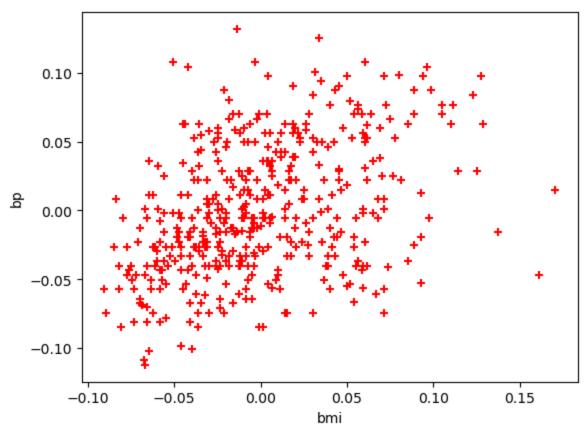
Experiment 8: Implement Decision Tree by using Diabetes Dataset

```
In [4]: import pandas as pd
       import numpy as np
       import matplotlib.pyplot as plt
       from sklearn.datasets import load diabetes
       diabetes = load diabetes()
In [5]: diabetes.feature names
       #diabetes.DESCR
Out[5]: ['aqe', 'sex', 'bmi', 'bp', 's1', 's2', 's3', 's4', 's5', 's6']
In [6]: df = pd.DataFrame(diabetes.data, columns=diabetes.feature names)
       df.head()
Out[6]:
               age
                         sex
                                  bmi
                                            bp
                                                      s1
                                                                s2
                                                                         s3
        0 0.038076
                    0.050680 0.061696
                                      0.021872 -0.044223 -0.034821 -0.043401 -0
        1 -0.001882 -0.044642 -0.051474 -0.026328 -0.008449 -0.019163
                                                                   0.074412 -0
        2 0.085299 0.050680 0.044451 -0.005670 -0.045599 -0.034194 -0.032356 -0
        3 -0.089063 -0.044642 -0.011595 -0.036656
                                                 0.012191 0.024991 -0.036038
        4 0.005383 -0.044642 -0.036385 0.021872 0.003935 0.015596 0.008142 -0
In [7]: df['target'] = diabetes.target
       df.head()
Out[7]:
                                  bmi
                                            bp
                                                      s1
                                                                s2
                                                                         s3
               age
                         sex
        0 0.038076
                    1 -0.001882 -0.044642 -0.051474 -0.026328 -0.008449 -0.019163 0.074412 -0
       2 0.085299 0.050680 0.044451 -0.005670 -0.045599 -0.034194 -0.032356 -0
       3 -0.089063 -0.044642 -0.011595 -0.036656 0.012191 0.024991 -0.036038
                                                                             0
          0.005383 -0.044642 -0.036385 0.021872 0.003935 0.015596 0.008142 -0
In [8]: diabetes.target filename
Out[8]: 'diabetes target.csv.gz'
In [9]:
       plt.xlabel('bmi')
       plt.ylabel('bp')
       plt.scatter(df['bmi'],df['bp'],color='red', marker='+')
```

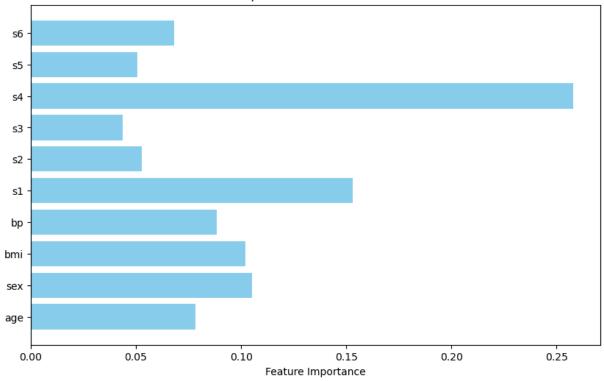


In [17]: model.score(x_test,y_test)

Out[17]: 0.46798286239277226

```
In [18]:
        model.predict(x test[:6])
Out[18]: array([[-0.02004471, -0.04464164,
                                          0.00457217,
                                                       0.09761511,
                                                                   0.0053108 ,
                              0.06336665, -0.03949338,
                 -0.02072908,
                                                                   0.01134862],
                                                       0.01255119,
                [ 0.02717829, 0.05068012, 0.09403057,
                                                       0.09761511, -0.03459183,
                 -0.03200243, -0.04340085, -0.00259226,
                                                       0.03664373, 0.10661708],
                [-0.00914709, 0.05068012, -0.03961813, -0.04009893, -0.00844872,
                                                       0.01776532, -0.067351411,
                  0.01622244, -0.06549067, 0.07120998,
                [ 0.05987114, -0.04464164, -0.00081689, -0.08485599,
                                                                   0.0754844 ,
                  0.07947843, 0.00446045, 0.03430886,
                                                       0.02337142,
                                                                   0.02791705],
                [-0.07453279, -0.04464164, 0.0433734, -0.03321323,
                                                                   0.01219057,
                  0.00025186, 0.06336665, -0.03949338, -0.02712902, -0.04664087],
                [-0.00551455, 0.05068012, -0.00836158, -0.00222757, -0.03321588,
                 -0.06363042, -0.03603757, -0.00259226, 0.08059005, 0.00720652]])
In [19]:
        x test[:6]
Out[19]:
                   age
                                      bmi
                                                 bp
                                                           s1
                                                                    s2
                                                                              s3
                            sex
         343 -0.020045 -0.044642
                                 0.018584
                                           0.090729
                                                     0.003935
                                                               0.008707
                                                                        0.037595
         163
              0.016281 0.050680 0.072474
                                           0.076958 -0.008449
                                                               0.005575 -0.006584
          0.012152 -0.072854
         282
              0.070769 -0.044642 -0.005128 -0.005670 0.087868
                                                               0.102965
                                                                        0.011824
         143 -0.030942 -0.044642 0.005650 -0.009113 0.019070 0.006828
                                                                        0.074412
         275 -0.005515
                        0.050680 -0.011595
                                           0.011544 -0.022208 -0.015406 -0.021311
In [20]: # Feature Importance
         importances = model.feature importances
         plt.figure(figsize=(10, 6))
         plt.barh(df.columns[:-1], importances, color="skyblue")
         plt.xlabel("Feature Importance")
         plt.title("Feature Importance in Decision Tree Model")
         plt.show()
```

Feature Importance in Decision Tree Model



```
In [40]: from sklearn.metrics import mean_absolute_error, mean_squared_error
import numpy as np

# Predictions with the best model
y_pred = best_model.predict(x_test)

# Evaluation metrics
mae = mean_absolute_error(y_test, y_pred)
rmse = np.sqrt(mean_squared_error(y_test, y_pred))
r2_score = best_model.score(x_test, y_test)

print("Mean Absolute Error (MAE):", mae)
print("Root Mean Squared Error (RMSE):", rmse)
print("R^2 Score:", r2_score)
```

Mean Absolute Error (MAE): 0.021066464678941133 Root Mean Squared Error (RMSE): 0.02900287696752611 R^2 Score: 0.6095040401103815

This notebook was converted with convert.ploomber.io