Untitled

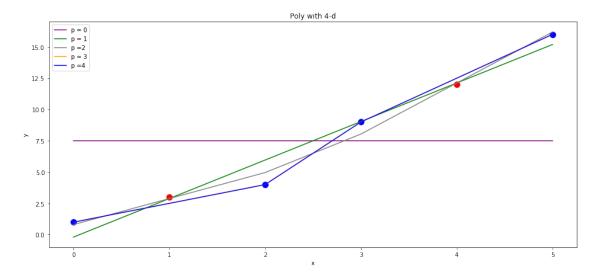
September 23, 2019

```
[24]: # QUESTION 1
   ______
   import numpy as np
   import matplotlib.pyplot as plt
   from sklearn.preprocessing import PolynomialFeatures
   from sklearn.linear_model import LinearRegression
   # Present the training and testing data points
   x_train = np.array([ [0],[2],[3],[5] ])
   y_{train} = [1,4,9,16]
   x_test = np.array([ [1],[4] ])
   y_{test} = [3,12]
   area = np.pi*30
   plt.figure(figsize=(16,7))
   plt.scatter(x_train, y_train, s=area, c='blue', alpha=1)
   plt.scatter(x_test, y_test, s=area, c='red', alpha=1)
   plt.title('Poly with 4-d')
   plt.xlabel('x')
   plt.ylabel('y')
   # LinearRegression
   # lin = LinearRegression()
   # lin.fit(x_train,y_train)
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# plt.plot(x_train, lin.predict(x_train), color = 'red')
\# D = 0
from sklearn.preprocessing import PolynomialFeatures
poly = PolynomialFeatures(degree = 0)
X poly = poly.fit transform(x train)
poly.fit(X_poly, y_train)
lin0 = LinearRegression()
lin0.fit(X_poly,y_train)
plt.plot(x_train, lin0.predict(poly.fit_transform(x_train)), color = 'purple',__
\rightarrowlabel = 'p = 0')
train_result0 = lin0.predict(poly.fit_transform(x_train))
test_result0 = lin0.predict(poly.fit_transform(x_test))
\# D = 1
from sklearn.preprocessing import PolynomialFeatures
poly = PolynomialFeatures(degree = 1)
X_poly = poly.fit_transform(x_train)
poly.fit(X_poly, y_train)
lin1 = LinearRegression()
lin1.fit(X_poly,y_train)
plt.plot(x_train, lin1.predict(poly.fit_transform(x_train)), color = 'green',__
\rightarrowlabel = 'p = 1')
train_result1 = lin1.predict(poly.fit_transform(x_train))
test_result1 = lin1.predict(poly.fit_transform(x_test))
\# D = 2
from sklearn.preprocessing import PolynomialFeatures
poly = PolynomialFeatures(degree = 2)
X_poly = poly.fit_transform(x_train)
poly.fit(X_poly, y_train)
lin2 = LinearRegression()
lin2.fit(X_poly,y_train)
plt.plot(x_train, lin2.predict(poly.fit_transform(x_train)), color = 'grey',__
\rightarrowlabel = 'p =2')
```

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train_result2 = lin2.predict(poly.fit_transform(x_train))
test_result2 = lin2.predict(poly.fit_transform(x_test))
\# D = 3
from sklearn.preprocessing import PolynomialFeatures
poly = PolynomialFeatures(degree = 3)
X_poly = poly.fit_transform(x_train)
poly.fit(X_poly, y_train)
lin3 = LinearRegression()
lin3.fit(X_poly,y_train)
plt.plot(x_train, lin3.predict(poly.fit_transform(x_train)), color = 'orange',_
\Rightarrowlabel = 'p = 3')
train_result3 = lin3.predict(poly.fit_transform(x_train))
test_result3 = lin3.predict(poly.fit_transform(x_test))
#D = 4
from sklearn.preprocessing import PolynomialFeatures
poly = PolynomialFeatures(degree = 4)
X_poly = poly.fit_transform(x_train)
poly.fit(X_poly, y_train)
lin4 = LinearRegression()
lin4.fit(X_poly,y_train)
plt.plot(x_train, lin4.predict(poly.fit_transform(x_train)), color = 'blue',_
 \rightarrowlabel = 'p =4')
train_result4 = lin4.predict(poly.fit_transform(x_train))
test_result4 = lin4.predict(poly.fit_transform(x_test))
\# D = 8
from sklearn.preprocessing import PolynomialFeatures
poly = PolynomialFeatures(degree = 8)
X_poly = poly.fit_transform(x_train)
poly.fit(X_poly, y_train)
lin20 = LinearRegression()
lin20.fit(X_poly,y_train)
test_result20 = lin20.predict(poly.fit_transform(x_test))
```

