

BILKENT UNIVERSITY

CS464 HOMEWORK 2

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Question 1.1

$$\min \frac{1}{2} \|y - XB\|^2 = \min tr(Y'Y) - 2tr(Y'XB) + tr(B'X'XB)$$
$$(\partial/\partial B) = 0 \Rightarrow \widehat{B} = (X'X)^{-1}X'Y$$

Question 1.2

There are rank(X'X) = 6 independent features in the model.

Question 1.3

Coefficients: [-7.68973700e-01 -2.66894243e-02 -3.62572818e-02 3.28973372e-05

-3.03106645e-01 5.28691557e-01] Train Error: 10.20164702644783 Test Error: 35.975047174669974

Question 1.4

If a coefficient is negative, an increase in corresponding feature value will result in a decrease in the prediction. Likewise if a coefficient is positive, an increase in corresponding feature value will result in an increase in the prediction.

As the magnitude of a coefficient increases, the effect of corresponding feature on the prediction will increase, making it a stronger factor in determining the prediction.

Question 1.5

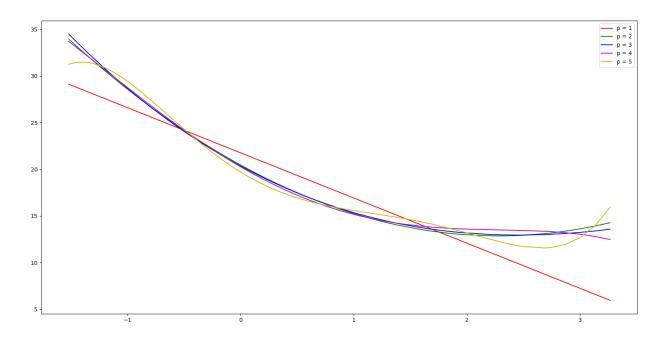
MPG is inversely proportional to horsepower(HP) value in this dataset, which explains the negative coefficient of beta values in Question 1.3.

Question 1.6

Matrix rank increases until p = 2, then stays constant at rank = 3: rank = p + 1, if p <= 2; else, 3. This means powers of HP greater than 2 can be written as a linear combination of lesser powers, and they powers cannot be used as features directly. After normalization, we observed that rank of X'X increases as we increase the parameter p, which implies centralization created linearly independent features that can be used to predict class label of train data. Then, rank function becomes: rank = p, p <= 6. However if X'X is not invertible, we cannot use the closed solution of beta directly. For example, when p = 0, we have a singular matrix.

Question 1.7

Since p = 0 gives a singular matrix for closed form of beta, we ignore it.



Results:

Coefficients of Beta for p = 1:

[21.75876359 -4.83692608]

Train Error for p = 1: 14.341899019749064

Test Error for p = 1: 73.83848307979181

Coefficients of Beta for p = 2:

[20.45431495 -6.65954213 1.46039737]

Train Error for p = 2: 10.839424149052432

Test Error for p = 2: 60.53249695797431

Coefficients of Beta for p = 3:

[20.34507117 -6.56475888 1.66498393 -0.08860305]

Train Error for p = 3: 10.816044462865847

Test Error for p = 3: 60.31291330318858

Coefficients of Beta for p = 4:

[20.26224255 -7.02669019 1.80373431 0.23874763 -0.10890227]

Train Error for p = 4: 10.771087052589538

Test Error for p = 4: 59.991301591670435

Coefficients of Beta for p = 5:

[19.70530656 -7.42906053 3.86834442 0.26457814 -1.07547209 0.24861662]

Train Error for p = 5: 10.383020895493294

Test Error for p = 5: 58.329788653047466

Question 1.8

Coefficients of Beta for p = 1:

[22.0255366 -4.69684017 0.71822189]

Train Error for p = 1: 14.066764652607663

Test Error for p = 1: 59.22618224681992

Coefficients of Beta for p = 2:

[20.21911804 -6.62573072 1.53009732 2.44046804 1.54598368]

Train Error for p = 2: 9.45156444953559

Test Error for p = 2: 22.293142924636175

Coefficients of Beta for p = 3:

[20.08490786 -6.5159226 1.76776157 -0.10331268 2.44994773 1.56194401]

Train Error for p = 3: 9.419849048918984

Test Error for p = 3: 21.94819248761814

We observe that since underlying distribution is not linear, using polynomials of a feature set gives better results than simple linear regression. Moreover, using many features (ignoring curse of dimensionality) results in lower test error for almost all cases. It is also clear that after p = 2, for 1.7, using higher powers do not result in significant performance improvement which means p = 2 is the most cost-effective predictor for Question 1.7 Similarly, we see that p = 2 is the most cost-effective predictor for Question 1.8 as well. Finally, observe that polynomials of multiple features is the best predictor we trained, which is suitable with the assumption that more features usually means better outcomes.

