```
In [1]: import numpy
        import pandas as pd
        import matplotlib.pyplot as plt
        import seaborn as sns
        from scipy.stats.mstats import normaltest
```

In [3]: data = pd.read_csv('../data/california_housing_pricing.csv') data.head()

Out[3]: longitude latitude housing_median_age total_rooms total_bedrooms population households 0 -122.23 37.88 41.0 880.0 126.0 129.0 322.0 -122.22 37.86 7099.0 2401.0 1 21.0 1106.0 1138.0 2 -122.24 37.85 52.0 1467.0 190.0 496.0 177.0 3 -122.25 37.85 52.0 558.0 1274.0 235.0 219.0 -122.25 37.85 52.0 280.0 565.0 259.0

data_dictionary = pd.read_csv('../data/data_dictionary.csv') In [4]: data_dictionary

1627.0

Out[4]:		Field	Meaning
	0	longitude	A measure of how far west a house is; a highe
	1	latitude	A measure of how far north a house is; a high
	2	housing_median_age	Median age of a house within a block; a lower
	3	total_rooms	Total number of rooms within a block
	4	total_bedrooms	Total number of bedrooms within a block
	5	population	Total number of people residing within a block
	6	households	Total number of households, a group of people
	7	median_income)	Median income for households within a block o
	8	median_house_value)	Median house value for households within a bl
	9	ocean_proximityea	Location of the house w.r.t ocean/s

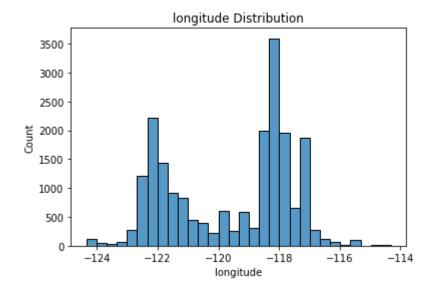
```
In [5]: data.info()
        <class 'pandas.core.frame.DataFrame'>
        RangeIndex: 20640 entries, 0 to 20639
        Data columns (total 10 columns):
         #
             Column
                                 Non-Null Count Dtype
             ----
                                 -----
        - - -
                                                 ----
             longitude
         0
                                 20640 non-null float64
             latitude
                                 20640 non-null float64
         1
         2
             housing_median_age 20640 non-null float64
         3
             total rooms
                                 20640 non-null float64
             total_bedrooms
         4
                                 20433 non-null float64
         5
             population
                                 20640 non-null float64
         6
             households
                                 20640 non-null float64
         7
             median income
                                 20640 non-null float64
         8
             median_house_value 20640 non-null float64
         9
             ocean proximity
                                 20640 non-null object
        dtypes: float64(9), object(1)
        memory usage: 1.6+ MB
In [6]: | data.total_bedrooms = data.total_bedrooms.fillna(data.total_bedrooms.median())
In [7]: numeric cols = list(data.select dtypes([int, float]).columns)
        numeric cols
Out[7]: ['longitude',
         'latitude',
         'housing median age',
         'total_rooms',
         'total bedrooms',
          'population',
         'households',
         'median income',
         'median house value']
```

EDA

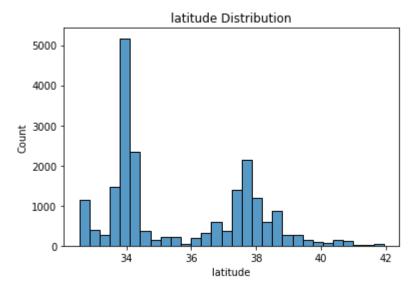
```
In [8]: | normaltest(data.median_house_value.values)
Out[8]: NormaltestResult(statistic=2430.931051066072, pvalue=0.0)
```

```
In [9]: | for col in numeric_cols:
          _, pvalue = normaltest(data[col].values)
          if pvalue < 0.05:</pre>
             result = 'not normally distributed'
          else:
             result = 'normally distributed'
          plt.figure(figsize=(6,4))
          sns.histplot(data=data, x = col, bins=30)
          plt.title(f'{col} Distribution')
          plt.show()
```