```
plt.legend()
plt.show()
```



```
[26]: # Training Result
    train result = ['','','','']
    train_result = np.row_stack((train_result,train_result0))
    train_result = np.row_stack((train_result,train_result1))
    train_result = np.row_stack((train_result, train_result2))
    train_result = np.row_stack((train_result, train_result3))
    train_result = np.row_stack((train_result, train_result4))
    train result = np.delete(train result, (0), axis=0)
    print("----- train_result -----")
    print(train_result)
    # Test Result
    \# x_train_matrix = [0,0,0]
    \# x_train_matrix = np.row_stack((x_train_matrix, x_1))
    # x_train_matrix = np.row_stack((x_train_matrix,x_2))
    \# x_train_matrix = np.row_stack((x_train_matrix, x_3))
    # x_train_matrix = np.row_stack((x_train_matrix,x_4))
    # x_train_matrix = np.delete(x_train_matrix, (0), axis=0)
```

```
test_result = ['','']
test_result = np.row_stack((test_result,test_result0))
test_result = np.row_stack((test_result, test_result1))
test_result = np.row_stack((test_result, test_result2))
test_result = np.row_stack((test_result, test_result3))
test_result = np.row_stack((test_result,test_result4))
test_result = np.delete(test_result, (0), axis=0)
print("----- test result -----")
print(test_result)
# Typical Bias
x_{test} = np.array([[1],[4]])
y_{test} = [3, 12]
Typical_Bias_0 = pow( (float(test_result[0][0]) - float(x_test[0])),2 ) + pow(__
\rightarrow(float(test_result[0][1]) - float(x_test[1])), 2)/2
Typical Bias 1 = pow((float(test result[1][0]) - float(x test[0])), 2) + pow(1)
\rightarrow (float(test_result[1][1]) - float(x_test[1])), 2)/2
Typical_Bias_2 = pow( (float(test_result[2][0]) - float(x_test[0])),2 ) + pow(__
\rightarrow (float(test_result[2][1]) - float(x_test[1])), 2)/2
Typical_Bias_3 = pow( (float(test_result[3][0]) - float(x_test[0])),2 ) + pow(__
\rightarrow (float(test_result[3][1]) - float(x_test[1])), 2)/2
Typical_Bias_4 = pow( (float(test_result[4][0]) - float(x_test[0])),2 ) + pow(__
 →(float(test_result[4][1]) - float(x_test[1])), 2)/2
Typical_Bias = []
Typical Bias.append(Typical Bias 0)
Typical_Bias.append(Typical_Bias_1)
Typical_Bias.append(Typical_Bias_2)
Typical_Bias.append(Typical_Bias_3)
Typical_Bias.append(Typical_Bias_4)
print(Typical_Bias)
# Variance
Variance = []
k = 0
print(len(Typical_Bias))
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for i in range(len(Typical_Bias)):
   Variance.append( Typical_Bias[k]/2 )
   k += 1
print(Variance)
# Total Error
Total_Error = []
for i in range(len(test_result)):
   Total_Error.append(float(test_result[i][0]) - float(x_test[0]) + _
→float(test_result[i][1]) - float(x_test[1]) )
print(Total_Error)
# Training Error
x_{train} = np.array([[0],[2],[3],[5]])
y_{train} = [1,4,9,16]
training_error = []
for i in train_result:# Get every training result
   k = 0
   training_error_sum = 0
   for j in i:
      training_error_sum = training_error_sum + pow( y_train[int(k)] -__
 \rightarrowfloat(i[k]), 2)
      k += 1
   training_error.append(training_error_sum/4)
print("----- Training Error -----")
print(training_error)
# Test Error
testing_error = []
for i in test_result:# Get every training result
   k = 0
   testing_error_sum = 0
```