Question 2.1

Hyper parameters: Iteration count -> 2500, Learning Rate -> 0.001

Confusion Matrix -> {'tp': 20, 'tn': 20, 'fp': 0, 'fn': 0}

Question 2.2

Current status of Question 2.2 includes forward selection and backward elimination implementations, however, because of the long execution time we are unable to acquire the final output (backward elimination only completed ~70 eliminations in approximately 4 hours, and forward selection completed ~1000 features). However, the program we provide correctly selects features except the first feature, which is a by-product of initialization step of the algorithms. Since we do not have the final output, we are unable to provide the confusion matrix, however, assuming execution is completed, the program will indeed print the confusion matrices for each feature set.

As a final note to their executions, we observed that intermediate outputs of two algorithms are different. This is so since backward elimination seeks features that, when removed from feature set, will decrease classification accuracy whereas forward selection seeks features that, when

inserted into a subset of the feature set, will improve the classification accuracy. From here, we see the distinction between the two as selecting non-optimal features that increase intermediate prediction accuracy vs. removing non-optimal features that do not reduce intermediate prediction accuracy. Hence, the execution of the algorithms above and their outputs are naturally different. To address this issue, we might run forward selection algorithm multiple times with random shuffling of the feature set in order to eliminate the effect of locations of features on the output.

Question 3.1

First option is line x1 + x2 = 3, with margin w = 1 / sqrt(2). In this case, optimization function is $||w||^2 + C * (number of errors) = ||w||^2$, since no error exists.

Another option is x1 + x2 = 5, with margin = sqrt(2). In this case, optimization function is $||w||^2 + C * 1$, since data point (2,2) violates the margin.

Last option is x1 + x2 = 7, with margin = 2sqrt(2). In this case, optimization function is $||w||^2 + C * 2$, since orange points violate the margin.

So, for any C value greater than 0, optimizer will select the hard margin SVM.

Question 3.2

If C is set to infinity, all constraints are enforced and the margin becomes a hard one, since cost of a margin violation becomes infinity.

Question 3.3

```
C -> 10.0
```

```
Confusion Matrix -> {'tp': 42, 'tn': 141, 'fp': 0, 'fn': 0}
Accuracy: 1.0
```

We used k-fold validation with iteration count 5, for statistically significant results, and small number of iterations.

Question 3.4

Accuracy: 0.9836065573770492

```
Linear SVM with C = 10.0

Confusion Matrix -> {'tp': 42, 'tn': 141, 'fp': 0, 'fn': 0}

Accuracy: 1.0

SVM with RBF kernel, gamma = 0.0625

Confusion Matrix -> {'tp': 42, 'tn': 138, 'fp': 3, 'fn': 0}
```

We used k-fold validation with iteration count 5, for statistically significant results, and small number of iterations.