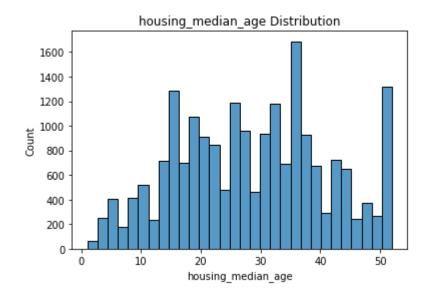
Normality Test P-value: 0.0, Decision: not normally distributed



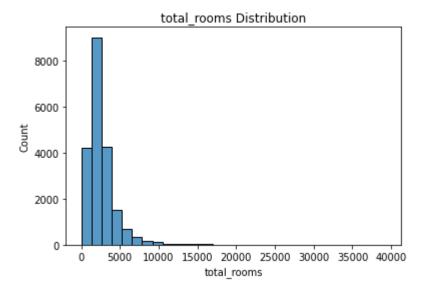
Normality Test P-value: 0.0 , Decision: not normally distributed



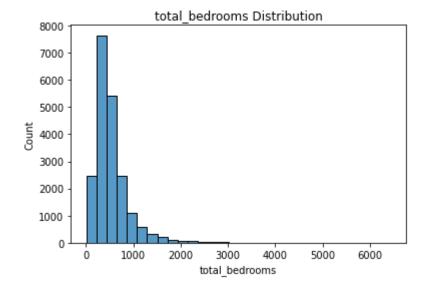
Normality Test P-value: 0.0 , Decision: not normally distributed



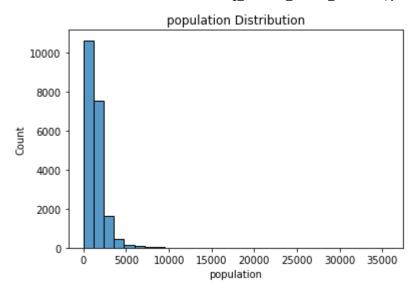
Normality Test P-value: 0.0 , Decision: not normally distributed



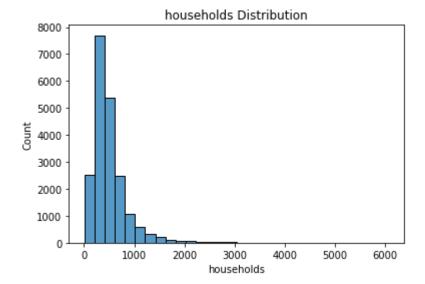
Normality Test P-value: 0.0 , Decision: not normally distributed



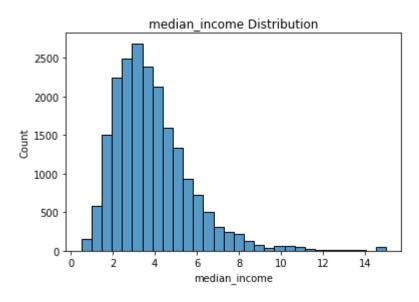
Normality Test P-value: 0.0 , Decision: not normally distributed



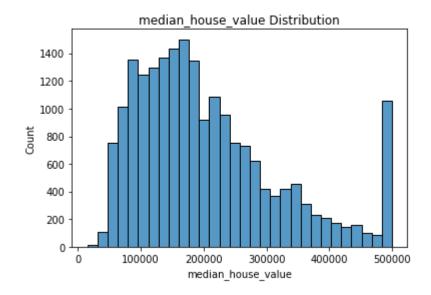
Normality Test P-value: 0.0, Decision: not normally distributed



Normality Test P-value: 0.0, Decision: not normally distributed



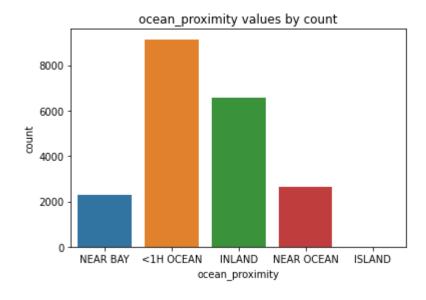
Normality Test P-value: 0.0 , Decision: not normally distributed



