# **Ridge Regression**

Here we will try to fit the dataset with a Ridge Regression model. The steps are

- Determine a class for the model supporting methods
  - fit
  - predict
  - score
- · Search for hyperparameters through trial and error
  - evaluate the average training and validating error for each hyperparameter
- Plot the distributions of weight on the features
  - Does Ridge Regression give us sparsity
- Threshold the values to compare zero/non-zero against the weights of the target function

### In [1]:

```
import numpy as np
import pandas as pd
import itertools
import matplotlib.pyplot as plt
%matplotlib inline

from scipy.optimize import minimize

from sklearn.base import BaseEstimator, RegressorMixin
from sklearn.linear_model import Ridge
from sklearn.model_selection import GridSearchCV, PredefinedSplit
from sklearn.model_selection import ParameterGrid
from sklearn.metrics import mean_squared_error, make_scorer
from sklearn.metrics import confusion_matrix

from load_data import load_problem

PICKLE_PATH = 'lasso_data.pickle'
```

### **Dataset**

### In [2]:

```
#load data

x_train, y_train, x_val, y_val, target_fn, coefs_true, featurize = load_problem(
PICKLE_PATH)

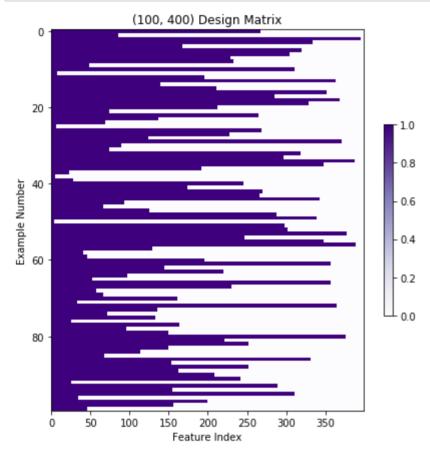
X_train = featurize(x_train)

X_val = featurize(x_val)
```

### In [6]:

```
#Visualize training data

fig, ax = plt.subplots(figsize = (7,7))
ax.set_title("({0}, {1}) Design Matrix".format(X_train.shape[0], X_train.shape[1]))
ax.set_xlabel("Feature Index")
ax.set_ylabel("Example Number")
temp = ax.imshow(X_train, cmap=plt.cm.Purples, aspect="auto")
plt.colorbar(temp, shrink=0.5);
```



## **Class for Ridge Regression**

#### In [12]:

```
class RidgeRegression(BaseEstimator, RegressorMixin):
        """ ridge regression"""
        def init__(self, l2reg=1):
                if 12req < 0:
                        raise ValueError('Regularization penalty should be at le
ast 0.')
                self.l2reg = l2reg
        def fit(self, X, y=None):
                n, num ftrs = X.shape
                # convert y to 1-dim array, in case we're given a column vector
                y = y.reshape(-1)
                def ridge obj(w):
                        predictions = np.dot(X,w)
                        residual = y - predictions
                        empirical risk = np.sum(residual**2) / n
                        12 norm squared = np.sum(w**2)
                        objective = empirical_risk + self.l2reg * l2_norm_square
d
                        return objective
                self.ridge obj = ridge obj
                w 0 = np.zeros(num ftrs)
                self.w = minimize(ridge obj, w 0).x
                return self
        def predict(self, X, y=None):
                try:
                        getattr(self, "w ")
                except AttributeError:
                        raise RuntimeError("You must train classifer before pred
icting data!")
                return np.dot(X, self.w )
        def score(self, X, y):
                # Average square error
                try:
                        getattr(self, "w_")
                except AttributeError:
                        raise RuntimeError("You must train classifer before pred
icting data!")
                residuals = self.predict(X) - y
                return np.dot(residuals, residuals)/len(y)
```

We can compare to the sklearn implementation.

