RegularizationAndGradientDescent

April 4, 2023

1 Polynomial Features and Regularization

Regularization and Gradient Descent

We will begin with a short tutorial on regression, polynomial features, and regularization based on a very simple, sparse data set that contains a column of x data and associated y noisy data. The data file is called $X_Y_Sinusoid_Data.csv$.

Task1

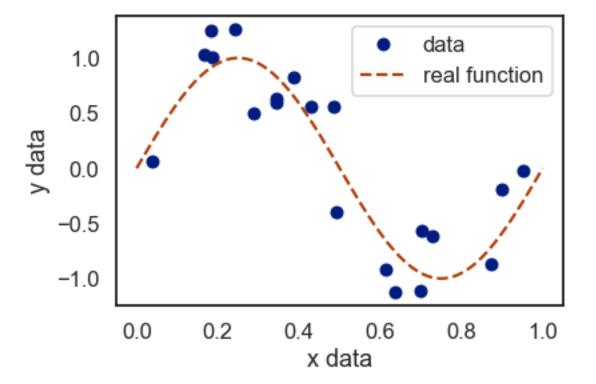
Import the data.

Also generate approximately 100 equally spaced x data points over the range of 0 to 1. Using these points, calculate the y-data which represents the "ground truth" (the real function) from the equation: $y = \sin(2\pi x)$

Plot the sparse data (x vs y) and the calculated ("real") data.

```
[1]: # Surpress warnings from using older version of sklearn:
    def warn(*args, **kwargs):
        pass
    import warnings
    warnings.warn = warn
```

```
[5]: x y
0 0.038571 0.066391
1 0.166776 1.027483
2 0.183153 1.245302
3 0.187359 1.004781
4 0.243116 1.264121
```



Task2

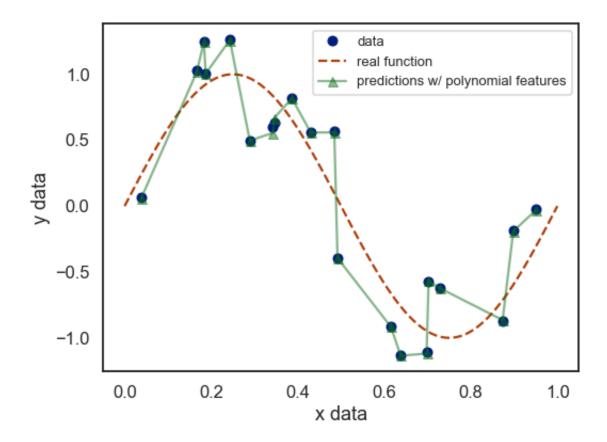
Using the PolynomialFeatures class from Scikit-learn's preprocessing library, create 20th order polynomial features.

Fit this data using linear regression.

Plot the resulting predicted value compared to the calculated data.

Note that PolynomialFeatures requires either a dataframe (with one column, not a Series) or a 2D array of dimension (X, 1), where X is the length.

```
[15]: from sklearn.preprocessing import PolynomialFeatures
      from sklearn.linear_model import LinearRegression
      # Setup the polynomial features
      degree = 20
      pf = PolynomialFeatures(degree)
      lr = LinearRegression()
      \# Extract the X- and Y- data from the dataframe
      X_data = data[['x']]
      Y_data = data['y']
      # Create the features and fit the model
      X_poly = pf.fit_transform(X_data)
      lr = lr.fit(X_poly, Y_data)
      Y_pred = lr.predict(X_poly)
      # Plot the result
      plt.figure(figsize=(8, 6))
      plt.plot(X_data, Y_data, marker='o', ls='', label='data', alpha=1)
      plt.plot(X_real, Y_real, ls='--', label='real function')
      plt.plot(X_data, Y_pred, marker='^', alpha=.5, label='predictions w/ polynomial_
       ⇔features')
     plt.legend(loc='upper right', bbox_to_anchor=(1.0, 1.0), fontsize=13)
      ax = plt.gca()
      ax.set(xlabel='x data', ylabel='y data');
```



Perform the regression on using the data with polynomial features using ridge regression (α =0.001) and lasso regression (α =0.0001).

