                                                               virginica
           147
                        6.5
                                                5.2
                                                                virginica
                                    3.0
                                                            2.0
           148
                        6.2
                                    3.4
                                                5.4
                                                            2.3
                                                               virginica
           149
                        5.9
                                    3.0
                                                5.1
                                                            1.8 virginica
In [53]:
          data.shape
Out[53]: (150, 5)
In [54]: data['species'].value counts()
Out[54]: species
                          50
          setosa
          versicolor
                          50
```

virginica

50

Name: count, dtype: int64

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	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	0
1	4.9	3.0	1.4	0.2	0
2	4.7	3.2	1.3	0.2	0
3	4.6	3.1	1.5	0.2	0
4	5.0	3.6	1.4	0.2	0
145	6.7	3.0	5.2	2.3	2
146	6.3	2.5	5.0	1.9	2
147	6.5	3.0	5.2	2.0	2
148	6.2	3.4	5.4	2.3	2
149	5.9	3.0	5.1	1.8	2

150 rows × 5 columns

Data Visualization

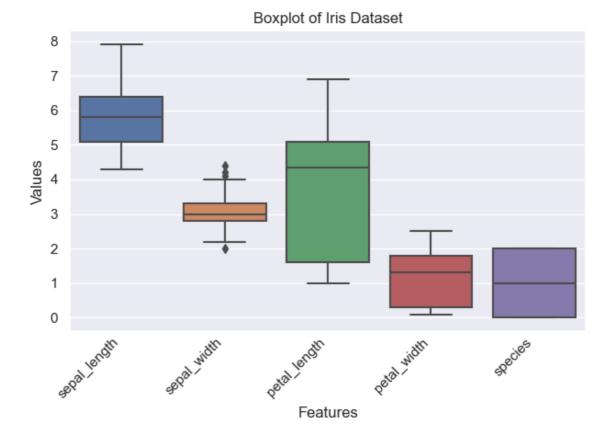
```
In [56]: import seaborn as sns
import matplotlib.pyplot as plt
```

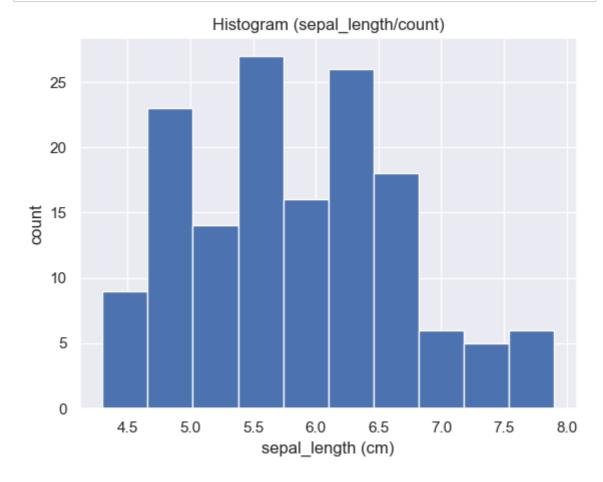
```
In [57]: sns.boxplot(data=data)

plt.tick_params(axis='x', which='both', length=0)
plt.xticks(rotation=45, ha='right')

# Add labels and title
plt.xlabel('Features')
plt.ylabel('Values')
plt.title('Boxplot of Iris Dataset')

# Display the plot
plt.tight_layout() # Ensures labels are not cut off
plt.show()
```





```
x=data.drop(['species'],axis=1)
In [59]:
Out[59]:
                 sepal_length sepal_width petal_length petal_width
              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
                                                     1.4
                                                                  0.2
                           5.0
                                        3.6
            145
                           6.7
                                        3.0
                                                     5.2
                                                                  2.3
                                        2.5
                                                     5.0
            146
                           6.3
                                                                  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 [60]: y=data['species']
Out[60]: 0
                    0
                    0
                    0
           3
                    0
           4
           145
                    2
           146
                    2
           147
                    2
                    2
           148
```

Data Splitting into Training data and Testing data

Name: species, Length: 150, dtype: int64

149