#### In [13]:

```
def compare our ridge with sklearn(X train, y train, 12 reg=1):
        # Fit with sklearn -- need to multiply 12 reg by sample size, since thei
r
        # objective function has the total square loss, rather than average squa
re
        # loss.
        n = X train.shape[0]
        sklearn ridge = Ridge(alpha=n*12 reg, fit intercept=False, normalize=Fal
se)
        sklearn ridge.fit(X train, y train)
        sklearn ridge coefs = sklearn ridge.coef
        # Now run our ridge regression and compare the coefficients to sklearn's
        ridge regression estimator = RidgeRegression(12reg=12 reg)
        ridge regression estimator.fit(X train, y train)
        our coefs = ridge regression estimator.w
        print("Hoping this is very close to 0:{}".format(np.sum((our_coefs - skl
earn ridge coefs)**2)))
```

### In [14]:

```
compare_our_ridge_with_sklearn(X_train, y_train, 12_reg=1.5)
```

Hoping this is very close to 0:4.6933160195738277e-11

### **Grid Search to Tune Hyperparameter**

Now let's use sklearn to help us do hyperparameter tuning GridSearchCv.fit by default splits the data into training and validation itself; we want to use our own splits, so we need to stack our training and validation sets together, and supply an index (validation\_fold) to specify which entries are train and which are validation.

#### In [15]:

```
default params = np.unique(np.concatenate((10.**np.arange(-6,1,1), np.arange(1,3
,.3))))
def do grid search ridge(X train, y train, X val, y val, params = default params
):
        X train val = np.vstack((X train, X val))
        y_train_val = np.concatenate((y_train, y_val))
        val fold = [-1]*len(X train) + [0]*len(X val) #0 corresponds to validati
on
        param grid = [{'l2reg':params}]
        ridge regression estimator = RidgeRegression()
        grid = GridSearchCV(ridge regression estimator,
                                                return train score=True,
                                                cv = PredefinedSplit(test fold=v
al fold),
                                                refit = True,
                                                scoring = make scorer(mean squar
ed error,
greater_is_better = False))
        grid.fit(X train val, y train val)
        df = pd.DataFrame(grid.cv results )
        # Flip sign of score back, because GridSearchCV likes to maximize,
        # so it flips the sign of the score if "greater is better=FALSE"
        df['mean_test_score'] = -df['mean_test_score']
        df['mean_train_score'] = -df['mean_train_score']
        cols to keep = ["param 12reg", "mean test score", "mean train score"]
        df toshow = df[cols to keep].fillna('-')
        df toshow = df toshow.sort values(by=["param l2reg"])
        return grid, df toshow
```

### In [16]:

```
grid, results = do_grid_search_ridge(X_train, y_train, X_val, y_val)
```

# In [17]:

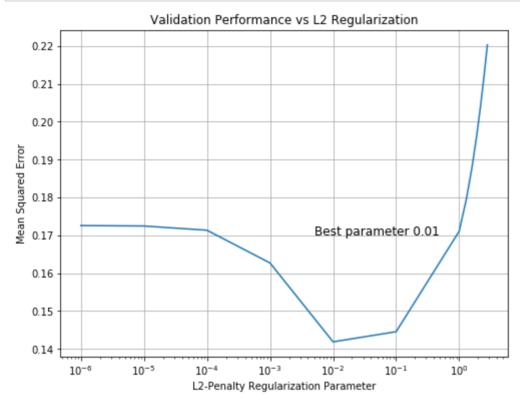
results

## Out[17]:

	param_l2reg	mean_test_score	mean_train_score
0	0.000001	0.172579	0.006752
1	0.000010	0.172464	0.006752
2	0.000100	0.171345	0.006774
3	0.001000	0.162705	0.008285
4	0.010000	0.141887	0.032767
5	0.100000	0.144566	0.094953
6	1.000000	0.171068	0.197694
7	1.300000	0.179521	0.216591
8	1.600000	0.187993	0.233450
9	1.900000	0.196361	0.248803
10	2.200000	0.204553	0.262958
11	2.500000	0.212530	0.276116
12	2.800000	0.220271	0.288422

### In [18]:

```
# Plot validation performance vs regularization parameter
fig, ax = plt.subplots(figsize = (8,6))
ax.grid()
ax.set_title("Validation Performance vs L2 Regularization")
ax.set_xlabel("L2-Penalty Regularization Parameter")
ax.set_ylabel("Mean Squared Error")
ax.semilogx(results["param_l2reg"], results["mean_test_score"])
ax.text(0.005,0.17,"Best parameter {0}".format(grid.best_params_['l2reg']), font
size = 12);
```



### **Comparing to the Target Function**

Let's plot prediction functions and compare coefficients for several fits and the target function.