Plot the results, as was done in Task1.

Also plot the magnitude of the coefficients obtained from these regressions, and compare them to those obtained from linear regression in the previous question. The linear regression coefficients will likely need a separate plot (or their own y-axis) due to their large magnitude.

What does the comparatively large magnitude of the data tell you about the role of regularization?

```
[18]: # Mute the sklearn warning about regularization
import warnings
warnings.filterwarnings('ignore', module='sklearn')

from sklearn.linear_model import Ridge, Lasso

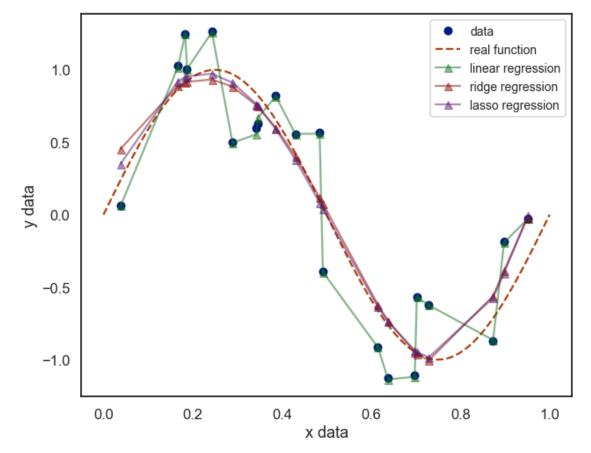
# The ridge regression model
rr = Ridge(alpha=0.001)
rr = rr.fit(X_poly, Y_data)
Y_pred_rr = rr.predict(X_poly)
```

```
# The lasso regression model
lassor = Lasso(alpha=0.0001)
lassor = lassor.fit(X_poly, Y_data)
Y_pred_lr = lassor.predict(X_poly)

# The plot of the predicted values
plt.figure(figsize=(10, 8))

plt.plot(X_data, Y_data, marker='o', ls='', label='data')
plt.plot(X_real, Y_real, ls='--', label='real function')
plt.plot(X_data, Y_pred, label='linear regression', marker='^', alpha=.5)
plt.plot(X_data, Y_pred_rr, label='ridge regression', marker='^', alpha=.5)
plt.plot(X_data, Y_pred_lr, label='lasso regression', marker='^', alpha=.5)
plt.legend(loc='upper right', bbox_to_anchor=(1.0, 1.0), fontsize=14)

ax = plt.gca()
ax.set(xlabel='x data', ylabel='y data');
```



