## Regularization and Gradient Descent Exercises



# Learning Objectives

- Explain cost functions, regularization, feature selection, and hyper-parameters
- Summarize complex statistical optimization algorithms like gradient descent and its application to linear regression
- Apply Intel® Extension for Scikit-learn\* to leverage underlying compute capabilities of hardware

#### scikit-learn\*

Frameworks provide structure that Data Scientists use to build code. Frameworks are more than just libraries, because in addition to callable code, frameworks influence how code is written.

A main virtue of using an optimized framework is that code runs faster. Code that runs faster is just generally more convenient but when we begin looking at applied data science and AI models, we can see more material benefits. Here you will see how optimization, particularly hyperparameter optimization can benefit more than just speed.

These exercises will demonstrate how to apply **the Intel® Extension for Scikit-learn\***, a seamless way to speed up your Scikit-learn application. The acceleration is achieved through the use of the Intel® oneAPI Data Analytics Library (oneDAL). Patching is the term used to extend scikit-learn with Intel optimizations and makes it a well-suited machine learning framework for dealing with real-life problems.

To get optimized versions of many Scikit-learn algorithms using a patch() approach consisting of adding these lines of code after importing sklearn:

- from sklearnex import patch\_sklearn
- patch\_sklearn()

### This exercise relies on installation of Intel® Extension for Scikit-learn\*

If you have not already done so, follow the instructions from Week 1 for instructions

#### Introduction

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.
from future import print function
import os
data_path = ['data']
from sklearn.preprocessing import PolynomialFeatures
from sklearn.linear_model import LinearRegression
from sklearn.linear_model import SGDRegressor
from sklearn.metrics import mean squared error
from sklearn.linear model import LinearRegression
from sklearn.model selection import train test split
from sklearn.linear model import Ridge, Lasso
from sklearn.linear model import RidgeCV
from sklearn.linear model import LassoCV
from sklearn.linear model import ElasticNetCV
from sklearn.linear model import SGDRegressor
from sklearn.preprocessing import MinMaxScaler
from sklearnex import patch sklearn
patch sklearn()
```

Intel(R) Extension for Scikit-learn\* enabled (https://github.com/intel/scikit-learn-intelex)

4

#### Question 1

- · 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.

```
import pandas as pd
import numpy as np

filepath = os.sep.join(data_path + ['X_Y_Sinusoid_Data.csv'])
data = pd.read_csv(filepath)

X_real = np.linspace(0, 1.0, 100)

Y_real = np.sin(2 * np.pi * X_real)

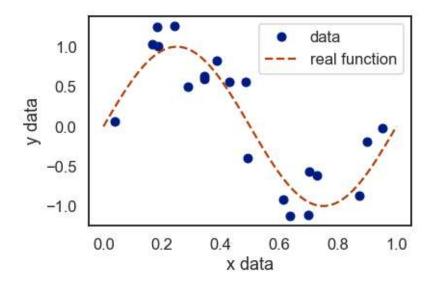
import matplotlib.pyplot as plt
import seaborn as sns

%matplotlib inline
```

```
sns.set_style('white')
sns.set_context('talk')
sns.set_palette('dark')

ax = data.set_index('x')['y'].plot(ls='', marker='o', label='data')
ax.plot(X_real, Y_real, ls='--', marker='', label='real function')

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



- 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.

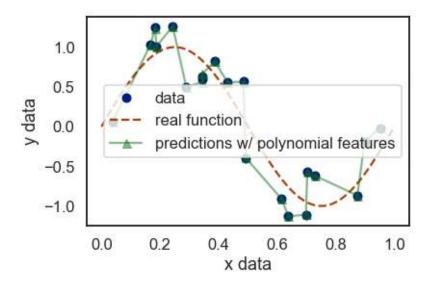
Note that PolynomialFeatures requires either a dataframe (with one column, not a Series) or a 2D array of dimension (x, 1), where x is the length.

```
degree = 20
pf = PolynomialFeatures(degree)
lr = LinearRegression()

X_data = data[['x']].to_numpy()
Y_data = data['y'].to_numpy()

X_poly = pf.fit_transform(X_data)
lr = lr.fit(X_poly, Y_data)
Y_pred = lr.predict(X_poly)

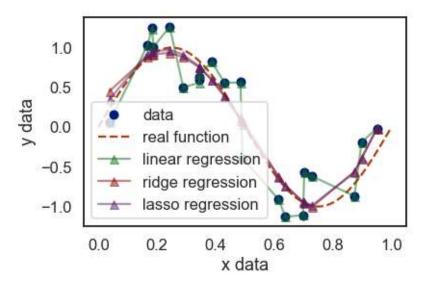
plt.plot(X_data, Y_data, marker='o', ls='', label='data', alpha=1)
plt.plot(X_real, Y_real, ls='--', label='real function')
plt.plot(X_data, Y_pred, marker='^', alpha=.5, label='predictions w/ polynomial features')
plt.legend()
ax = plt.gca()
ax.set(xlabel='x data', ylabel='y data');
```



- Perform the regression on using the data with polynomial features using ridge regression ( $\alpha$  =0.001) and lasso regression ( $\alpha$ =0.0001).
- Plot the results, as was done in Question 1.
- 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?

```
import warnings
warnings.filterwarnings('ignore', module='sklearn')
rr = Ridge(alpha=0.001)
rr = rr.fit(X poly, Y data)
Y_pred_rr = rr.predict(X_poly)
lassor = Lasso(alpha=0.0001)
lassor = lassor.fit(X poly, Y data)
Y_pred_lr = lassor.predict(X_poly)
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()
ax = plt.gca()
ax.set(xlabel='x data', ylabel='y data');
```