Let's create a list of dicts called <code>pred\_fns</code> . Each dict has a "name" key and a "preds" key. The value corresponding to the "preds" key is an array of predictions corresponding to the input vector x. x\_train and y\_train are the input and output values for the training data

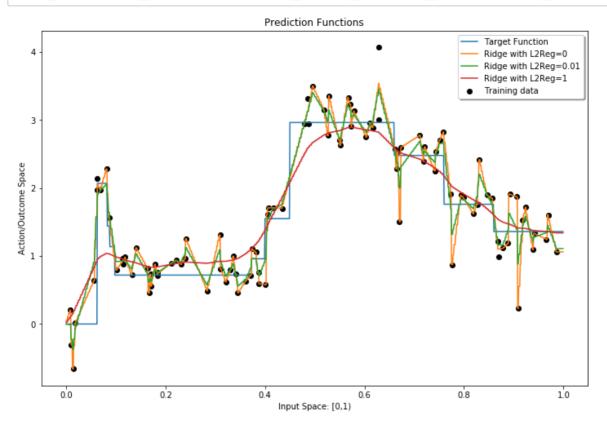
## In [19]:

### In [20]:

```
def plot_prediction_functions(x, pred_fns, x_train, y_train, legend_loc="best"):
    fig, ax = plt.subplots(figsize = (12,8))
    ax.set_xlabel('Input Space: [0,1)')
    ax.set_ylabel('Action/Outcome Space')
    ax.set_title("Prediction Functions")
    plt.scatter(x_train, y_train, color="k", label='Training data')
    for i in range(len(pred_fns)):
        ax.plot(x, pred_fns[i]["preds"], label=pred_fns[i]["name"])
    legend = ax.legend(loc=legend_loc, shadow=True)
    return fig
```

### In [21]:

plot\_prediction\_functions(x, pred\_fns, x\_train, y\_train, legend\_loc="best");



## **Visualizing the Weights**

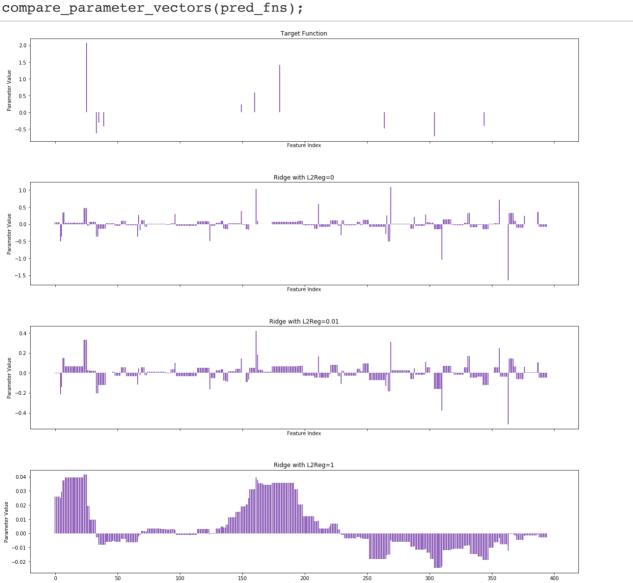
Using pred\_fns let's try to see how sparse the weights are...