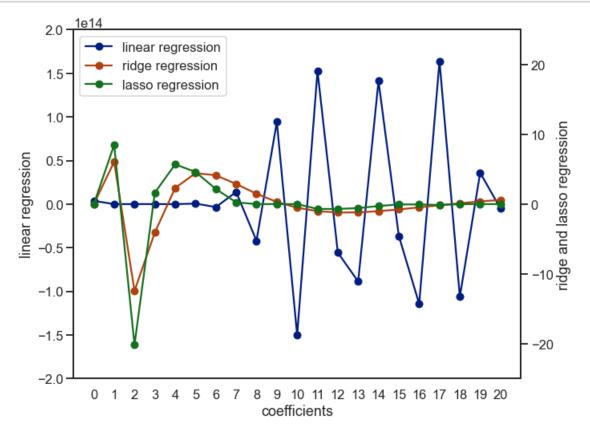
```
[19]: # let's look at the absolute value of coefficients for each model
      coefficients = pd.DataFrame()
      coefficients['linear regression'] = lr.coef_.ravel()
      coefficients['ridge regression'] = rr.coef_.ravel()
      coefficients['lasso regression'] = lassor.coef_.ravel()
      coefficients = coefficients.applymap(abs)
      coefficients.describe() # Huge difference in scale between non-regularized vs_
       ⇔regularized regression
[19]:
             linear regression ridge regression lasso regression
                  2.100000e+01
                                       21.000000
                                                         21.000000
      count
     mean
                  5.754304e+13
                                        2.169397
                                                          2.167284
     std
                  5.999233e+13
                                        2.900278
                                                          4.706731
     min
                  1.611590e+07
                                        0.000000
                                                          0.000000
     25%
                  3.403676e+12
                                        0.467578
                                                          0.000000
     50%
                  3.649017e+13
                                        1.017272
                                                          0.252181
     75%
                  1.061917e+14
                                        2.883507
                                                          1.641353
     max
                  1.639660e+14
                                       12.429635
                                                         20.176708
[22]: import matplotlib.pyplot as plt
      import seaborn as sns
      colors = sns.color_palette()
      # Setup the dual y-axes
      fig, ax1 = plt.subplots(figsize=(10, 8))
      ax2 = ax1.twinx()
      # Plot the linear regression data
      ax1.plot(lr.coef_.ravel(),
               color=colors[0], marker='o', label='linear regression')
      # Plot the regularization data sets
      ax2.plot(rr.coef_.ravel(),
               color=colors[1], marker='o', label='ridge regression')
      ax2.plot(lassor.coef_.ravel(),
               color=colors[2], marker='o', label='lasso regression')
      # Customize axes scales
      ax1.set_ylim(-2e14, 2e14)
      ax2.set_ylim(-25, 25)
      # Combine the legends
```

h1, l1 = ax1.get_legend_handles_labels()

```
h2, 12 = ax2.get_legend_handles_labels()
ax1.legend(h1+h2, 11+12)

ax1.set(xlabel='coefficients', ylabel='linear regression')
ax2.set(ylabel='ridge and lasso regression')

ax1.set_xticks(range(len(lr.coef_)));
```



For the remaining questions, we will be working with the data set, which is based on housing prices in Ames, Iowa. There are an extensive number of features—see the exercises from week three for a discussion of these features.

To begin:

Import the data with Pandas, remove any null values, and one hot encode categoricals. Either Scikit-learn's feature encoders or Pandas <code>get_dummies</code> method can be used.

Split the data into train and test sets.

Log transform skewed features.

Scaling can be attempted, although it can be interesting to see how well regularization works

without scaling features.

[23]: data = pd.read_csv("https://cf-courses-data.s3.us.cloud-object-storage.

appdomain.cloud/IBM-ML240EN-SkillsNetwork/labs/data/Ames_Housing_Sales.csv")

data.head(10)