The numeric columns including the targte are not normally distributed

```
sns.countplot(x = data.ocean_proximity)
In [10]:
         plt.title('ocean proximity values by count');
```

Out[10]: Text(0.5, 1.0, 'ocean_proximity values by count')



Features Selection

In this section, I'll conduct statistical stest to determine which features have significant relationships with the target in order to select them for the model.

FOr the numeric features, I'll use correlation test, and for the categorical features, I'll use ANOVA test.

```
import scipy.stats
In [11]:
```

Testing the significance of the numeric features

```
In [12]: numeric_cols_stats = [col for col in numeric_cols if col != 'median_house_value
```

```
In [13]: numeric_cols_stats
Out[13]: ['longitude',
           'latitude',
           'housing_median_age',
           'total_rooms',
           'total_bedrooms',
           'population',
           'households',
           'median_income']
```

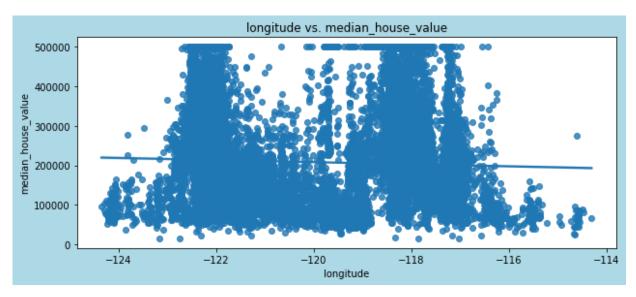
```
In [14]: significant features = []
         for col in numeric cols stats:
             corr, pvalue = scipy.stats.pearsonr(data[col],data['median_house_value'])
             print(f'{col} vs. median house value Results:\n')
             print('Correlation Coefficient:',corr)
             print('p_value:',pvalue)
             if pvalue < 0.05:</pre>
                  significant_features.append(col)
                 if corr > 0:
                      print(f'There is a significant evidence of a positive relationship be
                 else:
                      print(f'There is a significant evidence of a negative relationship be
             else:
                 print(f'There is no significant relationship between {col} and Median Hol
             plt.figure(figsize = (10,4), facecolor='lightblue')
             plt.title(f'{col} vs. median house value')
             sns.regplot(x=data[col], y=data['median_house_value'],data=data)
             plt.show()
```

longitude vs. median_house_value Results:

Correlation Coefficient: -0.04596661511797848

p value: 3.923322071106899e-11

There is a significant evidence of a negative relationship between longitude an d Median House Value

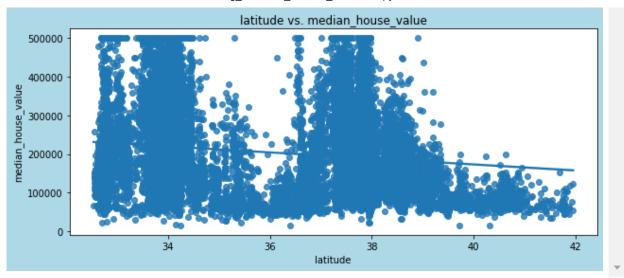


latitude vs. median house value Results:

Correlation Coefficient: -0.1441602768746593

p value: 2.9398592907424878e-96

There is a significant evidence of a negative relationship between latitude and Median House Value

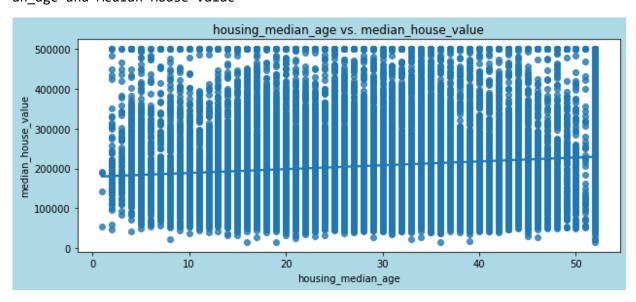


housing_median_age vs. median_house_value Results:

