CSCI 538 Final Project In this notebook, we take a look at housing data collected from King County, Washington from May 2014-2015. The data has a number of different features included in the data set that describe different houses that were sold during that time period. The problem to be solved is to see if we can train a machine learning algorithm to model the data that was collected and use that model to predict the prices of houses that could be sold in the future. While trying to solve this problem we will look at exploration of the data, cleaning the data to prepare for the training of the model, and finally we will train the model and evaluate how well it was trained. For this project we will be using the scikit-learn python library that includes many simple and efficient tools for predictive data analysis. Scikit-learn is an open source machine learning library that supports supervised and unsupervised learning. It also provides various tools for model fitting, data preprocessing, model selection and evaluation, and many other utilities. **Housing Data Exploration** In the first section the housing data will be explored by taking a look at the different features of the data and visualizing it using different plotting methods. Exploration into the correlation between the different features will show how each one affects the price of the house. Exploring the data is a vital part of the machine learning process because before you can train a model you have to know what kind of data your dealing with. Any missing data must be fixed otherwise it will affect the training and subsequently the performance of the model. In [1]: #Importing pyton libraries needed to load, explore, visualize, and manipulate the data import numpy as np import matplotlib as mpl import matplotlib.pyplot as plt import seaborn as sbn import pandas as pd In [2]: **from sklearn.linear_model import** LinearRegression from sklearn.linear_model import SGDRegressor from sklearn.linear_model import Ridge from sklearn.linear_model import Lasso from sklearn.linear_model import ElasticNet from sklearn.metrics import mean_squared_error from sklearn.metrics import mean_absolute_error from sklearn.metrics import r2_score from sklearn.model_selection import train_test_split In [3]: #Loading the data into a pandas dataframe so that we can better visualize and manipulate data = pd.read_csv('kingcoutyhousingdata.csv') Using the head() function lets us explore the first 5 instances of the data and outputs the values for each of the features. This allows a quick overview of the features and the data types that are present in the data In [4]: #Displaying the first 5 instances in the dataset data.head() Out[4]: price bedrooms bathrooms livingsq sqft_lot floors waterfront view condition grade sqft_above sqft_ment yr_built yr_re_ated zipcode 98178 47.5112 **0** 221900 1 1180 5650 1180 1955 **1** 538000 2 2570 7242 2 0 3 7 2170 400 1951 98125 47.7210 1 770 10000 770 0 1933 0 98028 47.7379 **2** 180000 1 0 0 3 6 **3** 604000 1960 5000 1050 910 1965 0 98136 47.5208 **4** 510000 2 1680 8080 0 0 1680 0 1987 0 98074 47.6168 1 Before any training on the data begins, information such as data types and the number of null entries for features is very important to review. Any features that are not numerical must be converted to numberical features because the regression algorithms will not train on those features. Also, any null entries will also affect the regression algorithm as it will actually think that the feature is 0 instead of missing. Looking at the info() function, we can see that all features are numerical and there are no non-null entries for each feature. There are a total of 19 different features with a breif description of each: **Price:** Sale price **Bedrooms:** Number of bedrooms Bathrooms: Number of bathrooms Livingsq: Size of living area in square feet **Sqft_lot:** Size of lot in square feet Floors: Number of floors Waterfront: '1' if the porperty has a waterfront, '0' if not **View:** An index from 0 to 4 of how good the view on the property is **Condition:** Condition of the house, ranked from 1 to 5 Grade: Classification by construction quality which refers to the types of materials used and the quality of workmanship; the higher the better. **Sqft_above:** Square feet above ground **Sqft_ment:** Square feet below ground Yr_built: Year built Yr_re_ated: Year renovated. '0' if never renovated **Zipcode:** 5 digit zip code **Lat:** Latitude Long: Longitude Sqft_ng15: Average size of interior housing living space for the closest 15 houses, in square feet **Sqft_lot15:** Average size of land lost for the closest 15 houses, in square feet In [5]: #Description of the different features with their non-null count and data types data.info() <class 'pandas.core.frame.DataFrame'> RangeIndex: 21613 entries, 0 to 21612 Data columns (total 19 columns): Column Non-Null Count Dtype - - -----------0 price 21613 non-null int64 bedrooms 21613 non-null int64 1 21613 non-null int64 bathrooms livingsq 21613 non-null int64 sqft_lot 21613 non-null int64 5 floors 21613 non-null int64 waterfront 21613 non-null int64 6 view 21613 non-null int64 8 condition 21613 non-null int64 9 grade 21613 non-null int64 sqft_above 