Source Code

Ouestion 1

```
# Cylinders, Displacement, Horsepower, Weight, Acceleration, Model Year and
MPG
import numpy as np
from numpy.linalg import inv, matrix rank
import matplotlib.pyplot as plt
data = np.loadtxt("../data/carbig.csv", delimiter=',')
print("Question 1.2: ")
(x, y) = data.shape
train sample count = 300
train data = data[:train sample count, : y - 1]
train labels = data[:train sample count, y - 1]
test data = data[train sample count : x + 1, : y - 1]
test labels = data[train sample count: x + 1, y - 1]
# print(matrix rank(np.transpose(train data).dot(train data)))
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
train predictions = train data.dot(beta)
train error = (np.square(train labels - train predictions)).mean(axis = None)
test predictions = test data.dot(beta)
test error = (np.square(test labels - test predictions)).mean(axis = None)
print("Coefficients of Beta: ")
print(beta)
print("Train Error: " + str(train error))
print("Test Error: " + str(test error))
print("************************")
print("Question 1.5: ")
(x, y) = data.shape
plt.plot(data[:, 2], data[:, y - 1], ".")
plt.xlabel("Horsepower")
plt.ylabel("MPG")
plt.show()
print("*************************")
print("Question 1.6: ")
features = np.column_stack((np.ones(x), data[:, 2], data[:, 2] ** 2, data[:, 2])
2] ** 3, data[:, 2] ** 4, data[:, 2] ** 5))
f1 = features[:, 0]
f2 = features[:, 0:2]
f3 = features[:, 0:3]
```

```
f4 = features[:, 0:4]
f5 = features[:, 0:5]
f6 = features[:, 0:6]
xTx1 = np.transpose(f1).dot(f1)
xTx2 = np.transpose(f2).dot(f2)
xTx3 = np.transpose(f3).dot(f3)
xTx4 = np.transpose(f4).dot(f4)
xTx5 = np.transpose(f5).dot(f5)
xTx6 = np.transpose(f6).dot(f6)
rank1 = matrix rank(xTx1)
rank2 = matrix rank(xTx2)
rank3 = matrix rank(xTx3)
rank4 = matrix rank(xTx4)
rank5 = matrix rank(xTx5)
rank6 = matrix rank(xTx6)
print("Ranks before normalization: ")
print("Rank for p = 0: " + str(rank1))
print("Rank for p = 1: " + str(rank2))
print("Rank for p = 2: " + str(rank3))
print("Rank for p = 3: " + str(rank4))
print("Rank for p = 4: " + str(rank5))
print("Rank for p = 5: " + str(rank6))
mean = features.mean(0)
stddev = features.std(0)
for i in range(x):
      for j in range (1, y - 1):
            features[i][j] = (lambda x, y, z, i, j: float(x - y[j]) / z[j])
(features[i][j], mean, stddev, i, j)
f1 = features[:, 0]
f2 = features[:, 0:2]
f3 = features[:, 0:3]
f4 = features[:, 0:4]
f5 = features[:, 0:5]
f6 = features[:, 0:6]
xTx1 = np.transpose(f1).dot(f1)
xTx2 = np.transpose(f2).dot(f2)
xTx3 = np.transpose(f3).dot(f3)
xTx4 = np.transpose(f4).dot(f4)
xTx5 = np.transpose(f5).dot(f5)
xTx6 = np.transpose(f6).dot(f6)
rank1 = matrix rank(xTx1)
rank2 = matrix rank(xTx2)
rank3 = matrix rank(xTx3)
rank4 = matrix rank(xTx4)
rank5 = matrix rank(xTx5)
rank6 = matrix rank(xTx6)
print("Ranks before normalization: ")
```

```
print("Rank for p = 0: " + str(rank1))
print("Rank for p = 1: " + str(rank2))
print("Rank for p = 2: " + str(rank3))
print("Rank for p = 3: " + str(rank4))
print("Rank for p = 4:" + str(rank5))
print("Rank for p = 5: " + str(rank6))
print("************************")
print("Question 1.7: ")
hp = data[:, 2]
mean hp = hp.mean(0)
stddev hp = hp.std(0)
for i in range(x):
    hp[i] = (lambda x: (x - mean hp) / stddev hp) (hp[i])
features_17_1 = np.column_stack((np.ones(x), hp))
features 17 2 = np.column stack((np.ones(x), hp, hp ** 2))
features 17 3 = np.column stack((np.ones(x), hp, hp ** 2, hp ** 3))
features 17 4 = np.column stack((np.ones(x), hp, hp ** 2, hp ** 3, hp ** 4))
features 17 5 = np.column stack((np.ones(x), hp, hp ** 2, hp ** 3, hp ** 4, hp
** 5))
\# p = 1
train sample count = 300
train data = features 17 1[:train sample count, : y - 1]
train labels = data[:train sample count, y - 1]
test_data = features_17_1[train_sample_count : x + 1, : y - 1]
test labels = data[train sample count: x + 1, y - 1]
beta 1
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
train predictions = train data.dot(beta 1)
train error = (np.square(train labels - train predictions)).mean(axis = None)
test predictions = test data.dot(beta 1)
test error = (np.square(test labels - test predictions)).mean(axis = None)
print("Coefficients of Beta for p = 1: ")
print(beta 1)
print("Train Error for p = 1: " + str(train error))
print("Test Error for p = 1: " + str(test error))
# p = 2
train sample count = 300
train data = features 17 2[:train sample count, : y - 1]