Correlation Coefficient: 0.10562341249320993

p_value: 2.7618606761502365e-52

There is a significant evidence of a positive relationship between housing_medi an age and Median House Value



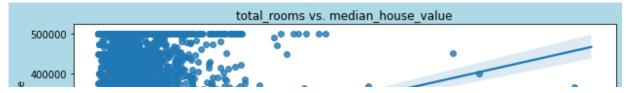
total_rooms vs. median_house_value Results:

Correlation Coefficient: 0.1341531138065631

p_value: 1.6893845634754333e-83

There is a significant evidence of a positive relationship between total_rooms

and Median House Value

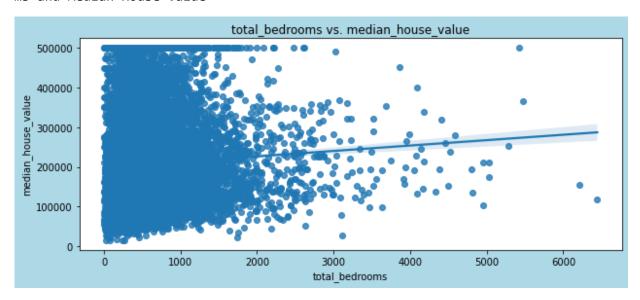


total_bedrooms vs. median_house_value Results:

Correlation Coefficient: 0.049456861920854564

p value: 1.1671461983715622e-12

There is a significant evidence of a positive relationship between total bedroo ms and Median House Value

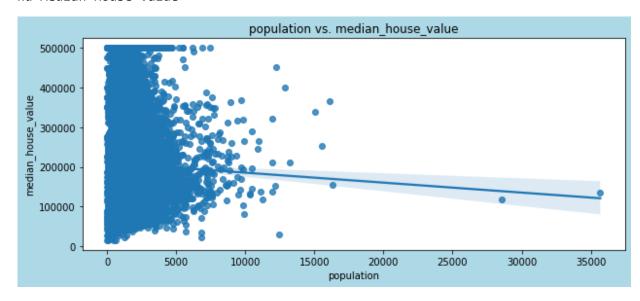


population vs. median house value Results:

Correlation Coefficient: -0.02464967888889489

p_value: 0.0003976307847911049

There is a significant evidence of a negative relationship between population a nd Median House Value

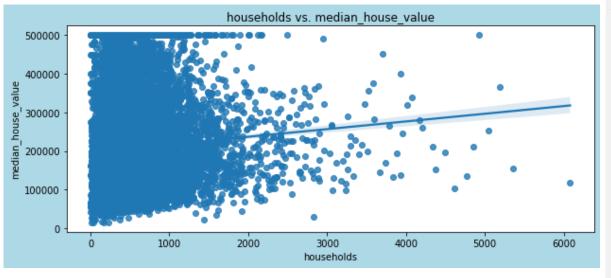


households vs. median house value Results:

Correlation Coefficient: 0.06584265057005645

p_value: 2.8234206519537093e-21

There is a significant evidence of a positive relationship between households a

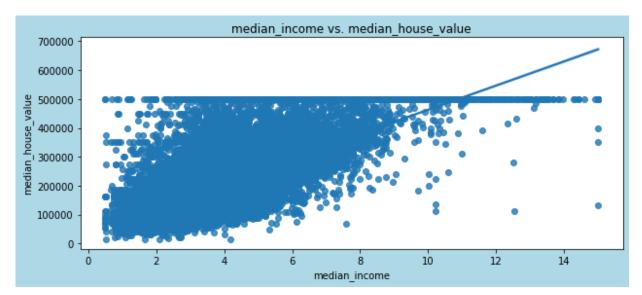


median_income vs. median_house_value Results:

