This is the boston housing Dataset. Without statistical training only an R^2 of 70% can be achieved. I've managed to acheive R^2 of 91%!!

Some of the things I did was to relfected the skews to be right skew. Then I scaled the data. Then I fulled in NAs with mean. Then I removed outliers. Then I applied transformation to tame the variance skewness.

Lastly, I applied PCA to get a features that capture most variance

After doing all of this I got R^2 of 91% for Linear regression and 91% for regression trees

```
In [569...
          import pandas as pd
          from sklearn.preprocessing import PolynomialFeatures
          import matplotlib.pyplot as plt
          from sklearn.model_selection import train_test_split
          from sklearn.ensemble import RandomForestRegressor
          from sklearn.metrics import mean_squared_error, r2_score
          from sklearn.preprocessing import MinMaxScaler
          from scipy import stats
          from sklearn.model_selection import cross_val_score
          from sklearn.metrics import make_scorer, mean_squared_error
          import numpy as np
          df = pd.read_csv("housing.csv")
          saved = df.copy()
          colnam = df.columns
          skewness = df.skew()
          left_tailed_columns = skewness.index[skewness < 0]</pre>
          df[left_tailed_columns] = -df[left_tailed_columns]
          scaler = MinMaxScaler((1, 4))
          df = pd.DataFrame(scaler.fit_transform(df), columns=colnam)
          for column in df.columns:
              mean_value = df[column].mean()
              df[column].fillna(mean_value, inplace=True)
          ## Filter out outliers
          import numpy as np
          import pandas as pd
          from scipy import stats
          # Assuming df is your DataFrame
          def filter_outliers_zscore(dataframe, threshold):
              z_scores = np.abs(stats.zscore(dataframe))
              filtered_data = dataframe[(z_scores < threshold).all(axis=1)]</pre>
              return filtered_data
          # Set the z-score threshold
          zscore_threshold = 3.5
          df = filter_outliers_zscore(df, zscore_threshold)
          skewness = df.skew()
          high_skew_columns = skewness.index[skewness > 2.2]
          for column in high_skew_columns:
              df[column] = np.reciprocal(df[column])
          df = df.dropna()
          y = df.MEDV
          from sklearn.decomposition import PCA
```

```
n_components = 5
pca = PCA(n_components=n_components)
df_pca = pca.fit_transform(df)
```

# Concatenate the PCA components with the target variable 'y'
import statsmodels.api as sm
df\_pca = pd.DataFrame(df\_pca, columns=[f'PC{i+1}' for i in range(n\_components)])
df\_pca.head(4)

Out[570...

0

	PC1	PC2	PC3	PC4	PC5
0	-1.542732	0.546593	0.637931	0.222229	-0.407906
1	-1.205496	0.673129	-0.220644	-0.291848	-0.339355
2	-1.674761	0.216983	0.414124	-0.635204	0.256809
3	-2.028978	-0.352560	0.127839	-0.677374	0.024618

```
In [571...
          df_pca.hist()
          array([[<Axes: title={'center': 'PC1'}>, <Axes: title={'center': 'PC2'}>],
Out[571...
                  [<Axes: title={'center': 'PC3'}>, <Axes: title={'center': 'PC4'}>],
                  [<Axes: title={'center': 'PC5'}>, <Axes: >]], dtype=object)
                            PC1
                                                                    PC2
                                                  100
           50
                                                   50
                                                    0
                  -2
                            ₽C3
                                     2
                                                              -1
                                                                    RC4
                                                                                  2
          100
                                                  100
           50
                                                   50
            0
                                                    0
                          0 PC51
          100
           50
```

## Below it shows that 90% of the information is captured by these 5 new features

```
explained_variance_ratio = pca.explained_variance_ratio_
    cumulative_explained_variance = explained_variance_ratio.cumsum()

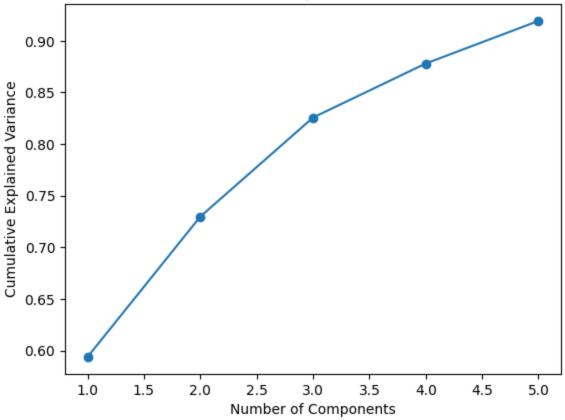
print("Explained Variance Ratios:")
    print(explained_variance_ratio)

plt.plot(range(1, n_components + 1), cumulative_explained_variance, marker='o')
    plt.title('Cumulative Explained Variance')
    plt.xlabel('Number of Components')
    plt.ylabel('Cumulative Explained Variance')
    plt.show()
```

Explained Variance Ratios: [0.5940003 0.13582333 0.09590953 0.05231835 0.04123233]