#### In [22]:

```
def compare parameter vectors(pred fns):
        fig, axs = plt.subplots(len(pred fns),1, sharex=True, figsize = (20,20))
        num ftrs = len(pred fns[0]["coefs"])
        for i in range(len(pred fns)):
                title = pred fns[i]["name"]
                coef vals = pred fns[i]["coefs"]
                axs[i].bar(range(num_ftrs), coef_vals, color = "tab:purple")
                axs[i].set_xlabel('Feature Index')
                axs[i].set ylabel('Parameter Value')
                axs[i].set title(title)
        fig.subplots_adjust(hspace=0.4)
        return fig
```

### In [23]:

## compare parameter vectors(pred fns);



#### **Confusion Matrix**

We can try to predict the features with corresponding weight zero. We will fix a threshold eps such that any value between -eps and eps will get counted as zero. We take the remaining features to have positive value. These predictions of can be compared to the weights for the target function.

### In [24]:

```
def plot confusion matrix(cm, title, classes):
         plt.imshow(cm, interpolation='nearest', cmap=plt.cm.Purples)
         plt.title(title)
         plt.colorbar()
         tick marks = np.arange(len(classes))
         plt.xticks(tick marks, classes, rotation=45)
         plt.yticks(tick marks, classes)
         thresh = cm.max() / 2.
         for i, j in itertools.product(range(cm.shape[0]), range(cm.shape[1])):
                 plt.text(j, i, format(cm[i, j], 'd'),
                                  horizontalalignment="center",
                                  color="white" if cm[i, j] > thresh else "blac
k")
         plt.tight_layout()
         plt.ylabel('True label')
         plt.xlabel('Predicted label')
```

# Q1

### In [62]:

```
lambdas = [1e-5,1e-4,1e-3,1e-2,1e-1,1,10]
losses = []
from tqdm import tqdm

for lambd in lambdas:
    model = RidgeRegression(lambd)
    model.fit(X_train,y_train)
    losses.append(model.score(X_val,y_val))
best_param = lambdas[np.argmin(np.array(losses))]
```

### In [63]:

```
result = pd.DataFrame({"lambda":lambdas, "avg loss on val set":losses})
result
```

### Out[63]:

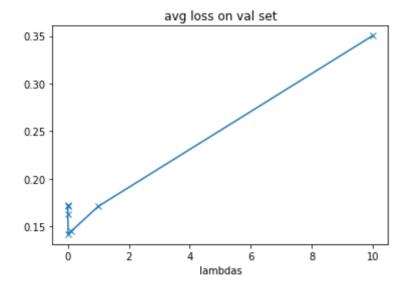
	lambda	avg loss on val set
0	0.00001	0.172464
1	0.00010	0.171345
2	0.00100	0.162705
3	0.01000	0.141887
4	0.10000	0.144566
5	1.00000	0.171068
6	10.00000	0.350321

## In [64]:

```
plt.plot(lambdas,losses,"x-")
plt.xlabel("lambdas")
plt.title("avg loss on val set")
```

### Out[64]:

Text(0.5, 1.0, 'avg loss on val set')