Correlation Coefficient: 0.6880752079585479

p_value: 0.0

There is a significant evidence of a positive relationship between median_incom e and Median House Value



```
In [15]: | significant_features
Out[15]: ['longitude',
           'latitude',
           'housing_median_age',
           'total_rooms',
           'total_bedrooms',
           'population',
           'households',
           'median_income']
```

```
In [16]: data.columns
Out[16]: Index(['longitude', 'latitude', 'housing median age', 'total rooms',
                  'total_bedrooms', 'population', 'households', 'median_income',
                 'median_house_value', 'ocean_proximity'],
                dtype='object')
In [17]: list(data.ocean proximity.value counts().index)
Out[17]: ['<1H OCEAN', 'INLAND', 'NEAR OCEAN', 'NEAR BAY', 'ISLAND']
          Testing the significance of the categorical feature
          State the hypothesis
           • H_0: \mu_1 = \mu_2 = \mu_3 = \mu_4 = \mu_5 (the population means are equal)
            • H_1: At least one of the means differ
In [18]: _, p_value = scipy.stats.f_oneway(data[data['ocean_proximity'] == '<1H OCEAN']['r</pre>
                                              data[data['ocean proximity'] == 'INLAND']['medi
                                              data[data['ocean_proximity'] == 'NEAR OCEAN'][
                                              data[data['ocean_proximity'] == 'NEAR BAY']['me
                                              data[data['ocean proximity'] == 'ISLAND']['medi
          print('P-values: ', str(p_value))
          if p value < 0.05:
              print('There is a statistical evidence that ocean proximity has a significant
              significant_features.append('ocean_proximity')
          else:
                  print('There is no statistical evidence that ocean proximity has a signif
          P-values: 0.0
          There is a statistical evidence that ocean proximity has a significant relation
          ship with median house value
In [19]: | significant features
Out[19]: ['longitude',
           'latitude',
           'housing median age',
           'total_rooms',
           'total bedrooms',
           'population',
           'households',
           'median_income',
           'ocean proximity']
```

As shown from the statistical analysis, ALL the features have significant relationship with the target.

Features Engineering

In this section, I'll use pandas to transfrom the categorical features "ocean proximity" to multiple features each for each distinct value in the original "ocean proximity" feature.

```
In [20]: # from sklearn.preprocessing import OneHotEncoder, LabelEncoder
In [21]: | df = data.copy()
In [22]: ocean proximity ohc = pd.get dummies(df['ocean proximity'], drop first=True)
In [23]: |df.drop('ocean_proximity', axis=1, inplace=True)
In [24]: | df = pd.concat([df, ocean_proximity_ohc], axis =1)
          df.head()
Out[24]:
              longitude latitude housing_median_age total_rooms total_bedrooms population households
           0
                -122.23
                          37.88
                                               41.0
                                                          880.0
                                                                                    322.0
                                                                                                126.0
                                                                         129.0
           1
                -122.22
                          37.86
                                               21.0
                                                         7099.0
                                                                        1106.0
                                                                                   2401.0
                                                                                               1138.0
           2
                -122.24
                          37.85
                                               52.0
                                                         1467.0
                                                                         190.0
                                                                                    496.0
                                                                                                177.0
           3
                -122.25
                          37.85
                                               52.0
                                                         1274.0
                                                                         235.0
                                                                                    558.0
                                                                                                219.0
                -122.25
                          37.85
                                               52.0
                                                         1627.0
                                                                         280.0
                                                                                    565.0
                                                                                                259.0
          Selecting the features and target
In [25]: # select the features
          X = df.drop('median house value',axis=1)
          # select the target
          y = df.median house value
          Although it is not necessary to normalize the targte varibale, sometimes it may lead to better
          model.
```