```
for j in i:
       testing_error_sum = testing_error_sum + pow( y_test[int(k)] -___
 \rightarrowfloat(i[k]), 2)
       k += 1
   testing error.append(testing error sum/4)
print("----- Testing Error -----")
print(testing_error)
# Plot
x_{point} = [0,1,2,3,4]
plt.figure(figsize=(14,7))
plt.plot(x_point, Typical_Bias , color = 'yellow', label = 'Typical_Bias')
plt.plot(x point, Variance , color = 'blue', label = 'Variance')
plt.plot(x_point, Total_Error , color = 'pink', label = 'Total_Error')
plt.plot(x_point, training_error , color = 'black', label = 'Training_error')
plt.plot(x_point, testing_error , color = 'grey', label = 'Testing_error')
plt.title('Error')
plt.xlabel('x')
plt.ylabel('y')
plt.legend()
plt.show()
# (3) Explain why each of the five curves has the shape displayed in part (2).
# --> Here we can see that when d=2 the error come to the minumum. Why? Because
→1) when d is 0 or 1, it is not enough to fit the data probably.2) While when
\rightarrow d > 2, the curve is overfited - which means the curve is fluctuate too muc -
which away from the ground truth more and more when d become large
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns # data visualization library
from scipy.optimize import minimize
from sklearn.datasets import load_boston
from sklearn.model_selection import KFold
```

```
boston_dataset = load_boston()
df = pd.DataFrame(boston_dataset.data, columns=boston_dataset.feature_names)
print (df.head(5))
#the data can be downloaded from "https://github.com/jcrouser/islr-python/blob/
→master/data/Smarket.csv"
df = pd.read_csv('Smarket.csv', usecols=range(0,10), index_col=0,__
→parse_dates=True)
X = df[['Lag1','Lag2']].values
Y = df['Today'].values
print(X[:,1])
print(len(X))
print(len(Y))
def forward(X_1,X_2,params):
   \# 3d \ Line \ Z = aX_1 + bX_2 + c
   return X_1.dot(params[:1]) + X_2.dot(params[1:2]) + params[2]
def cost_function(params, X_1, X_2, y, p):
   error_vector = y - forward(X_1, X_2, params)
   return np.linalg.norm(error vector, ord=p)
kf = KFold(n_splits=5)
kf.get_n_splits(X)
#print(kf)
\#KFold(n\_splits=5, random\_state=None, shuffle=False)
#print("----")
#print(type(kf.split(X)))
L2_Norm_P1 = 0
L2_Norm_P2 = 0
for train_index, test_index in kf.split(X):
  # print("TRAIN:", train_index, "TEST:", test_index)
   ### For X part
   X_train, X_test = X[train_index], X[test_index]
```