[00]		1 -+ E1 CE O	4E1 GE	20 D1-	477	- D-		. A 1 C	D1 -1 T	D+ C		
[23]:	^		ndFlrSF	3SsnPorch	-		earoom					
	0	856.0	854.0	0.0				3	1Fam		ΓA ΓA	
	1	1262.0	0.0	0.0				3	1Fam		ΓA	
	2	920.0	866.0	0.0				3	1Fam		ΓA	
	3	961.0	756.0	0.0				3	1Fam		d	
	4	1145.0	1053.0	0.0				4	1Fam		Γ A	
	5	796.0	566.0	320.0				1	1Fam		ΓA	
	6	1694.0	0.0	0.0				3	1Fam		ΓΑ	
	7	1107.0	983.0	0.0				3	1Fam		ΓΑ	
	8	1022.0	752.0	0.0				2	1Fam			
	9	1077.0	0.0	0.0	None	Э		2	2fmCon	Ί	ΓA	
		BsmtExposure	BsmtFir	ıSF1 Bsmt	BsmtFinSF2		ScreenPorch		Street	TotRmsA	AbvGrd	\
	0	No		06.0	0.0			0.0			8	
	1	Gd		8.0	0.0			0.0			6	
	2	Mn		86.0	0.0			0.0			6	
	3	No		6.0	0.0			0.0			7	
	4	Av		55.0	0.0			0.0			9	
	5	No		32.0	0.0			0.0			5	
	6	Av		9.0	0.0			0.0			7	
	7	Mn		9.0	32.0)		0.0) Pave		7	
	8	None		0.0	0.0)		0.0			8	
	9	No	85	51.0	0.0)		0.0) Pave		5	
		TotalBsmtSF	Utilitie	s WoodDe	ckSF Y	/earE	Built	YearRe	emodAdd Y:	rSold Sa	lePri	ce
	0	856.0	AllPu	ıb	0.0		2003		2003	2008 2	208500	.0
	1	1262.0	AllPu	ıb 2	98.0		1976		1976	2007 1	181500	.0
	2	920.0	AllPu	ıb	0.0		2001		2002	2008 2	223500	.0
	3	756.0	AllPu	ıb	0.0		1915		1970	2006 1	40000	.0
	4	1145.0	AllPu	ıb 1	92.0		2000		2000	2008 2	250000	.0
	5	796.0	AllPu	ıb	40.0		1993		1995	2009 1	43000	.0
	6	1686.0	AllPu	ıb 2	55.0		2004		2005	2007 3	307000	.0
	7	1107.0	AllPu	ıb 2	35.0		1973		1973	2009 2	200000	.0
	8	952.0	AllPu	ıb	90.0		1931		1950	2008 1	29900	.0
	9	991.0	AllPu	ıb	0.0		1939		1950	2008 1	18000	.0

[10 rows x 80 columns]

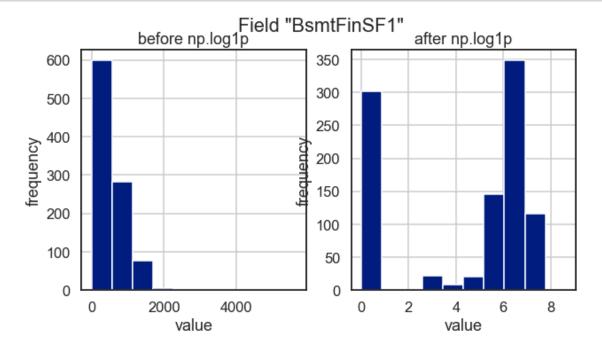
Create a list of categorial data and one-hot encode. Pandas one-hot encoder (get_dummies) works well with data that is defined as a categorical.

```
[25]: # Get a Pd. Series consisting of all the string categoricals
      one_hot_encode_cols = data.dtypes[data.dtypes == object] # filtering by string_
       \hookrightarrow categoricals
      \verb|one_hot_encode_cols = one_hot_encode_cols.index.tolist()| \textit{# list of categorical}|
       \hookrightarrow fields
      # Here we see another way of one-hot-encoding:
      \# Encode these columns as categoricals so one hot encoding works on split data_\sqcup
       \hookrightarrow (if desired)
      for col in one hot encode cols:
          data[col] = pd.Categorical(data[col])
      # Do the one hot encoding
      data = pd.get_dummies(data, columns=one_hot_encode_cols)
[26]: # split the data in train and test data sets.
      from sklearn.model_selection import train_test_split
      train, test = train_test_split(data, test_size=0.3, random_state=42)
     There are a number of columns that have skewed features – a log transformation can be applied to
     them. Note that this includes the SalePrice, our predictor. However, let's keep that one as is.
[27]: # Create a list of float colums to check for skewing
      mask = data.dtypes == float
      float_cols = data.columns[mask]
[28]: skew_limit = 0.75
      skew_vals = train[float_cols].skew()
      skew_cols = (skew_vals
                    .sort_values(ascending=False)
                    .to_frame()
                    .rename(columns={0:'Skew'})
                    .query('abs(Skew) > {0}'.format(skew_limit)))
      skew cols
[28]:
                            Skew
      MiscVal
                      26.915364
      PoolArea
                      15.777668
      LotArea
                      11.501694
      LowQualFinSF
                      11.210638
      3SsnPorch
                      10.150612
      ScreenPorch
                       4.599803
      BsmtFinSF2
                       4.466378
```