```
Flower Species prediction using Random Forest - Jupyter Notebook
           from sklearn.model_selection import train_test_split
In [61]:
           x_test,x_train,y_test,y_train=train_test_split(x,y,test_size=0.3,random_sta
           x_test.head(10)
Out[61]:
                 sepal_length sepal_width petal_length petal_width
             81
                                       2.4
                          5.5
                                                    3.7
                                                                1.0
            133
                          6.3
                                       2.8
                                                    5.1
                                                                1.5
            137
                          6.4
                                       3.1
                                                    5.5
                                                                1.8
             75
                          6.6
                                       3.0
                                                    4.4
                                                                1.4
                          7.2
                                                    6.1
                                                                2.5
            109
                                       3.6
                          5.7
                                       2.9
                                                    4.2
                                                                1.3
             96
            105
                          7.6
                                       3.0
                                                    6.6
                                                                2.1
                          5.6
                                                    4.5
             66
                                       3.0
                                                                1.5
              0
                          5.1
                                       3.5
                                                    1.4
                                                                0.2
            122
                          7.7
                                       2.8
                                                    6.7
                                                                2.0
In [62]:
           x_train.head()
Out[62]:
                 sepal_length sepal_width petal_length petal_width
             73
                          6.1
                                       2.8
                                                    4.7
                                                                1.2
             18
                          5.7
                                       3.8
                                                    1.7
                                                                0.3
            118
                          7.7
                                       2.6
                                                                2.3
                                                    6.9
             78
                          6.0
                                       2.9
                                                    4.5
                                                                1.5
             76
                          6.8
                                       2.8
                                                    4.8
                                                                1.4
In [63]:
           y_test.head()
Out[63]: 81
                    1
                    2
           133
           137
                    2
                    1
           75
           109
                    2
           Name: species, dtype: int64
In [64]:
           y_train.head()
Out[64]: 73
                    1
           18
                    0
                    2
           118
           78
                    1
           76
                    1
           Name: species, dtype: int64
```

In [65]: x_train.shape

Out[65]: (45, 4)

```
In [66]: x_test.shape
Out[66]: (105, 4)
In [67]: y_train.shape
Out[67]: (45,)
In [68]: y_test.shape
Out[68]: (105,)
```

Random Forest

```
In [69]: #from sklearn.model_selection import GridSearchCV #GridSearchCV is for para
from sklearn.ensemble import RandomForestRegressor
reg=RandomForestRegressor(random_state=0)
reg.fit(x_train,y_train)
```

Out[69]: RandomForestRegressor(random_state=0)

In a Jupyter environment, please rerun this cell to show the HTML representation or trust the notebook.

On GitHub, the HTML representation is unable to render, please try loading this page with nbviewer.org.

Accuracy for Random Forest

```
In [71]: from sklearn.metrics import r2_score
    r2_score(y_test,y_pred)
```

Out[71]: 0.9481385135135135

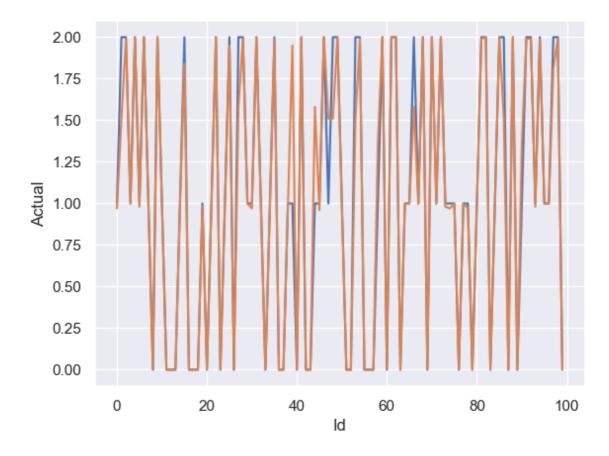
```
In [72]: Results= pd.DataFrame(columns=['Actual','Predicted'])
Results['Actual']=y_test
Results['Predicted']=y_pred
#Results['km']=X_test['km']
Results=Results.reset_index()
Results['Id']=Results.index
Results.head(10)
```

Out[72]:

	index	Actual	Predicted	ld
0	81	1	0.97	0
1	133	2	1.51	1
2	137	2	1.99	2
3	75	1	1.00	3
4	109	2	2.00	4
5	96	1	0.98	5
6	105	2	2.00	6
7	66	1	0.98	7
8	0	0	0.00	8
9	122	2	2.00	9

```
import seaborn as sns
import matplotlib.pyplot as plt
sns.lineplot(x='Id',y='Actual',data=Results.head(100))
sns.lineplot(x='Id',y='Predicted',data=Results.head(100))
plt.plot()
```

Out[73]: []



```
In [ ]:

In [ ]:
```