Spliting the Data and Buidling the Model

In [26]: # normalize the target (as much as possible)

y = numpy.log1p(y)

```
In [61]: from sklearn.preprocessing import StandardScaler, PolynomialFeatures
         from sklearn.model selection import KFold , cross val predict
         from sklearn.linear model import LinearRegression, Lasso, Ridge
         from sklearn.linear model import ElasticNet
         from sklearn.pipeline import Pipeline
         from sklearn.model_selection import GridSearchCV
         from sklearn.metrics import r2 score
In [28]: kf = KFold(shuffle=True, random_state = 2, n_splits = 5)
```

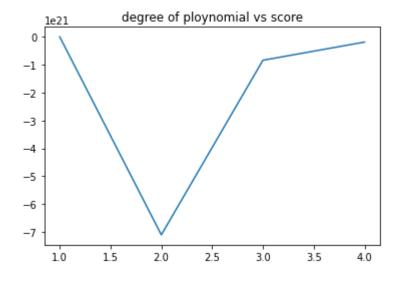
Linear Model without Regularization

```
In [29]: estimator = Pipeline([("polynomial_features", PolynomialFeatures()),
                                ("scaler", StandardScaler()),
                                ("ridge regression", LinearRegression())])
         params = {'polynomial_features__degree': [1, 2, 3]}
         grid = GridSearchCV(estimator, params, cv=kf)
In [30]: grid.fit(X, y)
Out[30]: GridSearchCV(cv=KFold(n_splits=5, random_state=2, shuffle=True),
                      estimator=Pipeline(steps=[('polynomial features',
                                                  PolynomialFeatures()),
                                                 ('scaler', StandardScaler()),
                                                 ('ridge regression',
                                                  LinearRegression())]),
                      param_grid={'polynomial_features__degree': [1, 2, 3]})
In [31]: |grid.best_score_.round(2), grid.best_params_
Out[31]: (0.66, {'polynomial features degree': 1})
```

```
In [32]: degrees = [1,2,3,4]
         lin_dict = {'degree':degrees, 'score':[]}
         lr = LinearRegression()
         s = StandardScaler()
         for d in degrees:
             pf = PolynomialFeatures(degree=d)
             estimator = Pipeline([("polynomial_features", pf),
                                    ("scaler", s),
                                    ("linear_regression", lr)])
             y_red = cross_val_predict(estimator, X, y, cv = kf)
             score = numpy.round(r2_score(y,y_red),2)
             lin_dict['score'].append(score)
```

```
In [33]: lin df = pd.DataFrame(lin dict)
         print(lin_df.sort_values('score', ascending=False))
         plt.plot(lin_df['degree'], lin_df['score'])
         plt.title('degree of ploynomial vs score');
```

```
degree
                   score
0
        1 6.600000e-01
3
        4 -1.896192e+20
2
        3 -8.410814e+20
1
        2 -7.105933e+21
```