EnclosedPorch 3.218303

```
LotFrontage
                3.138032
MasVnrArea
                2.492814
OpenPorchSF
                2.295489
SalePrice
                2.106910
BsmtFinSF1
                2.010766
TotalBsmtSF
                1.979164
1stFlrSF
                1.539692
GrLivArea
                1.455564
WoodDeckSF
                1.334388
BsmtUnfSF
                0.900308
GarageArea
                0.838422
2ndFlrSF
                0.773655
```

Transform all the columns where the skew is greater than 0.75, excluding "SalePrice".



```
[30]: # Mute the setting wtih a copy warnings
pd.options.mode.chained_assignment = None

for col in skew_cols.index.tolist():
    if col == "SalePrice":
        continue
    train[col] = np.log1p(train[col])
    test[col] = test[col].apply(np.log1p) # same thing
```

```
[31]: # Separate features from predictor.
feature_cols = [x for x in train.columns if x != 'SalePrice']
X_train = train[feature_cols]
y_train = train['SalePrice']

X_test = test[feature_cols]
y_test = test['SalePrice']
```

Write a function rmse that takes in truth and prediction values and returns the root-mean-squared error. Use sklearn's mean_squared_error.

```
[32]: from sklearn.metrics import mean_squared_error

def rmse(ytrue, ypredicted):
    return np.sqrt(mean_squared_error(ytrue, ypredicted))
```

Fit a basic linear regression model

print the root-mean-squared error for this model

plot the predicted vs actual sale price based on the model.

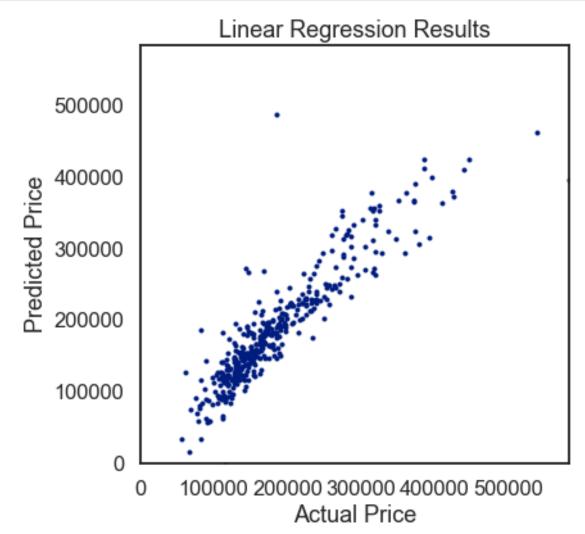
```
[33]: from sklearn.linear_model import LinearRegression

linearRegression = LinearRegression().fit(X_train, y_train)

linearRegression_rmse = rmse(y_test, linearRegression.predict(X_test))

print(linearRegression_rmse)
```

306369.6834231985



Ridge regression uses L2 normalization to reduce the magnitude of the coefficients. This can be helpful in situations where there is high variance. The regularization functions in Scikit-learn each contain versions that have cross-validation built in.

Fit a regular (non-cross validated) Ridge model to a range of α values and plot the RMSE using the cross validated error function you created above.

Use

[0.005, 0.05, 0.1, 0.3, 1, 3, 5, 10, 15, 30, 80]

as the range of alphas.

Then repeat the fitting of the Ridge models using the range of α values from the prior section. Compare the results.

Now for the RidgeCV method. It's not possible to get the alpha values for the models that weren't selected, unfortunately. The resulting error values and α values are very similar to those obtained above.

ridgeCV alpha: 15.0

ridgeCV rmse: 32169.17620567245

Task7

Much like the RidgeCV function, there is also a LassoCV function that uses an L1 regularization function and cross-validation. L1 regularization will selectively shrink some coefficients, effectively performing feature elimination.