0

## **Cumulative Explained Variance**



```
from sklearn.linear_model import LinearRegression
In [573...
                              from sklearn.metrics import r2_score
                              from sklearn.model_selection import train_test_split
                              # Assuming X_train, X_test, y_train, y_test are already defined
                              df_pca_with_const = sm.add_constant(df_pca)
                              df_pca_with_const
                              # Number of iterations
                              num_iterations = 6
                              # Initialize the linear regression model
                              linear_reg_model = LinearRegression()
                              for i in range(num_iterations):
                                         # Split the data into training and testing sets for each iteration
                                         X_train, X_test, y_train, y_test = train_test_split(df_pca_with_const, y, test_size=0.20, rain_value = train_test_split(df_pca_with_const, y, 
                                         # Train the model
                                         linear_reg_model.fit(X_train, y_train)
                                         # Make predictions on the test set
                                         y_pred = linear_reg_model.predict(X_test)
                                         # Calculate R-squared
                                         r2 = r2_score(y_test, y_pred)
                                         # Calculate adjusted R-squared
                                         n = X_test.shape[0] # Number of samples
                                         p = X_test.shape[1] # Number of features
                                         adjusted_r2 = 1 - (1 - r2) * ((n - 1) / (n - p - 1))
                                         # Display the adjusted R-squared for each iteration
                                         print(f'Iteration {i + 1}: Adjusted R-squared: {adjusted_r2:.4f}')
```

```
Iteration 1: Adjusted R-squared: 0.8875
Iteration 2: Adjusted R-squared: 0.9137
Iteration 3: Adjusted R-squared: 0.8613
Iteration 4: Adjusted R-squared: 0.9161
Iteration 5: Adjusted R-squared: 0.9060
Iteration 6: Adjusted R-squared: 0.8842
```

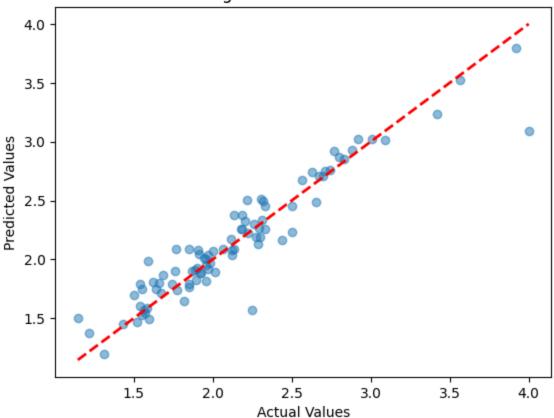
## Standardized MSE = 0.1079

```
import matplotlib.pyplot as plt
from sklearn.preprocessing import StandardScaler
# Compute standardized MSE
mse_std = (mean_squared_error(y_test, y_pred))/(np.var(y_test))
print(f'Standardized Mean Squared Error: {mse_std:.4f}')
```

```
plt.scatter(y_test, y_pred, alpha=0.5)
plt.plot([min(y_test), max(y_test)], [min(y_test), max(y_test)], color='red', linestyle='--', lwi
plt.title('Linear Regression: Actual vs Predicted')
plt.xlabel('Actual Values')
plt.ylabel('Predicted Values')
plt.show()
```

Standardized Mean Squared Error: 0.1079

## Linear Regression: Actual vs Predicted



```
In [575...
from sklearn.ensemble import RandomForestRegressor
from sklearn.metrics import mean_squared_error, r2_score
from sklearn.model_selection import train_test_split

# Assuming you have X_train, X_test, y_train, y_test defined

X_train, X_test, y_train, y_test = train_test_split(df_pca, y, test_size=0.20, random_state=3)

# Initialize the Random Forest Regressor with default parameters
random_forest_model = RandomForestRegressor()
random_forest_model.fit(X_train, y_train)
y_pred_rf = random_forest_model.predict(X_test)
mse_rf = mean_squared_error(y_test, y_pred_rf)
r2_rf = r2_score(y_test, y_pred_rf)
print(f'Random Forest Mean Squared Error: {mse_rf:.4f}')
print(f'Random Forest R-squared: {r2_rf:.4f}')
```

Random Forest Mean Squared Error: 0.0270 Random Forest R-squared: 0.9114

```
In [576...
          import matplotlib.pyplot as plt
          import numpy as np
          # Scatter plot for actual vs predicted values with diagonal line
          plt.scatter(y_test, y_pred_rf, alpha=0.5)
          plt.plot([min(y_test), max(y_test)], [min(y_test), max(y_test)], color='red', linestyle='--', lw
          plt.title('Random Forest: Actual vs Predicted')
          plt.xlabel('Actual Values')
          plt.ylabel('Predicted Values')
          plt.show()
          # Residual plot
          residuals = y_test - y_pred_rf
          plt.scatter(y_pred_rf, residuals, alpha=0.5)
          plt.title('Random Forest: Residual Plot')
          plt.xlabel('Predicted Values')
          plt.ylabel('Residuals')
          plt.axhline(y=0, color='r', linestyle='--')
          plt.show()
```