This verifies that the gridserach method output the best model prediction

Linear Regression with L2 Regularization (Ridge)

```
In [34]: rr estimator = Pipeline([("polynomial features", PolynomialFeatures()),
                               ("scaler", StandardScaler()),
                               ("ridge_regression", Ridge())])
         rr params = {'polynomial features degree': [1, 2, 3],
                   'ridge_regression__alpha': numpy.geomspace(0.00001, 100000, 11)
         rr grid = GridSearchCV(rr estimator, rr params, cv=kf)
In [35]: rr grid.fit(X,y)
Out[35]: GridSearchCV(cv=KFold(n_splits=5, random_state=2, shuffle=True),
                      estimator=Pipeline(steps=[('polynomial features',
                                                PolynomialFeatures()),
                                                ('scaler', StandardScaler()),
                                                ('ridge_regression', Ridge())]),
                      param_grid={'polynomial_features__degree': [1, 2, 3],
                                  'ridge_regression__alpha': array([1.e-05, 1.e-04, 1.e-
         03, 1.e-02, 1.e-01, 1.e+00, 1.e+01, 1.e+02,
                1.e+03, 1.e+04, 1.e+05])})
In [36]: numpy.round(rr grid.best score ,3), rr grid.best params
Out[36]: (0.713, {'polynomial_features__degree': 3, 'ridge_regression__alpha': 1000.0})
         Linear Regression with L1 Regularization (LASSO)
```

```
In [41]: | lr_estimator = Pipeline([("polynomial_features", PolynomialFeatures()),
                                   ("scaler", StandardScaler()),
                                   ("lasso_regression", Lasso(max_iter = 100000))])
         lr_params = {'polynomial_features__degree': [1,2,3],
                       'lasso regression alpha': [0.001, 0.01, 0.1,1,10]
         lr grid = GridSearchCV(lr estimator, lr params, cv=kf)
In [42]: |lr_grid.fit(X,y)
Out[42]: GridSearchCV(cv=KFold(n splits=5, random state=2, shuffle=True),
                       estimator=Pipeline(steps=[('polynomial_features',
                                                  PolynomialFeatures()),
                                                 ('scaler', StandardScaler()),
                                                 ('lasso regression',
                                                  Lasso(max iter=100000))]),
                       param grid={'lasso regression alpha': [0.001, 0.01, 0.1, 1, 10],
                                   'polynomial features degree': [1, 2, 3]})
```

```
In [40]:
          numpy.round(lr_grid.best_score_,3), lr_grid.best_params_
Out[40]: (0.716, {'lasso_regression__alpha': 0.001, 'polynomial_features__degree': 3})
```

In [65]: alphas = numpy.array([0.005, 0.05, 0.1, 1, 5, 20, 50, 80, 100, 120,140])

ElasticNet

```
l1 ratios = numpy.linspace(0.1, 0.9, 9)
In [68]: en estimator = Pipeline([("polynomial features", PolynomialFeatures()),
                                   ("scaler", StandardScaler()),
                                   ("elasticNet_regression", ElasticNet(max_iter = 100000))
         en params = {'polynomial features degree': [1,2,3],
                       'elasticNet_regression__alpha': alphas,
                       'elasticNet regression l1 ratio': l1 ratios
         en grid = GridSearchCV(en estimator, en params, cv=kf)
In [69]: en grid.fit(X,y)
Out[69]: GridSearchCV(cv=KFold(n_splits=5, random_state=2, shuffle=True),
                       estimator=Pipeline(steps=[('polynomial features',
                                                  PolynomialFeatures()),
                                                  ('scaler', StandardScaler()),
                                                  ('elasticNet regression',
                                                  ElasticNet(max iter=100000))]),
                       param_grid={'elasticNet_regression__alpha': array([5.0e-03, 5.0e-0
         2, 1.0e-01, 1.0e+00, 5.0e+00, 2.0e+01, 5.0e+01,
                 8.0e+01, 1.0e+02, 1.2e+02, 1.4e+02]),
                                   'elasticNet regression l1 ratio': array([0.1, 0.2, 0.
         3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9]),
                                    'polynomial features degree': [1, 2, 3]})
In [70]: numpy.round(en grid.best score ,3), en grid.best params
Out[70]: (0.715,
          {'elasticNet_regression__alpha': 0.005,
            'elasticNet regression l1 ratio': 0.2,
            'polynomial features degree': 3})
         From the scores registered, the lasso regression model has the best score. However, lasso model
         has tendency to bias, therefore, I will use ElsticNet.
In [71]: from sklearn.model selection import train test split
In [72]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.1, random_s
```

```
In [74]: best_estimator = Pipeline([("polynomial_features", PolynomialFeatures(degree=3))]
                                    ("scaler", StandardScaler()),
                                    ("elasticNet_regression", ElasticNet(alpha = 0.005, 11
         best_estimator.fit(X_train, y_train)
         best_estimator.score(X_train, y_train).round(2)
Out[74]: 0.73
In [75]: y_predict = best_estimator.predict(X_test)
In [76]: r2_score(y_predict, y_test)
Out[76]: 0.6013673693934608
In [77]: import joblib
 In [ ]: joblib.dump(en_grid, 'en_grid.pkl')
In [ ]:
```