

```
import numpy as np
import pandas as pd
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn import svm
from sklearn import metrics
import matplotlib.pyplot as plt
import seaborn as sns
import itertools
from sklearn.linear_model import LinearRegression
```

```
:Number of Instances: 150 (50 in each of three classes)
:Number of Attributes: 4 numeric, predictive attributes and the class
:Attribute Information:
  - sepal length in cm
  - sepal width in cm
  - petal length in cm
  - petal width in cm
  - class:
    - Iris-Setosa
    - Iris-Versicolour
    - Iris-Virginica
```

```
['setosa' 'versicolor' 'virginica']
```

[illegible]

[5.2	4.1	1.5	0.1]
[5.5	4.2	1.4	0.2]
[4.9	3.1	1.5	0.2]
[5.	3.2	1.2	0.2]
[5.5	3.5	1.3	0.2]
[4.9	3.6	1.4	0.1]
[4.4	3.	1.3	0.2]
[5.1	3.4	1.5	0.2]
[5.	3.5	1.3	0.3]
[4.5	2.3	1.3	0.3]
[4.4	3.2	1.3	0.2]
[5.	3.5	1.6	0.6]
[5.1	3.8	1.9	0.4]
[4.8	3.	1.4	0.3]
[5.1	3.8	1.6	0.2]
[4.6	3.2	1.4	0.2]
[5.3	3.7	1.5	0.2]
[5.	3.3	1.4	0.2]
[7.	3.2	4.7	1.4]
[6.4	3.2	4.5	1.5]

```
['sepal length (cm)', 'sepal width (cm)', 'petal length (cm)', 'petal width (cm)']
```

[illegible]

```
iris_df = pd.DataFrame(data= data.data, columns= data.feature_names)
target_df = pd.DataFrame(data= data.target, columns= ['species'])
def converter(specie):
    if specie == 0:
        return 'setosa'
    elif specie == 1:
        return 'versicolor'
    else:
        return 'virginica'
target_df['species'] = target_df['species'].apply(converter)
# Concatenate the DataFrames
iris_df = pd.concat([iris_df, target_df], axis= 1)
```

```
target_df = pd.DataFrame(columns= ['species'], data= data.target)
iris_df = pd.concat([iris_df, target_df], axis= 1)
```

```
# Variables
X= iris_df.drop(labels = ['sepal length (cm)', 'petal length (cm)', 'petal width (cm)', 'species'], axis= 1)
y= iris_df['sepal length (cm)']
print(X)
print(y)

# Splitting the Dataset
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size= 0.33, random_state= 101)
```

```

    sepal width (cm)
0          3.5
1          3.0
2          3.2
3          3.1
4          3.6
..         ...
145        3.0
146        2.5
147        3.0
148        3.4
149        3.0

[150 rows x 1 columns]
0          5.1
1          4.9
2          4.7
3          4.6
4          5.0
..         ...
145        6.7
146        6.3
147        6.5
148        6.2
149        5.9
Name: sepal length (cm), Length: 150, dtype: float64
```

```
# Instantiating LinearRegression() Model
lr = LinearRegression()

# Training/Fitting the Model
lr.fit(X_train, y_train)
```

```
▼ LinearRegression
LinearRegression()
```

In [93]:

```
print(lr.score(X_test, y_test))
# The coefficients
print('Coefficients: \n', lr.coef_)
```

0.006469855487622134

```
Coefficients:
[-0.22561002]
```

In [50]:

```
# Making Predictions
lr.predict(X_test)
pred = lr.predict(X_test)
print(X_test[:5])
print(pred[:5])
print(y_test[:5])
```

```
      sepal width (cm)
33          4.2
16          3.9
43          3.5
129         3.0
50          3.2
[5.61492654  5.68260954  5.77285355  5.88565856  5.84053656]
33          5.5
16          5.4
43          5.0
129         7.2
50          7.0
Name: sepal length (cm), dtype: float64
```

/var/folders/kb/2qtwss7n3y3091dclgrcn9hm0000gn/T/ipykernel_97907/2836971499.py:6: FutureWarning: The behavior of `series[i:j]` with an integer-dtype index is deprecated. In a future version, this will be treated as *label-based* indexing, consistent with e.g. `series[i]` lookups. To retain the old behavior, use `series.iloc[i:j]`. To get the future behavior, use `series.loc[i:j]`.

```
print(y_test[:5])
```