The LassoCV function does not allow the scoring function to be set. However, the custom error function (rmse) created above can be used to evaluate the error on the final model.

Similarly, there is also an elastic net function with cross validation, ElasticNetCV, which is a combination of L2 and L1 regularization.

Fit a Lasso model using cross validation and determine the optimum value for α and the RMSE using the function created above. Note that the magnitude of α may be different from the Ridge model.

Repeat this with the Elastic net model.

Compare the results via table and/or plot.

Use the following alphas: [1e-5, 5e-5, 0.0001, 0.0005]

```
lassoCV_rmse = rmse(y_test, lassoCV.predict(X_test))
print(lassoCV.alpha_, lassoCV_rmse) # Lasso is slower
```

0.0005 39257.393991449186

We can determine how many of these features remain non-zero.

```
[39]: print('Of {} coefficients, {} are non-zero with Lasso.'.format(len(lassoCV. coef_), len(lassoCV. coef_.nonzero()[0])))
```

Of 294 coefficients, 273 are non-zero with Lasso.

Now try the elastic net, with the same alphas as in Lasso, and l1_ratios between 0.1 and 0.9

elasticNetCV alpha: 0.0005 elasticNetCV l1_ratio: 0.1

elasticNetCV rmse: 35001.23429607459

Comparing the RMSE calculation from all models is easiest in a table.

```
[42]: rmse_vals = [linearRegression_rmse, ridgeCV_rmse, lassoCV_rmse, uselasticNetCV_rmse]

labels = ['Linear', 'Ridge', 'Lasso', 'ElasticNet']

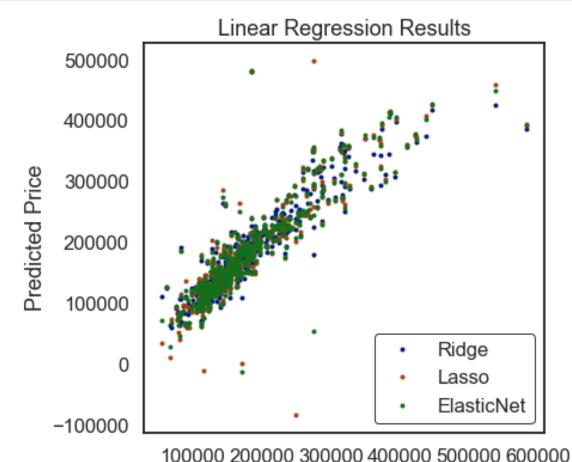
rmse_df = pd.Series(rmse_vals, index=labels).to_frame()

rmse_df.rename(columns={0: 'RMSE'}, inplace=1)

rmse_df
```

```
[42]: RMSE
Linear 306369.683423
Ridge 32169.176206
Lasso 39257.393991
ElasticNet 35001.234296
```

We can also make a plot of actual vs predicted housing prices as before.



Actual Price

Let's explore Stochastic gradient descent in this exercise.

Recall that Linear models in general are sensitive to scaling. However, SGD is *very* sensitive to scaling.

Moreover, a high value of learning rate can cause the algorithm to diverge, whereas a too low value may take too long to converge.

Fit a stochastic gradient descent model without a regularization penalty (the relevant parameter is penalty).

Now fit stochastic gradient descent models with each of the three penalties (L2, L1, Elastic Net) using the parameter values determined by cross validation above.

Do not scale the data before fitting the model.