21613 non-null int64 11 sqft_ment 21613 non-null int64 12 yr_built 21613 non-null int64 13 yr_re_ated 21613 non-null int64 14 zipcode 21613 non-null int64 15 lat 21613 non-null float64 16 long 21613 non-null float64 17 sqft__ng15 21613 non-null int64 18 sqft_lot15 21613 non-null int64 dtypes: float64(2), int64(17) memory usage: 3.1 MB The describe() function gives a more statistical description of the data providing details such as the mean, standard devation (std), minimum value, maximum value, and the values that represents the 25, 50, and 75 percentiles. In [6]: data.describe().T Out[6]: 25% **50**% **75**% count mean std min max **price** 21613.0 540088.141767 367127.196483 75000.0000 321950.000 450000.0000 645000.000 7.700000e+06 **bedrooms** 21613.0 4.000 1.100000e+01 3.369454 0.907964 0.0000 3.000 3.0000 bathrooms 21613.0 1.749734 0.734873 0.0000 1.000 2.0000 2.000 8.000000e+00 2550.000 1.354000e+04 livingsq 21613.0 2079.899736 918.440897 290.0000 1427.000 1910.0000 **sqft_lot** 21613.0 41420.511515 15106.967566 520.0000 5040.000 7618.0000 10688.000 1.651359e+06 floors 21613.0 1.446213 0.551894 1.0000 1.000 1.0000 2.000 3.000000e+00 waterfront 21613.0 0.007542 0.086517 0.0000 0.000 0.0000 0.000 1.000000e+00 view 21613.0 0.000 4.000000e+00 0.234303 0.766318 0.0000 0.000 0.0000 condition 21613.0 3.409430 0.650743 1.0000 3.000 3.0000 4.000 5.000000e+00 7.656873 8.000 1.300000e+01 grade 21613.0 1.175459 1.0000 7.000 7.0000 sqft_above 21613.0 2210.000 9.410000e+03 1788.390691 828.090978 290.0000 1190.000 1560.0000 442.575043 **sqft_ment** 21613.0 291.509045 0.0000 0.000 0.0000 560.000 4.820000e+03 **yr_built** 21613.0 1971.005136 29.373411 1900.0000 1951.000 1975.0000 1997.000 2.015000e+03 yr_re_ated 21613.0 84.402258 401.679240 0.0000 0.000 0.0000 0.000 2.015000e+03 53.505026 98001.0000 **zipcode** 21613.0 98033.000 98065.0000 98118.000 9.819900e+04 98077.939805 lat 21613.0 47.560053 0.138564 47.1559 47.471 47.5718 47.678 4.777760e+01 long 21613.0 -122.213896 0.140828 -122.5190 -122.328 -122.2300 -122.125 -1.213150e+02 1986.552492 1490.000 1840.0000 2360.000 6.210000e+03 **sqft__ng15** 21613.0 685.391304 399.0000 **sqft_lot15** 21613.0 12768.455652 27304.179631 651.0000 5100.000 7620.0000 10083.000 8.712000e+05 Visualizing the histograms of the data can reveal a few things which can then be used during the data cleanup phase. Looking at the data here we can see that some of the features are very tail-heavy and do not exhibit a normal distribution. Many of the features also have very different scales. Bedrooms range from 0 to 11 but livingsq ranges from 290 to 2550. Different scales like this are problems for some machine learning models, and thus we may need to scale attributes like these with widely different ranges to have a common similar range of values. In [7]: data.hist(bins=50, figsize=(20, 15)) plt.show() bathrooms bedrooms condition floors 10000 12500 10000 12500 8000 10000 8000 10000 6000 7500 6000 7500 4000 5000 -4000 5000 2000 2500 -2000 2500 1.5 2.0 livingsq 3000 2000 1000 8000 800 1500 6000 2000 600 1000 -4000 400 1000 500 2000 47.2 47.3 47.4 47.5 47.6 47.7 47.8 2500 5000 7500 10000 12500 -122.4-122.2-122.0-121.8-121.6-121.4 sqft_ng15 sqft_above 20000 6000 -2500 1500 15000 2000 4000 1500 1000 10000 -1000 2000 500 5000 -1000 2000 3000 4000 5000 6000 2000 4000 sqft_lot15 waterfront 20000 20000 12500 20000 15000 15000 10000 15000 7500 10000 10000 10000 5000 5000 5000 -5000 2500 200000 400000 600000 800000 2000 3000 0.4 zipcode 1250 20000 1500 15000 10000 Using a the scatter_matrix() function will plot different features against each other to show if there is any linear correlation. Including the price feature, the one that is used as the target features, into the scatter matrix will show which features will be the most promising in predicting the price of the houses. In [9]: #Import the scatter_matrix function from pandas.plotting import scatter_matrix attributes = ['price', 'livingsq', 'yr_built', 'sqft_above', 'long', 'lat'] scatter_matrix(data[attributes], alpha=0.1, figsize=(18, 14)) plt.show() 일 4000000 2000000 12500 10000 7500 1975 1925 6000 분 4000 · -121.50 -121.75 -122.00 -122.25 -122.50 le6 price yr_built sqft_above Similarly, the correlation function corr_matrix will give us numerical values to look at to help decide which functions provide a good linear correlation and which ones don't. Correlation matrix values range from -1 to 1. When it is close to 1 (a positive correlation) it means that there is a strong linear correlation between the features. This means that when one feature increase, the other usually increases in some standard ratio. A negative correlation, near to -1, means a correlation but in the opposite direction. When one increases, the other decreases in some standard linear ratio. A correlation of 0 means that there doesn't appear to be a relationship, when one increases, the other is as likely to increase or decrease or stay the same. In [9]: corr_matrix = data.corr() corr_matrix['price'].sort_values(ascending=False) Out[9]: price 1.000000 livingsq 0.702035 grade 0.667434 sqft_above 0.605567 0.585379 sqft__ng15 bathrooms 0.510072 view 0.397293 sqft_ment 0.323816 bedrooms 0.315438 lat 0.307003 waterfront 0.266369 floors 