```
### For Y part
   Y_train, Y_test = Y[train_index], Y[test_index]
   ############################ P = 1
   output1 = minimize( cost_function, [0.5,0.5,1], args=( np.c_[X_train[:
→,0]], np.c_[X_train[:,1]],Y_train,1
                                    ) )
   y_hat1 = forward(np.c_[X_train[:,0]],np.c_[X_train[:,1]], output1.x) #_
 \rightarrow print (y_hat1)
   ### Validation
   L2_Norm_Sum = 0
   for i in range(len(Y_test)):
        L2 = np.power((y_hat1[i] - Y_test[i]),2)
        L2_Norm_Sum = L2_Norm_Sum + L2
   # Add all together
   L2_Norm_P1 = L2_Norm_P1 + np.sqrt(L2_Norm_Sum)
   ### Training
   output2 = minimize( cost_function, [0.5,0.5,1], args=( np.c_[X_train[:
 →,0]], np.c_[X_train[:,1]],Y_train,2 )
   y_hat2 = forward(np.c_[X_train[:,0]],np.c_[X_train[:,1]], output2.x) #_
 \rightarrow print (y_hat2)
   ### Validation
   L2_Norm_Sum = 0
   for i in range(len(Y_test)):
        L2 = np.power((y_hat2[i] - Y_test[i]),2)
        L2_Norm_Sum = L2_Norm_Sum + L2
   # Add all together
   L2_Norm_P2 = L2_Norm_P2 + np.sqrt(L2_Norm_Sum)
L2_Norm_P1_Mean = L2_Norm_P1/5
L2_Norm_P2_Mean = L2_Norm_P2/5
print(L2_Norm_P1_Mean)
print(L2_Norm_P2_Mean)
```

# Please compare the mean values across all the 5 folds together. Justify your \rightarrow results.	
#> I believe the reason is that, after cross validation, the error can be \Box	
\hookrightarrow minimized. Thus even we trained with different P value, the result is similar	
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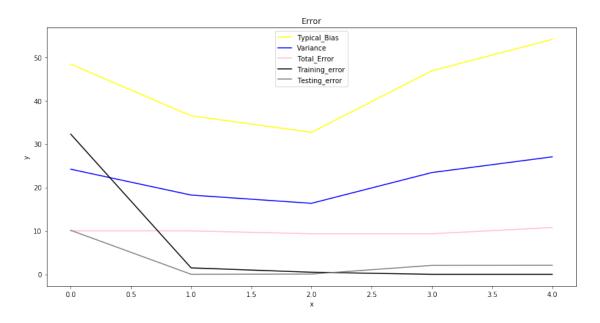
```
# (2) Logistic Regression Regularization Comparison with Bootstrapping:
#Using 80% of the data as a training set and 20% as a testing set in each
 →bootstrap repeated 1000 times each, please implement and compare the average
 →and the standard deviation of coefficients obtained from Ridge regression
 →and LASSO regularized logistic regression.
#By averaging the coefficients it is meant that all the different coefficient \sqcup
 →values for a specific variable is averaged. Coefficient average values then
 → are to be compared by plotting them against each other. Are all input,
 → features necessary? Please describe how categorical features are handled.
# (3) Please plot the ROC curve for both models for a single bootstrap data. \Box
 →What are the area under the curve measurements?
# (4) What is the optimal decision threshold to maximize the f1 score?
# (5) Please provide a mean and standard deviation for the AUROCs for each
 →model . (6) Please provide a mean and standard deviation for the f1 score
 \rightarrow for each model.
# -----
from sklearn.linear_model import LogisticRegression, Ridge, Lasso
from sklearn.model_selection import GridSearchCV
from sklearn.model selection import StratifiedKFold
from sklearn.metrics import roc_auc_score
----- train result -----
[['7.5' '7.5' '7.5' '7.5']
 ['-0.19230769230769518' '5.961538461538461' '9.038461538461538'
 '15.192307692307695']
 ['0.8076923076923119' '4.961538461538469' '8.038461538461542'
 '16.19230769230768']
 ['1.0' '3.9999999999992' '9.00000000000007' '16.00000000000004']
['1.000000000000226' '4.0000000000016' '9.0000000000007'
 '15.999999999995']]
----- test_result -----
[['7.5' '7.5']
 ['2.884615384615383' '12.115384615384617']
['2.551282051282058' '11.78205128205128']
```