Compare the results to those obtained without using stochastic gradient descent.

```
[44]: # Import SGDRegressor and prepare the parameters
      from sklearn.linear_model import SGDRegressor
      model_parameters_dict = {
          'Linear': {'penalty': 'none'},
          'Lasso': {'penalty': '12',
                 'alpha': lassoCV.alpha },
          'Ridge': {'penalty': 'l1',
                 'alpha': ridgeCV_rmse},
          'ElasticNet': {'penalty': 'elasticnet',
                         'alpha': elasticNetCV.alpha_,
                         'l1_ratio': elasticNetCV.l1_ratio_}
      }
      new_rmses = {}
      for modellabel, parameters in model_parameters_dict.items():
          # following notation passes the dict items as arguments
          SGD = SGDRegressor(**parameters)
          SGD.fit(X_train, y_train)
          new_rmses[modellabel] = rmse(y_test, SGD.predict(X_test))
      rmse_df['RMSE-SGD'] = pd.Series(new_rmses)
      rmse df
```

```
[44]: RMSE RMSE-SGD
Linear 306369.683423 1.074298e+16
Ridge 32169.176206 6.422933e+15
Lasso 39257.393991 4.864477e+15
```

```
ElasticNet 35001.234296 8.694994e+15
```

Notice how high the error values are! The algorithm is diverging. This can be due to scaling and/or learning rate being too high. Let's adjust the learning rate and see what happens.

Pass in eta0=1e-7 when creating the instance of SGDClassifier.

Re-compute the errors for all the penalties and compare.

```
[45]: # Import SGDRegressor and prepare the parameters
      from sklearn.linear_model import SGDRegressor
      model_parameters_dict = {
          'Linear': {'penalty': 'none'},
          'Lasso': {'penalty': '12',
                 'alpha': lassoCV.alpha },
          'Ridge': {'penalty': 'l1',
                 'alpha': ridgeCV rmse},
          'ElasticNet': {'penalty': 'elasticnet',
                         'alpha': elasticNetCV.alpha_,
                         'l1 ratio': elasticNetCV.l1 ratio }
      }
      new_rmses = {}
      for modellabel, parameters in model_parameters_dict.items():
          # following notation passes the dict items as arguments
          SGD = SGDRegressor(eta0=1e-7, **parameters)
          SGD.fit(X_train, y_train)
          new_rmses[modellabel] = rmse(y_test, SGD.predict(X_test))
      rmse_df['RMSE-SGD-learningrate'] = pd.Series(new_rmses)
      rmse df
```

```
[45]:
                           RMSE
                                     RMSE-SGD
                                               RMSE-SGD-learningrate
     Linear
                  306369.683423 1.074298e+16
                                                        78006.308526
     Ridge
                   32169.176206 6.422933e+15
                                                        76905.859017
     Lasso
                   39257.393991
                                 4.864477e+15
                                                        81633.200993
     ElasticNet
                   35001.234296 8.694994e+15
                                                        72000.166118
```

Now let's scale our training data and try again.

Fit a MinMaxScaler to X train create a variable X train scaled.

Using the scaler, transform X_test and create a variable X_test_scaled.

Apply the same versions of SGD to them and compare the results. Don't pass in a eta0 this time.

```
[46]: from sklearn.preprocessing import MinMaxScaler scaler = MinMaxScaler()
```

```
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)

new_rmses = {}
for modellabel, parameters in model_parameters_dict.items():
    # following notation passes the dict items as arguments
    SGD = SGDRegressor(**parameters)
    SGD.fit(X_train_scaled, y_train)
    new_rmses[modellabel] = rmse(y_test, SGD.predict(X_test_scaled))

rmse_df['RMSE-SGD-scaled'] = pd.Series(new_rmses)
rmse_df
```

[46]:		RMSE	RMSE-SGD	RMSE-SGD-learningrate	\
	Linear	306369.683423	1.074298e+16	78006.308526	
	Ridge	32169.176206	6.422933e+15	76905.859017	
	Lasso	39257.393991	4.864477e+15	81633.200993	
	ElasticNet	35001.234296	8.694994e+15	72000.166118	
		RMSE-SGD-scale	d		
	Linear	32785.21134	1		

Lasso 32791.985461 ElasticNet 32881.304540

77776.160547

Ridge