0.237211 yr_re_ated 0.126434 sqft_lot 0.089661 sqft_lot15 0.082447 yr_built 0.054012 condition 0.036362 long 0.021626 zipcode -0.053203 Name: price, dtype: float64 Just by looking at the numbers, it shows that livingsq is the most linearly correlated feature with the target price and we can visualize this by looking at the pair on a scatter plot. By looking at the scatter plot below, it is fairly obvious that there is a strong linear correlation between these two features. You can see what looks like an upward trend such that an increase in livingsq means there is an increase in the price. In [16]: data.plot(kind='scatter', x='livingsq', y='price', alpha=0.1, figsize=(12,8)); 14000 2000 4000 6000 8000 10000 12000 livingsq Although this set of data has a good number of features to train our model on, there may be times when we look at the correlation matrix data and there seems to be not many features that strongly correlate with our target label. Feature engineering is something that data scientists can do to provide significant improvements to the machine learning modeling. After initial exploration of data, you might begin to ask yourself what information might be useful for a model to learn to predict from. We can look at the existing features and create features with a clearer signal from existing data. Below you can see that the bedrooms and bathrooms features provide a decent correlation with the target label. Just to experiment, we can create another feature called rooms_per_sqft by adding the bedroom and bathroom features together and dividing them by the total livingsq. In [11]: data['rooms_per_sqft'] = (data['bedrooms'] + data['bathrooms']) / data['livingsq'] By running the corr_matrix function again with this new feature we can see that we've created a new feature that has a good negative correlation with the target label. In [12]: corr_matrix = data.corr() corr_matrix['price'].sort_values(ascending=False) Out[12]: price 1.000000 livingsq 0.702035 grade 0.667434 sqft_above 0.605567 sqft__ng15 0.585379 bathrooms 0.510072 view 0.397293 sqft_ment 0.323816 bedrooms 0.315438 lat 0.307003 waterfront 0.266369 floors 0.237211 yr_re_ated 0.126434 0.089661 sqft_lot sqft_lot15 0.082447 yr_built 0.054012 condition 0.036362 long 0.021626 zipcode -0.053203 rooms_per_sqft -0.461442 Name: price, dtype: float64 The downward trend is obvious here but it looks more like a polynomial correlation more than a linear one. This feature engineering was just for example purposes and we won't be actually using it in the modeling of our regression algorithm. In [13]: data.plot(kind='scatter', x='rooms_per_sqft', y='price', alpha=0.1, figsize=(12,8)); 0.000 0.001 0.002 0.003 0.004 0.005 0.006 0.007 rooms_per_sqft In [14]: |#Dropping the 'rooms_per_sqft' feature and showing that our data is back to its original state data.drop('rooms_per_sqft', axis=1, inplace=True) **Preparing the Data for Linear Regression Algorithms** Once the data been explored, the next step is to prepare it so that we can train our machine learning models with it. The first thing that can be done is to drop features from the data set. Dropping features that don't have strong correlations with the target label can help the linear regression algorithms fit the training data without overfitting it. Overfitting happens when the algorithms tries too hard to fit the training data that it doesn't generalize well to unseen data. Looking at the correlation matrix, there are a number of features that a very low correlation with the target label for each instance. Features such as zipcode, yr_re_ated, sqft_lot15, can be dropped to without very little hinderance to the accuracy and precision of our model and will also speed up the training. In [15]: #Dropping the 3 features that are not needed then printing the first 5 data instances to show those features are gon data.drop(['zipcode', 'yr_re_ated', 'sqft_lot15'], axis=1, inplace=True) data.head() Out[15]: price bedrooms bathrooms livingsq sqft_lot floors waterfront view condition grade sqft_above sqft_ment yr_built long sqft_ng15 **0** 221900 0 1955 47.5112 -122.257 1 1180 5650 1180 **1** 538000 2570 7242 2 7 2170 1951 47.7210 -122.319 2 0 3 **2** 180000 770 10000 1 0 0 3 6 770 1933 47.7379 -122.233 **3** 604000 1960 1050 1965 47.5208 -122.393 3 5000 1 0 0 5 **4** 510000 2 1680 8080 1680 0 1987 47.6168 -122.045 1 0 0 The next step in preparing the data is to split the dataset up into a training set and a testing it. This is done so that we can help train our model so that it generalizes well to unseen data. If we were to train our model on our whole data set, we wouldn't be able to evaluate the performance of it because it would always predict correctly when asked to make a prediction. Training on only a portion of the data allows for the evaluation of the learned hypotesis function on that test data that was not seen during the training. Once we see how the hypothesis function performs on the unseen data, we can then start to make adjustments on the model to get it to the best it can be. Using the sklearn train_test_split function allows the data to be split into a training set (which will be used to train the model with), and a test set which will be used to evaulate the model to gauge how successful the training was. The test size 0.2 specifies how big we want the testing data set to be. The parameter random_state which is set to 42 will ensure that anyone else who runs this notebook will have the same instances split into the training and testing sets and therefore have the same results in their evaulation of the model. In [16]: data_train, data_test = train_test_split(data, test_size=0.2, random_state=42) print(data_train.shape) print(data_test.shape) (17290, 16)(4323, 16) The data is now split into a training set with 17290 instances and a testing set with 4323 instances and the next step is to further prepare our training dataset by dropping the price feature and putting it into a separate vector. This will let us train different models with different parameters so that the models can learn to predict the target label price based on the training data. In [17]: housing = data_train.drop('price', axis=1) housing_labels = data_train['price'].copy() housing_test = data_test.drop('price', axis=1) housing_test_labels = data_test['price'].copy() print(housing.shape) print(housing_labels.shape) print(housing_test.shape) print(housing_test_labels.shape) (17290, 15) (17290,)(4323, 15)(4323,)At this point, there dataset is pretty clean and is ready to be used to train machine learning models. **Training Linear Regression Algorithms** We now can train a number of machine learning models, evaluate, and then tune them a bit to try and get the best one that we can then use to predict future price of houses in the same area. All models from sklearn are full estimator - predictors. The purpose of a supervised learning machine learning algorithm is to generate a hypothesis function h() from the training data, and use the learned hypothesis to predict the target labels of values of the unseen data. In [18]: linear regression = LinearRegression() linear_regression.fit(housing, housing_labels) Out[18]: LinearRegression() Now that the model has been trained (fitted), we can now test out and evaluate how well it makes predictions. Before we do that though it would be nice to run some formal evaluations on our model to see how well it performs. Using performance measures such as the R-squared score and confidence intervals we can see how well the model is fitted and is making perdictions on the unseen data. The R-squared score is a measure of the scatter of the data points around the fitted regression line. It is also called the coefficient of determination, or the coefficient of multiple determination for multiple regression. For the same data set, higher R-squared values represent smaller differences between the observed data and the fitted values. R-squared scores are in the range of 0 to 1 and is normally measured as a percentage of the variance in the dependent variable that the independent variables explain collectively. In [19]: lin_reg_predict = linear_regression.predict(housing_test) print("R^2 Score: ",r2_score(housing_test_labels, lin_reg_predict)) R^2 Score: 0.6955638499670669 The Linear Regression algorithm model we trained can explain about 69.5% of the total variance between the predicted values and the actual values. This number alone doesn't really tell us much about well of a number the model is predicting because of different biases in the model. For instance, some predictions could be really close to the regression line but some could be very far which would explain a R^2 score of 69.5%. The R^2 score isn't the only metric we can use to evaluate a model. Calulating the model's Mean Absolute Error (MAE) will give us a sum of the difference between the data and our model's prediction (whether it be over predicting or under predicting) for each instance of data. Using the Linear Regression model we can see that our MAE for the entirety of our test data that it was off by over 128,000. MAE isn't always a good metric to use because the under-predicted prices will work to cancel our the over-predicted ones. In [20]: print("Mean Absolute Error: ", mean_absolute_error(housing_test_labels, lin_reg_predict)) Mean Absolute Error: 128673.0942688452 Yet another metric we can use is the Root Mean Squared Error (RMSE) which squares all of the residual values. This gives us a better overall picture of the model and will penalize our model more heavily for any big outliers in the predictions. The sum is made over all of the residual vs. actual values so that prices that are under-predicted will add on the total error of the model instead of working with the over-predicted ones like in the MAE metric. The RMSE should actually be higher than what we computed with the MAE and we can see that below with a RMSE score of 214,531. In [21]: lin_reg_mse = mean_squared_error(housing_test_labels, lin_reg_predict) lin_reg_rmse = np.sqrt(lin_reg_mse) print("Root Mean Squared Error: ", lin_reg_rmse) Root Mean Squared Error: 214531.24773355035 Another evaluation that works well in evaluating a model is it's confidence interval. We can gauge performance if we compute a 95% confidence interval. This 95% confidence interval is for the generalization error of the target label. Computing the 95% confidence interval on our Linear Regression model

1340

1690

2720

1360

1800

95% Confidence Level: [193852.91520895 233384.57483994] We can see here that by using a standard Linear Regression algorithm our model's predictions were off by an average of about \$200,000. This is quite a large amount which would never work in the standard business model. It is hard to know exactly why the RMSE was so high sometimes its that there is not enough features or enough strongly correlated features. The linear regression algorithm is not a very powerful algorithm and was likely underfitting the training data which is why we see such a big difference between the predicted priced and actual target labels. We can next try a much more powerful model to see if we

tree_reg_predict = tree_regression.predict(housing_test)

95% Confidence Level: [171272.3383085 240612.67416067]

-3.20648074e+10 -4.00955861e+10 -2.48642057e+10 -3.34804650e+10

forest_reg_mse = mean_squared_error(housing_test_labels, forest_predict)

95% Confidence Interval: [123850.01432183 172323.51458866]

print("Score: ",r2_score(housing_test_labels, tree_reg_predict))

n(), scale=stats.sem(squared_errors))))

techniques that we did with our Linear Regression model.

tree_regression = DecisionTreeRegressor() tree_regression.fit(housing, housing_labels)

n(), scale=stats.sem(squared_errors))))

Mean Absolute Error: 105698.45327318991 Root Mean Squared Error: 208840.45685213033

-4.27870614e+10 -2.98706042e+10]

Stanard Deviation: 6355606124.193297

exhibits better performance than the single decision trees.

In [29]: from sklearn.ensemble import RandomForestRegressor

could be a confident predictor of housing prices.

unlikely that we would have seen something that would be used.