```
['0.666666666666536' '13.666666666668']
['1.3849813115868361' '14.384981311586802']]
[48.375, 36.48150887573965, 32.686637080867854, 46.83333333333348,
54.07212903127469]
5
[24.1875, 18.240754437869825, 16.343318540433927, 23.41666666666674,
27.036064515637346]
[10.0, 10.0, 9.333333333333337, 9.3333333333334, 10.769962623173639]
------ Training Error -----
```

[32.25, 1.4807692307692308, 0.48076923076923095, 3.1751651435145725e-29, 8.232256584047021e-28]

----- Testing Error -----

[10.125, 0.006656804733727994, 0.06221236028928212, 2.05555555555555552, 2.07410530513552]



	CRIM	ZN	INDUS	CHAS	NOX	RM	AGE	DIS	RAD	TAX	\
0	0.00632	18.0	2.31	0.0	0.538	6.575	65.2	4.0900	1.0	296.0	
1	0.02731	0.0	7.07	0.0	0.469	6.421	78.9	4.9671	2.0	242.0	
2	0.02729	0.0	7.07	0.0	0.469	7.185	61.1	4.9671	2.0	242.0	
3	0.03237	0.0	2.18	0.0	0.458	6.998	45.8	6.0622	3.0	222.0	
4	0.06905	0.0	2.18	0.0	0.458	7.147	54.2	6.0622	3.0	222.0	
	PTRATIO	Е	B LSTAT	Γ							
0	15.3	396.90	4.98	3							
1	17.8	396.90	9.14	4							
2	17.8	392.83	4.03	3							

3 18.7 394.63 2.94 4 18.7 396.90 5.33

```
[-0.192 0.381 0.959 ... 0.043 -0.955 0.13]

1250

1250

16.972221525679736

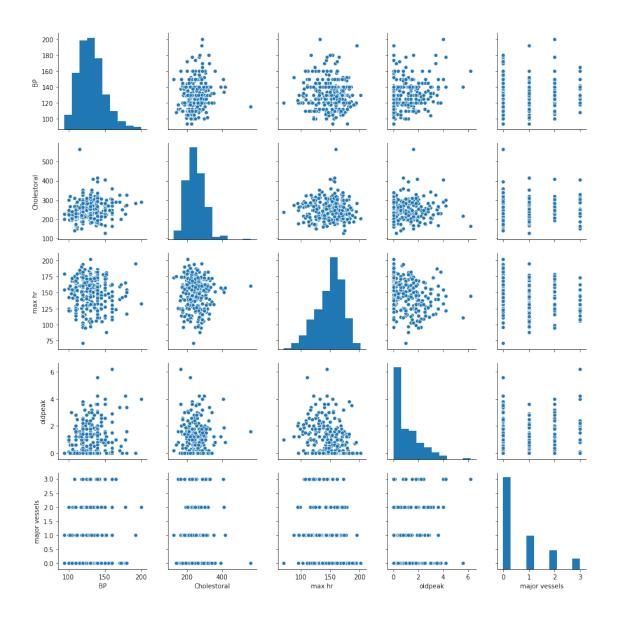
16.926771309143838

[27]: import numpy as np
import pandas as pd
import seaborn as sns

df = pd.read_csv('hw1_input.csv', usecols=range(0,14), index_col=0,___
--parse_dates=True)

[18]: sns.pairplot(df)
# this is to give relationship between different variables. For example, here__
--we have "BP" as the x-axis and Cholestoral as the y-axis, to plot an image
```

[18]: <seaborn.axisgrid.PairGrid at 0x1a25d656d8>



[9]: df.des	cribe()			
[9]:	ВР	Cholestoral	max hr	oldpeak
count	270.000000	270.000000	270.000000	270.00000
mean	131.344444	249.659259	149.677778	1.05000
std	17.861608	51.686237	23.165717	1.14521
min	94.000000	126.000000	71.000000	0.00000
25%	120.000000	213.000000	133.000000	0.00000
50%	130.000000	245.000000	153.500000	0.80000
75%	140.000000	280.000000	166.000000	1.60000
max	200.000000	564.000000	202.000000	6.20000

```
[13]:
                             BP Cholestoral fasting blood sugar > 120 \
            Sex Chest Pain
    Age
    70
         Female
                  Abnormal 130
                                         322
                                                                     No
        resting ECG max hr angina oldpeak slope major vessels defect \
    Age
     70
              hyper
                         109
                                No
                                         2.4 Flat
                                                                3 Normal
        heart disease
     Age
     70
                  Yes
[20]: from sklearn.linear_model import LogisticRegression, Ridge, Lasso
     from sklearn.model_selection import GridSearchCV
     from sklearn.model_selection import StratifiedKFold
     from sklearn.metrics import roc_auc_score
 []:
```