In [51]:

```
# Evaluating Model's Performance
print('Mean Absolute Error:', metrics.mean_absolute_error(y_test, pred))
print('Mean Squared Error:', metrics.mean_squared_error(y_test, pred))
print('Mean Root Squared Error:', np.sqrt(metrics.mean_squared_error(y_test, pred)))
```

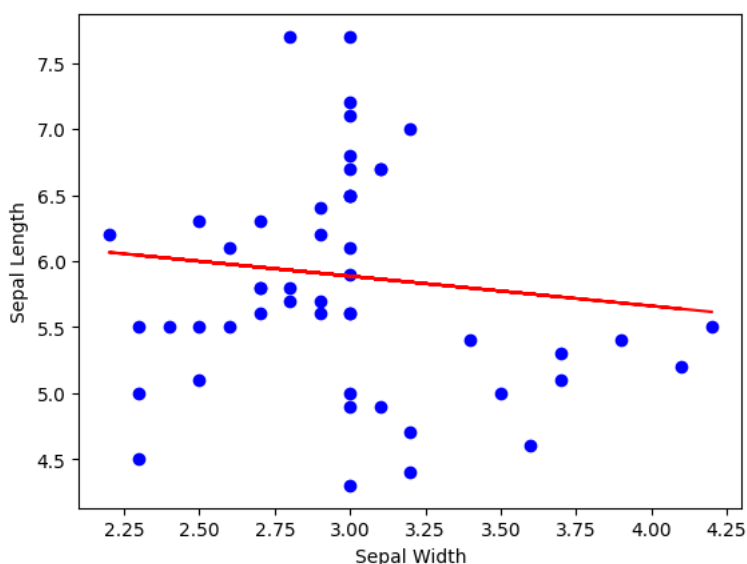
Mean Absolute Error: 0.6655951111688988

Mean Squared Error: 0.6598987478645434

Mean Root Squared Error: 0.8123415216917964

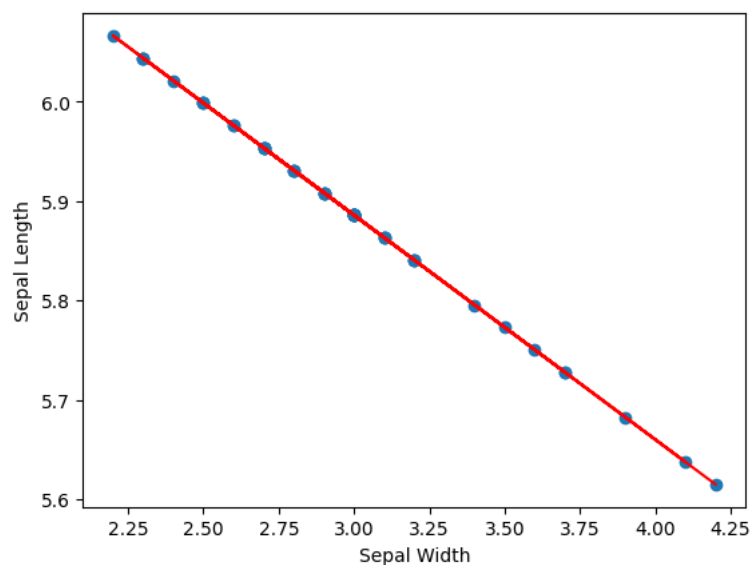
In [94]:

```
plt.scatter(X_test,y_test, color = 'b')
plt.plot(X_test,lr.predict(X_test),color = 'r')
plt.xlabel("Sepal Width")
plt.ylabel("Sepal Length")
plt.show()
```



In [104]:

```
plt.scatter(X_test,pred)
plt.plot(X_test,lr.predict(X_test),color = 'r')
plt.xlabel("Sepal Width")
plt.ylabel("Sepal Length")
plt.show()
```



In [90]:

```
#predicting values
d = {
    'sepal width (cm)' : [5.3]
}
testing = pd.DataFrame(data = d)
y_predicted_value = lr.predict(testing)
print(y_predicted_value)
```

[5.36675552]

In [105]:

```
#Coefficient of determination
r_squared = lr.score(X,y)
print(r_squared)

#slope
slope = lr.coef_
print(slope)

#intercept
intercept = lr.intercept_
print(intercept)

#SSR(sum of squared residuals)
residuals = y_test - pred

SSR = np.sum(residuals**2)
print(SSR)
```

```
0.012553070673583133
[-0.22561002]
6.562488623065922
32.99493739322717
```

In []: