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CE475

Fundamentals and Applications of Machine Learning

Project Report

Identification and Significance of the Problem

In this project, the goal is to find the best model for first 100 row in the data which has response values and then predict the last 20 row's response values with that method.

Methodology

Multiple Linear Regression

I've tried backward elimination in MLR and eliminated features x1 and x4 based on their p_values. After that, to prove that I've eliminated right features, I compare the root mean squared errors with cross-validation of all subsets (2^n). Model without x1 and x4 has the best rmse value.

Polynomial Regression

In Polynomial Regression, I've tried to find the best degree. RMSE significantly increased after degree 3, so I choose degree 3 for my polynomial model. However, It was still high and it was close to the MLR model's rmse value.

Lasso

I've used Lasso since it enhances the prediction accuracy and interpretability of the statistical model it produces. LassoCV selects the best model with cross-validation. Since it's rmse value is not satisfying, I've skipped to my next model.

Decision Tree Regressor

Decision tree builds regression or classification models in the form of a tree structure. It breaks down a dataset into smaller and smaller subsets while at the same time an associated decision tree is incrementally developed. The final result is a tree with decision nodes and leaf nodes. I've used GridSearchCV to find best DTR model. RMSE value was better.

Random Forest Regressor

A random forest is a meta estimator that fits a number of classifying decision trees on various sub-samples of the dataset and uses averaging to improve the predictive accuracy and control over-fitting. I've used GridSearchCV with 1000 estimators. RMSE value is so much better so I decided to use Random Forest Regressor to predict last 20 rows.

Implementation

In the implementation part, I've used spyder as my integrated development environment since I like using it's variable explorer feature and the jupyter notebook for presentation purposes. I've also used the scikit-learn library, pandas, numpy and seaborn. Implementation process can be seen below in appended the jupyter notebook.

Results

MLR		Polynomial R.		Lá	Lasso		DTR		RFR	
Activities ≡ Spyder ▼										
civici	эручен									
0		0			0		0		0	
0	1110.36	0	2189.62	0	1017.96	0	600	0	982.485	
1	850.545	1	-1462.19	1	846.275	1	93	1	91.559	
2	1238.38	2	3773.82	2	1128.54	2	90	2	92.261	
3	-403.082	3	-259.932	3	-173.536	3	71	3	84.238	
4	1990.11	4	2111.59	4	1760.51	4	1266	4	1973.33	
5	-1237.67	5	1921.64	5	-845.137	5	89	5	-208.434	
6	140.48	6	-2355.81	6	260.435	6	90	6	96.606	
7	882.786	7	474.287	7	979.274	7	64	7	69.494	
В	136.991	8	2224.77	8	235.615	8	-1611	8	-727.437	
9	1781.11	9	1335.33	9	1603.99	9	4896	9	3214.39	
10	2483.61	10	-1181.4	10	2144.2	10	1602	10	1795	
11	-966.046	11	-1221.83	11	-638.138	11	-1611	11	-715.243	
12	-381.436	12	-308.988	12	-229.874	12	-1138	12	-1055.69	
13	692.774	13	994.62	13	700.111	13	0	13	280.137	
14	16.2324	14	-703.001	14	118.793	14	61	14	48.481	
15	1337.49	15	5482.77	15	1301.7	15	94	15	81.8	
16	284.679	16	-327.534	16	479.006	16	90	16	89.579	
17	2353.18	17	4755.47	17	2119.81	17	4449	17	3831.15	
18	-96.2455	18	-5201.23	18	94.673	18	-1638	18	-747.102	
19	51.0311	19	-2157.38	19	184.795	19	71	19	130.192	
Format Re Format Re					Format Re	esi	Format Re	s F	Format Re	

Conclusion

To conclude, I've used various models and choose the one with the lowest root mean squared error value. In this case, it was Random Forest Regression for me.

There are a couple of lessons I've learned during the process. I've always forget to train my data at the end with all the data and use a model built upon training data from train_test_split.

I've also learned I've to %reset my ipython to not get confused with variables related to the previous model.

Sometimes DataFrame gets the previous row as a header and the code hard to debug if I don't notice it early on.

I have to pay more attention to documentation, for example, I've tried to build over 200 polynomial features on my own because I thought PolynomialFeatures is doing something else rather than that.

And I've learned that train_test_split is not enough, I've to use cross-validation if I can.

In [1]:

```
# %% Import Libraries
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import numpy as np
from sklearn import metrics
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import train_test_split
from sklearn.model_selection import GridSearchCV
from sklearn.linear model import LinearRegression
from sklearn.preprocessing import PolynomialFeatures
from sklearn.tree import DecisionTreeRegressor
from sklearn.metrics import make scorer
from sklearn.ensemble import RandomForestRegressor
import warnings
warnings.filterwarnings('ignore')
```

/opt/anaconda/anaconda3/lib/python3.7/site-packages/sklearn/ensemble/weight_boosting.py:29: Deprecat ionWarning: numpy.core.umath_tests is an internal NumPy module and should not be imported. It will be removed in a future NumPy release.

from numpy.core.umath tests import inner1d

In [2]:

```
# %% Read CSV
df = pd.read_csv('data.csv', index_col=0, nrows=100)
df.index = np.arange(0, len(df))
willbepredicted = pd.read_csv('data.csv', index_col=0, skiprows=range(1, 101), nrows=20)

feature_cols = ['x1', 'x2', 'x3', 'x4', 'x5']

X = df[feature_cols]
y = df.Y
x_find = willbepredicted[feature_cols]
```

In [3]:

```
# %% Train Test Split
X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.20, random_state=0)
```

In [4]:

```
linreg = LinearRegression()
linreg.fit(X, y)
y_pred = linreg.predict(X)
```

In [5]:

```
print("x1: ", linreg.coef_[0])
print("x2: ", linreg.coef_[1])
print("x3: ", linreg.coef_[2])
print("x4: ", linreg.coef_[3])
print("x5: ", linreg.coef_[4])
```

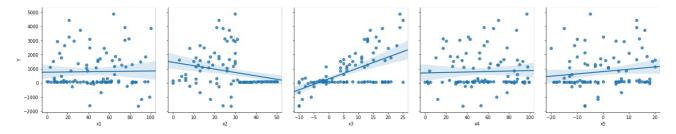
x1: -0.14813529290736643 x2: -32.1007066048948 x3: 82.67435716452609 x4: 0.5973716561125914 x5: 12.214009071344265

In [6]:

```
%matplotlib inline sns.pairplot(df, x_vars=['x1', 'x2', 'x3', 'x4', 'x5'], y_vars='Y', height=4, aspect=1.0, kind='reg')
```

Out[6]:

<seaborn.axisgrid.PairGrid at 0x7fe4771c92e8>



0.467

R-squared:

In [7]:

```
# Building the optimal model using Backward Elimination
import statsmodels.formula.api as sm
X.insert(0, "x0", 1)

X_opt = X[['x0', 'x1', 'x2', 'x3', 'x4', 'x5']]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
```

Out[7]:

OLS Regression Results

Dep. Variable:

	Model:				Adj.	R-s	quared:	0.439
	Meth	east Squ	iares	F-statistic:		16.47		
	Da	04 Dec 2018		Prob (F-statistic):		atistic):	1.20e-11	
	Time:			10:43:37		Log-Likelihood:		-825.21
No. Observations:				100			AIC:	1662.
Df Residuals:			94				BIC:	1678.
	Df Mod		5					
Co	variance Ty	nonro	bust					
	coef	std err	t	P> t	[0.0	025	0.975]
x0	1113.1040	349.862	3.182	0.002	418.	446	1807.762	2
x1	-0.1481	3.556	-0.042	0.967	-7.	208	6.911	L
x2	-32.1007	7.468	-4.298	0.000	-46.	929	-17.272	2
х3	82.6744	10.004	8.264	0.000	62.	810	102.538	3
x4	0.5974	3.631	0.165	0.870	-6.	612	7.807	7
x5	12.2140	8.437	1.448	0.151	L -4.538		28.966	5
_	Omnibus: 6.029 Durbin-Wa					1.8		
Prob(Omnibus): 0.04			Jarque-	-Bera	(JB):	5.5	01	

Warnings:

Skew: 0.555

Kurtosis: 3.298

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

296.

p value of x1 is 0.967 and it's greater than my SL. I've eliminated x1 $\,$

Prob(JB): 0.0639

Cond. No.

In [8]:

```
X_opt = X[['x0', 'x2', 'x3', 'x4', 'x5']]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
```

Out[8]:

OLS Regression Results

old Regression Results								
Dep. Variable:			Υ		R-squared		0.467	
	Mod	el:	OLS		Adj. R	squared:	0.444	
	Method:			ıares	F-statistic:		20.80	
	Da	te: Tue,	04 Dec	2018	Prob (F-s	statistic):	2.44e-12	
	Tin	ne:	10:4	13:37	Log-Li	kelihood:	-825.21	
No. Observations:				100		AIC:	1660.	
	Df Residua	ıls:		95		BIC:	1673.	
	Df Mod	el:		4				
Co	variance Ty	pe:	nonro	bust				
	coef	std err	t	P> t	[0.025	0.975]	
х0	1106.7271	312.948	3.536	0.001	485.447	1728.00	В	
x2	-32.0779	7.409	-4.330	0.000	-46.787	-17.36	9	
х3	82.6731	9.952	8.307	0.000	62.916	102.43	0	
х4	0.5788	3.585	0.161	0.872	-6.538	7.69	6	
х5	12.2223	8.390	1.457	0.148	-4.434	28.87	9	
	Omnibus	: 5.964	Durbi	n-Wats	son: 1.	889		
Pro	b(Omnibus)	: 0.051	Jarque	-Bera (JB): 5.	432		
	Skew	: 0.552		Prob(JB): 0.0	661		
	Kurtosis	3.295		Cond.	No. 2	217.		

Warnings:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

x4's p value (0.872) is greater than my SL. I've eliminated x4

In [9]:

```
X_opt = X[['x0', 'x2', 'x3', 'x5']]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
```

Out[9]:

OLS Regression Results

```
Dep. Variable:
                                Υ
                                                         0.467
                                          R-squared:
          Model:
                              OLS
                                                         0.450
                                      Adj. R-squared:
         Method:
                      Least Squares
                                          F-statistic:
                                                         28.01
            Date: Tue, 04 Dec 2018 Prob (F-statistic): 4.24e-13
           Time:
                          10:43:37
                                      Log-Likelihood:
                                                       -825.22
No. Observations:
                              100
                                                AIC:
                                                         1658.
    Df Residuals:
                               96
                                                BIC:
                                                         1669.
       Df Model:
                                3
Covariance Type:
                         nonrobust
         coef std err
                            t P>|t|
                                     [0.025
                                                0.9751
x0 1141.8474 223.861
                       5.101 0.000 697.486 1586.209
     -32.2098
                 7.326 -4.396 0.000
                                     -46.753
                                               -17.667
x2
      82.6821
                 9.901 8.351 0.000
                                      63.029
                                               102.335
x3
     12.1522
                 8.336 1.458 0.148
                                      -4.395
                                                28.700
x5
     Omnibus: 6.187
                                         1.884
                        Durbin-Watson:
Prob(Omnibus): 0.045 Jarque-Bera (JB):
                                          5.662
         Skew: 0.562
                              Prob(JB): 0.0589
      Kurtosis: 3.310
                              Cond. No.
                                           73.9
```

Warnings:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

remaining features' p values are less than my significance level 0.05

In [10]:

-381.43565364

284.67917056 2353.1827379

692.77445583

```
feature cols = [ 'x2', 'x3', 'x5' ]
X_train = df[feature_cols]
y_train = df.Y
X test = willbepredicted[feature cols]
linreg.fit(X train, y train)
y_pred = linreg.predict(X_test)
print(y_pred)
[ 1110.36291321
                 850.545366
                                1238.37578502 -403.08248109
 1990.10754432 -1237.67131187
                                 140.48020518
                                               882.78590327
  136.99100333 1781.10677895
                                2483.61099772
                                               -966.04607179
```

then I've decided find all the subsets of my data, and compare their rmse values to understand the effects of my features on my model

51.03107948]

16.23241308 1337.48754781

-96.24550126

In [11]:

```
scores = cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_error')
# fix the sign of MSE scores
mse_scores = -scores
# convert from MSE to RMSE
rmse_scores = np.sqrt(mse_scores)
# calculate the average RMSE
print("Mean of rmse with all features included: " , rmse_scores.mean())
```

Mean of rmse with all features included: 1000.0719831858908

Let's find rmse for all subsets of our features

['x5'],['x4'],['x4', x5'],['x3', 'x5'],['x3', 'x5'],['x3', 'x4'],['x3', 'x4'],['x2', 'x4'],['x2', 'x5'],['x2', 'x4'],['x2', 'x4', 'x5'],['x2', 'x3'],['x2', 'x3', 'x5'],['x2', 'x3', 'x5'],['x2', 'x3', 'x5'],['x2', 'x3', 'x5'],['x1', 'x5'],[

In [12]:

```
feature_cols = [ 'x5' ]
X = df[feature cols]
print("Mean of rmse [ 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_error'))
.mean())
feature cols = [ 'x4' ]
X = df[feature cols]
print("Mean of rmse [ 'x4' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_error'))
.mean())
feature cols = [ 'x4', 'x5' ]
X = df[feature cols]
print("Mean of rmse [ 'x4', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_er
ror')).mean())
feature cols = [ 'x3' ]
X = df[feature cols]
print("Mean of rmse [ 'x3' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_error'))
.mean())
feature cols = [ 'x3', 'x5' ]
X = df[feature_cols]
print("Mean of rmse [ 'x3', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_er
ror')).mean())
feature cols = [ 'x3', 'x4' ]
X = df[feature cols]
print("Mean of rmse [ 'x3', 'x4' ]: ", np.sqrt(-cross val score(linreg, X, y, cv=10, scoring='neg mean squared er
ror')).mean())
feature cols = [ 'x3', 'x4', 'x5' ]
X = df[feature_cols]
print("Mean of rmse [ 'x3', 'x4', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squa
red_error')).mean())
feature_cols = [ 'x2' ]
X = df[feature cols]
print("Mean of rmse [ 'x2' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_error'))
.mean())
feature cols = [ 'x2', 'x5' ]
X = df[feature cols]
print("Mean of rmse [ 'x2', 'x5' ]: ", np.sqrt(-cross val score(linreg, X, y, cv=10, scoring='neg mean squared er
ror')).mean())
feature cols = [ 'x2', 'x4' ]
X = df[feature_cols]
print("Mean of rmse [ 'x2', 'x4' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_er
ror')).mean())
feature_cols = [ 'x2', 'x4', 'x5' ]
X = df[feature_cols]
print("Mean of rmse [ 'x2', 'x4', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squa
red error')).mean())
feature_cols = [ 'x2', 'x3' ]
X = df[feature cols]
print("Mean of rmse [ 'x2', 'x3' ]: ", np.sqrt(-cross val score(linreq, X, y, cv=10, scorinq='neq mean squared er
```

```
ror')).mean())
feature_cols = [ 'x2', 'x3', 'x5' ]
X = df[feature cols]
print("Mean of rmse [ 'x2', 'x3', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squa
red error')).mean())
feature_cols = [ 'x2', 'x3', 'x4' ]
X = df[feature cols]
print("Mean of rmse [ 'x2', 'x3', 'x4' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squa
red error')).mean())
feature cols = [ 'x2', 'x3', 'x4', 'x5' ]
X = df[\overline{feature cols}]
print("Mean of rmse [ 'x2', 'x3', 'x4', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mea
n squared error')).mean())
feature cols = [ 'x1' ]
X = df[feature_cols]
print("Mean of rmse [ 'x1' ]: ", np.sqrt(-cross val score(linreg, X, y, cv=10, scoring='neg mean squared error'))
.mean())
feature_cols = [ 'x1', 'x5' ]
X = df[feature_cols]
print("Mean of rmse [ 'x1', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_er
ror')).mean())
feature cols = [ 'x1', 'x4' ]
X = df[feature cols]
print("Mean of rmse [ 'x1', 'x4' ]: ", np.sqrt(-cross val score(linreg, X, y, cv=10, scoring='neg mean squared er
ror')).mean())
feature cols = [ 'x1', 'x4', 'x5']
X = df[feature_cols]
print("Mean of rmse [ 'x1', 'x4', 'x5']: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squar
ed error')).mean())
feature_cols = [ 'x1', 'x3' ]
X = df[feature cols]
 print("Mean of rmse [ 'x1', 'x3' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squared_er, but it is not also as a simple of the context of the 
ror')).mean())
feature cols = [ 'x1', 'x3', 'x5' ]
X = df[feature cols]
print("Mean of rmse [ 'x1', 'x3', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squa
red error')).mean())
feature_cols = [ 'x1', 'x3', 'x4' ]
X = df[feature_cols]
print("Mean of rmse [ 'x1', 'x3', 'x4' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squa
red error')).mean())
feature cols = [ 'x1', 'x3', 'x4', 'x5' ]
X = df[feature_cols]
print("Mean of rmse [ 'x1', 'x3', 'x4', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mea
n_squared_error')).mean())
feature_cols = [ 'x1', 'x2' ]
X = df[feature cols]
print("Mean of rmse [ 'x1', 'x2' ]: ", np.sqrt(-cross val score(linreg, X, y, cv=10, scoring='neg mean squared er
ror')).mean())
feature_cols = [ 'x1', 'x2', 'x5' ]
X = df[feature_cols]
print("Mean of rmse [ 'x1', 'x2', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squa
red_error')).mean())
feature cols = [ 'x1', 'x2', 'x4' ]
X = df[feature cols]
print("Mean of rmse [ 'x1', 'x2', 'x4' ]: ", np.sqrt(-cross val score(linreg, X, y, cv=10, scoring='neg mean squa
red error')).mean())
feature cols = [ 'x1', 'x2', 'x4', 'x5' ]
X = df[feature_cols]
print("Mean of rmse [ 'x1', 'x2', 'x4', 'x5' ]: ", np.sqrt(-cross val score(linreg, X, y, cv=10, scoring='neg mea
n_squared_error')).mean())
feature_cols = [ 'x1', 'x2', 'x3' ]
X = df[feature cols]
print("Mean of rmse [ 'x1', 'x2', 'x3' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mean_squa
red error')).mean())
feature_cols = [ 'x1', 'x2', 'x3', 'x5' ]
```

```
print("Mean of rmse [ 'x1', 'x2', 'x3', 'x5' ]: ", np.sqrt(-cross val score(linreg, X, y, cv=10, scoring='neg mea
n squared error')).mean())
feature_cols = [ 'x1', 'x2', 'x3', 'x4' ]
X = df[feature cols]
print("Mean of rmse [ 'x1', 'x2', 'x3', 'x4' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='neg_mea
n squared error')).mean())
feature cols = [ 'x1', 'x2', 'x3', 'x4', 'x5' ]
X = df[feature_cols]
print("Mean of rmse [ 'x1', 'x2', 'x3', 'x4', 'x5' ]: ", np.sqrt(-cross_val_score(linreg, X, y, cv=10, scoring='n
eg_mean_squared_error')).mean())
Mean of rmse [ 'x5' ]: 1233.816965091099
Mean of rmse [ 'x4' ]: 1262.2824118519077
                    'x4', 'x5']: 1254.3868143284667
Mean of rmse [
                    'x3']: 1044.8402437111006
Mean of rmse [
                    'x3', 'x5' ]: 1045.5269498128887
'x3', 'x4' ]: 1073.407916205778
Mean of rmse [
Mean of rmse [
                    'x3', 'x4', 'x5']: 1074.0602249108588
Mean of rmse [
Mean of rmse [
                    'x2' ]: 1206.1829154024713
                    'x2', 'x5' ]: 1198.30099/3312
'x2', 'x4' ]: 1224.3957612088545
Mean of rmse [
Mean of rmse [
Mean of rmse [ 'x2', 'x4', 'x5' ]: 1216.4591428441017
Mean of rmse [ 'x2', 'x3' ]: 963.3214406915713

Mean of rmse [ 'x2', 'x3', 'x5' ]: 962.5889085840488

Mean of rmse [ 'x2', 'x3', 'x4' ]: 992.1879646685354

Mean of rmse [ 'x2', 'x3', 'x4', 'x5' ]: 991.8323083830168
                    'x1']: 1252.0266242901002
'x1', 'x5']: 1243.8427178517418
Mean of rmse [
Mean of rmse [
Mean of rmse [ 'x1', 'x4' ]: 1275.1254642411577
Mean of rmse [ 'x1', 'x4', 'x5']: 1266.5025472346201

Mean of rmse [ 'x1', 'x3' ]: 1058.7370593626356

Mean of rmse [ 'x1', 'x3', 'x5' ]: 1058.5477092551932
Mean of rmse [ 'x1', 'x3', 'x4' ]: 1089.3511823080048
                   'x1', 'x3', 'x4', 'x5']: 1089.4269954
'x1', 'x2']: 1214.777753151737
'x1', 'x2', 'x5']: 1206.7205443053754
                                    'x4', 'x5']: 1089.4269954875585
Mean of rmse [
Mean of rmse [
Mean of rmse [
Mean of rmse [ 'x1', 'x2', 'x4' ]: 1234.172616251501
Mean of rmse [ 'x1', 'x2', 'x4', 'x5' ]: 1225.8884062809059

Mean of rmse [ 'x1', 'x2', 'x4', 'x5' ]: 969.9825044197854

Mean of rmse [ 'x1', 'x2', 'x3' ]: 969.5297732894338
Mean of rmse [ 'x1', 'x2', 'x3', 'x5' ]: 969.5297732894338
Mean of rmse [ 'x1', 'x2', 'x3', 'x4' ]: 1000.131410389532
Mean of rmse [ 'x1', 'x2', 'x3', 'x4', 'x5' ]: 1000.0719831858908
```

now y mlr contains my Multiple Linear Regression predictions

In [13]:

X = df[feature_cols]

```
feature cols = [ 'x2', 'x3', 'x5' ]
X = df[feature cols]
v = df.Y
X test = willbepredicted[feature cols]
linreg.fit(X, y)
y_mlr = linreg.predict(X_test)
```

after that I've tried decision tree regression, random forest regression and lasso method.

In [14]:

```
X = df[feature_cols]
y = df.Y
x_find = willbepredicted[feature_cols]
# % Train Test Split
X_train, X_test, y_train, y_test = train_test_split(X, y,
test size=0.20, random state=0)
```

In [15]:

```
from sklearn.preprocessing import PolynomialFeatures
for i in range(1, 7):
    poly_reg = PolynomialFeatures(degree = i)
    X_poly = poly_reg.fit_transform(X)
    X_train_poly = poly_reg.fit_transform(X_train)
    lin_reg = LinearRegression()
    lin_reg.fit(X_train_poly, y_train)
    pol_scores = np.sqrt(-cross_val_score(lin_reg, X_poly, y, cv = 10, scoring='neg_mean_squared_error'))
    print("mean cross validation score(degree {}): {}".format(i,np.mean(pol_scores)))
    print("RMSE without CV(degree {}): {}".format(i,np.sqrt(np.sum(np.square(y_test - lin_reg.predict(poly_reg.fit_transform(X_test))))/
    print("")

mean cross validation score(degree 1): 962.5889085840487
RMSE without CV(degree 1): 828.3654902357964
```

```
mean cross validation score(degree 1): 962.5889085840487 RMSE without CV(degree 1): 828.3654902357964

mean cross validation score(degree 2): 929.3727870100718 RMSE without CV(degree 2): 847.7239333513262

mean cross validation score(degree 3): 994.7451650509689 RMSE without CV(degree 3): 771.7619208312487

mean cross validation score(degree 4): 1923.619778983891 RMSE without CV(degree 4): 1171.0306116996853

mean cross validation score(degree 5): 4671.6716115755935 RMSE without CV(degree 5): 4169.884535249694

mean cross validation score(degree 6): 37001.83191438672 RMSE without CV(degree 6): 28568.331650381257
```

RMSE increases after degree 3, so I choose degree 3 for my polynomial model

In [16]:

```
poly_reg = PolynomialFeatures(degree = 3)
X_poly = poly_reg.fit_transform(X)
lin_reg = LinearRegression()
lin_reg.fit(X_poly, y)
y_pol = lin_reg.predict(poly_reg.fit_transform(x_find))
```

```
In [17]:
# %% Decision Tree Regressor Model
dt = DecisionTreeRegressor(random_state=0, criterion="mse")
dt_fit = dt.fit(X_train, y_train)
# dt scores now consist RMSE
dt_scores = np.sqrt(-cross_val_score(dt_fit, X, y, cv = 10, scoring='neg_mean_squared_error'))
print("mean cross validation score: {}".format(np.mean(dt_scores)))
# RMSE without cross-validation
print("score without cv: {}".format(np.sqrt(np.sqm(np.square(y test - dt fit.predict(X test))))/len(X test))))
# GridSearchCV
from sklearn.metrics import mean squared error
scoring = make scorer(mean squared error)
g_cv = GridSearchCV(DecisionTreeRegressor(random_state=0),
              param_grid={'min_samples_split': range(2, 10)},
              scoring=scoring, cv=10, refit=True)
g cv.fit(X_train, y_train)
g_cv.best_params_
result = g_cv.cv_results_
# RMSE of GridSearchCV
print("RMSE of best estimator: {}".format(np.sqrt(np.sqm(np.square(y_test - g_cv.best_estimator_.predict(X_test))
)/len(X test))))
# fit all data before prediction
g_cv.fit(X, y)
# Prediction of last 20 row by Decision Tree Regressor Model
y_dtr = g_cv.best_estimator_.predict(x_find)
mean cross validation score: 547.5454703092869
score without cv: 853.5270353070254
RMSE of best estimator: 853.5270353070254
In [18]:
# %% Random Forest Regression
rfr = RandomForestRegressor(random state=0, criterion="mse")
rfr_fit = rfr.fit(X_train, y_train)
```

```
# rfr scores now consist RMSE
rfr_scores = np.sqrt(-cross_val_score(rfr_fit, X, y, cv = 10, scoring='neg_mean_squared_error'))
print("mean cross validation score: {}".format(np.mean(rfr scores)))
# RMSE without cross-validation
print("score without cv: {}".format(np.sqrt(np.sum(np.square(y test - rfr fit.predict(X test))))/len(X test))))
# RMSE of GridSearchCV
from sklearn.metrics import mean_squared_error
param grid = {
     'n estimators': [1000]
scoring = make_scorer(mean_squared_error)
g_cv = GridSearchCV(RandomForestRegressor(random_state=0),
              param_grid=param_grid,
              scoring=scoring, cv=10, refit=True)
g cv.fit(X_train, y_train)
g_cv.best_params_
result = g cv.cv results
# print(result)
print("RMSE \ of \ best \ estimator: \ \{\}".format(np.sqrt(np.sum(np.square(y\_test \ - \ g\_cv.best\_estimator\_.predict(X\_test)) \} \\
)/len(X test))))
# fit all data before prediction
g cv.fit(X, y)
y_rfr = g_cv.best_estimator_.predict(x_find)
```

mean cross validation score: 497.7086291841295 score without cv: 572.635035166379 RMSE of best estimator: 585.0795790109667

In [19]:

```
# %% Lasso
from sklearn.linear_model import LassoCV
reg = LassoCV(cv=5, random_state=0).fit(X_train, y_train)
lasso_scores = np.sqrt(-cross_val_score(reg, X, y, cv = 10, scoring='neg_mean_squared_error'))
print("mean cross validation score: {}".format(np.mean(lasso_scores)))

# RMSE without cross-validation
print("score without cv: {}".format(np.sqrt(np.sum(np.square(y_test - reg.predict(X_test)))/len(X_test))))
# fit all data before prediction
reg.fit(X, y)
y_lasso = reg.predict(x_find)
```

mean cross validation score: 962.9765376122477 score without cv: 828.0371207159719