Conclusion

Mean: -34208384007.385353

In [23]: from sklearn.tree import DecisionTreeRegressor

can get better performance from it.

Score: 0.7115009595915565

In [22]: **from scipy import** stats confidence = 0.95

\$230,384! This is a huge error and not something that we want our model to be doing.

squared_errors = (lin_reg_predict - housing_test_labels)**2

In [24]: print("Mean Absolute Error: ", mean_absolute_error(housing_test_labels, tree_reg_predict)) tree_reg_mse = mean_squared_error(housing_test_labels, tree_reg_predict) tree_reg_rmse = np.sqrt(tree_reg_mse) print("Root Mean Squared Error: ", tree_reg_rmse) squared_errors = (tree_reg_predict - housing_test_labels)**2

print("95% Confidence Level: ", np.sqrt(stats.t.interval(confidence, len(squared_errors) - 1, loc=squared_errors.mea

Using the Decision Tree Regressor our MAE and RMSE performance metrics are down a little bit from the Linear Regression model but it looks like there is

Regressor model was overfitting the data. For k-fold cross-validatoin, we split the training data up into roughly k equal pieces. The model is then fitted and evaluated k different times then the average and variance of the performance is computed across the training runs. Each run will take out 1 of the

Wow! You can see here how the Decision Tree Regressor has done very poorly with a cross-validation evaluation. The model is highly overfitting the training

Lastly, we can the Random Forest Regressor which is actually a collection of many decision trees. This is an ensemble machine learning model and normally

data and you can see how well it won't generalize well to the unseen data. This is one of the downfalls to the Decision Tree Regression algorithms.

still a lot of room for improvement. More formal evaluation of the model using k-fold cross-validation can help us to see if the Decision Tree

shows that the model is 95% certain the predicted price of an instance of data will be between 193,852 - 233,384 and have an error no larger than

print("95% Confidence Level: ", np.sqrt(stats.t.interval(confidence, len(squared_errors) - 1, loc=squared_errors.mea

The Decision Tree Regressor is a much more powerful algorithm that we can use to see if we can get a better outcome. We will use the same evaluation

```
k fols, train with the data from the other k-1 folds and evaluate performance on the held out fold. This gives a much better estimate of how well a model wil
          generalize on unseen data.
In [28]: from sklearn.model_selection import cross_val_score
          scores = cross_val_score(tree_regression, housing, housing_labels, scoring='neg_mean_squared_error', cv=10)
          tree_rmse_scores = np.sqrt(-scores)
          print("Scores: ", scores)
          print("Mean: ", scores.mean())
          print("Stanard Deviation: ", scores.std())
          Scores: [-2.82174726e+10 -4.59547707e+10 -3.09898252e+10 -3.37590419e+10
```

forest_reg = RandomForestRegressor(n_estimators=100, random_state = 42) forest_reg.fit(housing, housing_labels) forest_predict = forest_reg.predict(housing_test) print("Score: ",r2_score(housing_test_labels, forest_predict)) Score: 0.8510543751897715 In [30]: print("Mean Absolute Error: ", mean_absolute_error(housing_test_labels, forest_predict))

forest_reg_rmse = np.sqrt(forest_reg_mse) print("Root Mean Squared Error: ", forest_reg_rmse) squared_errors = (forest_predict - housing_test_labels)**2 print("95% Confidence Interval: ", np.sqrt(stats.t.interval(confidence, len(squared_errors) - 1, loc=squared_errors. mean(), scale=stats.sem(squared_errors)))) Mean Absolute Error: 73710.66701127964 Root Mean Squared Error: 150057.02204113491

We can see from the above metrics that the models we are training are trending in the correct direction but we are not to the point where there are models that

After exploring the housing data, cleaning it some, and training a few models the model that were trained didn't end up being useful enough to be able to use it

to predict any future housing prices. The models we looked at could have been tweaked some but with how far off our performance metrics were it was

Further investigation into the data could reveal that we didn't have enough of a linear relationship between the target label and the features we used to train the model. Improvements could be made to our model by dropping more features or engineering new ones. More complete examination of the data might

determine that there a polynomal fit to our selected features which would require a different set of algorithms and features that are fit to a polynomial set of theta parameters.