## In [65]:

```
print("Best parameter is: ", best_param)
```

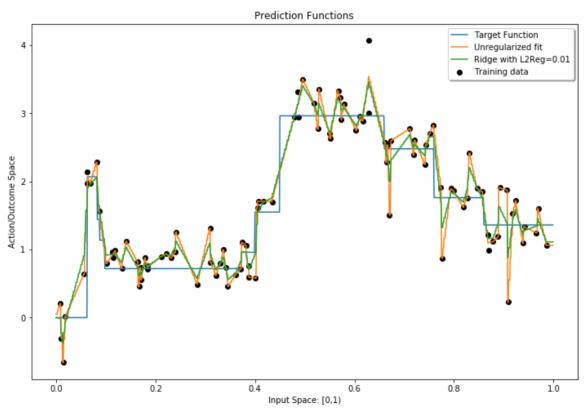
Best parameter is: 0.01

# **Q2**

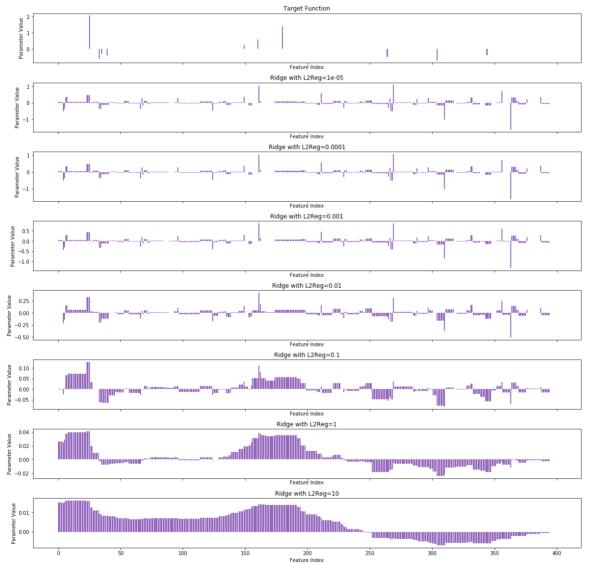
Plot the training data, the target function, an unregularized least squares fit (still using the featurizeddata), and the prediction function chosen in the previous proble

#### In [66]:

```
pred fns = []
x = np.sort(np.concatenate([np.arange(0,1,.001), x train]))
pred fns.append({"name": "Target Function", "coefs": coefs true, "preds": target
_fn(x)})
12regs = [0, grid.best params ['12reg'], 1]
X = featurize(x)
lambdas = [0, best_param]
for l2reg in lambdas:
    ridge regression estimator = RidgeRegression(12reg=12reg)
    ridge regression estimator.fit(X train, y train)
    if 12reg != 0:
        name = "Ridge with L2Reg="+str(12reg)
    else:
        name = "Unregularized fit"
    pred fns.append({"name":name,
                     "coefs":ridge regression estimator.w ,
                     "preds": ridge regression estimator.predict(X) })
plot_prediction_functions(x, pred_fns, x_train, y_train, legend_loc="best");
```



```
In [70]:
```



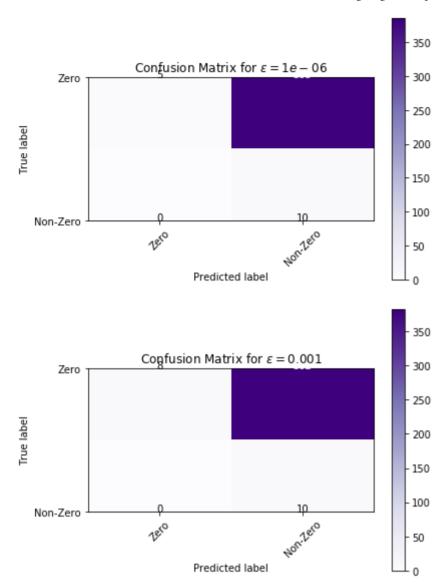
As we can see, as the regularization gets larger and larger, the coefficients will become smaller and smaller in absolute value. This confirms with how regularization works. Also, as regularization becomes smaller and closer to 0, the coefficients approach to those of the target function more and more.

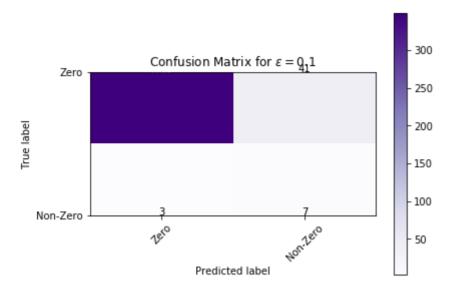
#### In [84]:

```
bin_coefs_true = coefs_true != 0 # your code goes here
eps_list = [1e-6,1e-3,1e-1]
ridge_regression_estimator = RidgeRegression(l2reg=best_param)
ridge_regression_estimator.fit(X_train, y_train)
w_tilde = ridge_regression_estimator.w_

for eps in eps_list:
    bin_coefs_estimated = np.abs(w_tilde) > eps # your code goes here
    cnf_matrix = confusion_matrix(bin_coefs_true, bin_coefs_estimated)
    plt.figure()
    plot_confusion_matrix(cnf_matrix, title="Confusion Matrix for $\end{a}\end{b}\end{b}\text{epsilon} = {\}
$".format(eps), classes=["Zero", "Non-Zero"])
```

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