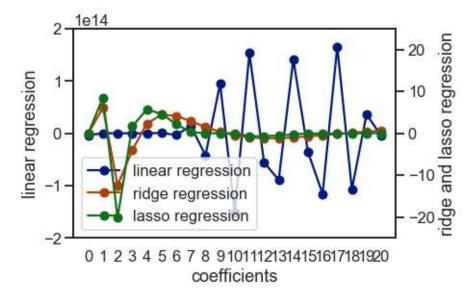


```
coefficients = pd.DataFrame()
coefficients['linear regression'] = lr.coef_.ravel()
coefficients['ridge regression'] = rr.coef_.ravel()
coefficients['lasso regression'] = lassor.coef_.ravel()
coefficients = coefficients.applymap(abs)
```

coefficients.describe() # difference in scale between non-regularized vs regularized regression

	linear regression	ridge regression	lasso regression
count	2.100000e+01	21.000000	21.000000
mean	5.783911e+13	2.169397	2.167284
std	6.038244e+13	2.900278	4.706731
min	1.622914e+07	0.000000	0.000000
25%	3.421984e+12	0.467578	0.000000
50%	3.623240e+13	1.017272	0.252181
75%	1.071032e+14	2.883507	1.641353
max	1.655707e+14	12.429635	20.176708

```
colors = sns.color_palette()
# Setup the dual y-axes
ax1 = plt.axes()
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, l1+l2)
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 <u>data set</u> from last lesson, which is based on housing prices in Ames, lowa. 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 get dummies 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.

```
filepath = os.sep.join(data_path + ['Ames_Housing_Sales.csv'])
data = pd.read_csv(filepath, sep=',')
```

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.

```
one_hot_encode_cols = data.dtypes[data.dtypes == object]
one_hot_encode_cols = one_hot_encode_cols.index.tolist()

for col in one_hot_encode_cols:
    data[col] = pd.Categorical(data[col])

data = pd.get_dummies(data, columns=one_hot_encode_cols)
```

Next, split the data in train and test data sets.

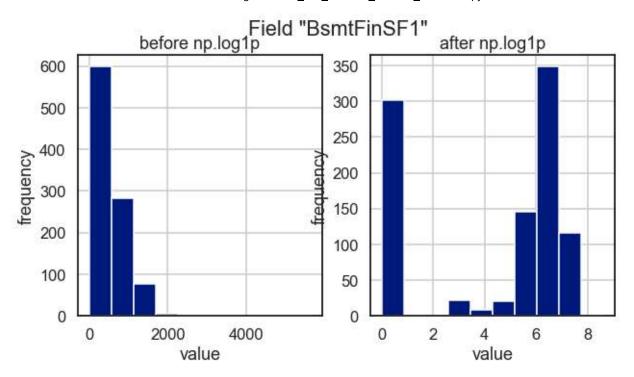
```
train, test = train_test_split(data, test_size=0.3, random_state=42)
```

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.

	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".

```
field = "BsmtFinSF1"
fig, (ax_before, ax_after) = plt.subplots(1, 2, figsize=(10, 5))
train[field].hist(ax=ax_before)
train[field].apply(np.log1p).hist(ax=ax_after)
ax_before.set(title='before np.log1p', ylabel='frequency', xlabel='value')
ax_after.set(title='after np.log1p', ylabel='frequency', xlabel='value')
fig.suptitle('Field "{}"'.format(field));
```



```
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
```

Separate features from predictor.

```
feature_cols = [x for x in train.columns if x != 'SalePrice']
X_train = train[feature_cols].to_numpy()
y_train = train['SalePrice'].to_numpy()

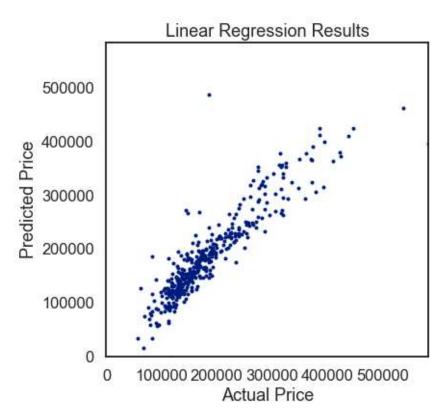
X_test = test[feature_cols].to_numpy()
y_test = test['SalePrice'].to_numpy()
```

#### Question 5

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



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  $\alpha$  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  $\alpha$  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  $\alpha$  values are very similar to those obtained above.

#### Question 7

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  $\alpha$  and the RMSE using the function created above. Note that the magnitude of  $\alpha$  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:
```

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

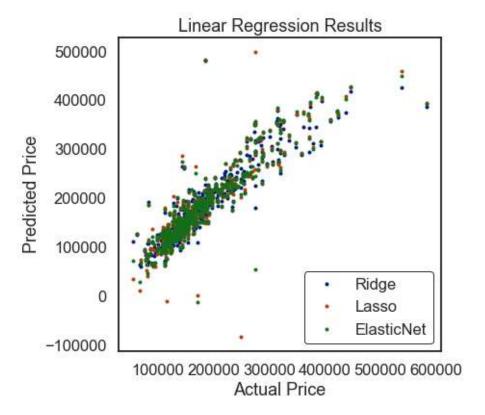
Now try the elastic net, with the same alphas as in Lasso, and I1\_ratios between 0.1 and 0.9

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

```
rmse_vals = [linearRegression_rmse, ridgeCV_rmse, lassoCV_rmse, elasticNetCV_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
```

	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.



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.

```
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():
    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)
```

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.

```
# Import SGDRegressor and prepare the parameters
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)
```