train labels = data[:train sample count, y - 1]
test data = features 17 2[train sample count : x + 1, : y - 1]
test labels = data[train sample count: x + 1, y - 1]
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
```

```
train predictions = train data.dot(beta 2)
train error = (np.square(train labels - train predictions)).mean(axis = None)
test predictions = test data.dot(beta 2)
test error = (np.square(test labels - test predictions)).mean(axis = None)
print("Coefficients of Beta for p = 2: ")
print(beta 2)
print("Train Error for p = 2: " + str(train error))
print("Test Error for p = 2: " + str(test error))
# p = 3
train sample count = 300
train data = features 17 3[:train sample count, : y - 1]
train labels = data[:train sample count, y - 1]
test data = features 17 3[train sample count : x + 1, : y - 1]
test labels = data[train sample count: x + 1, y - 1]
beta 3
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
train predictions = train data.dot(beta 3)
train error = (np.square(train labels - train predictions)).mean(axis = None)
test predictions = test data.dot(beta 3)
test_error = (np.square(test_labels - test_predictions)).mean(axis = None)
print("Coefficients of Beta for p = 3: ")
print(beta 3)
print("Train Error for p = 3: " + str(train error))
print("Test Error for p = 3: " + str(test error))
# p = 4
train sample count = 300
train data = features 17 4[:train sample count, : y - 1]
train labels = data[:train sample count, y - 1]
test data = features 17 4[train sample count : x + 1, : y - 1]
test labels = data[train sample count: x + 1, y - 1]
beta 4
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
train predictions = train data.dot(beta 4)
train error = (np.square(train labels - train predictions)).mean(axis = None)
test predictions = test data.dot(beta 4)
test error = (np.square(test labels - test predictions)).mean(axis = None)
print("Coefficients of Beta for p = 4: ")
print(beta 4)
print("Train Error for p = 4: " + str(train error))
print("Test Error for p = 4: " + str(test error))
# p = 5
train sample count = 300
train data = features 17 5[:train sample count, : y - 1]
train labels = data[:train sample count, y - 1]
```

```
test data = features 17 5[train sample count : x + 1, : y - 1]
test labels = data[train sample count: x + 1, y - 1]
beta 5
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
train predictions = train data.dot(beta 5)
train error = (np.square(train labels - train predictions)).mean(axis = None)
test predictions = test data.dot(beta 5)
test error = (np.square(test labels - test predictions)).mean(axis = None)
print("Coefficients of Beta for p = 5: ")
print(beta 5)
print("Train Error for p = 5: " + str(train error))
print("Test Error for p = 5: " + str(test error))
shp = np.sort(hp)
plt.plot(shp, (lambda x: beta 1[0] + beta 1[1] * x)(shp), "r", label = "p = "p"
plt.plot(shp, (lambda x: beta 2[0] + beta 2[1] * x + beta 2[2] * (x ** 2))
(shp), "g", label = "p = 2")
plt.plot(shp, (lambda x: beta 3[0] + beta 3[1] * x + beta 3[2] * (x ** 2) +
beta 3[3] * (x ** 3)) (shp), "b", label = "p = 3")
plt.plot(shp, (lambda x: beta_4[0] + beta_4[1] * x + beta_4[2] * (x ** 2) +
beta 4[3] * (x ** 3) + beta 4[4] * (x ** 4))(shp), "m", label = "p = 4")
plt.plot(shp, (lambda x: beta 5[0] + beta 5[1] * x + beta 5[2] * (x ** 2) +
beta 5[3] * (x ** 3) + beta_5[4] * (x ** 4) + beta_5[5] * (x ** 5))(shp), "y",
label = "p = 5")
plt.legend()
plt.show()
print("*********************************
print("Question 1.8: ")
my = data[:, 5]
mean my = my.mean(0)
stddev my = my.std(0)
for i in range(x):
    my[i] = (lambda x: (x - mean my) / stddev my) (my[i])
features 18 1 = np.column stack((np.ones(x), hp, my))
features 18 2 = np.column stack((np.ones(x), hp, hp ** 2, my, my ** 2))
features_18_3 = np.column_stack((np.ones(x), hp, hp ** 2, hp ** 3, my, my **
2, my ** 3))
\# p = 1
train sample count = 300
train data = features 18 1[:train sample count, : y - 1]
train labels = data[:train sample count, y - 1]
test data = features 18 1[train sample count : x + 1, : y - 1]
test labels = data[train sample count: x + 1, y - 1]
beta
```

```
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
train predictions = train data.dot(beta)
train_error = (np.square(train labels - train predictions)).mean(axis = None)
test predictions = test data.dot(beta)
test error = (np.square(test labels - test predictions)).mean(axis = None)
print("Coefficients of Beta for p = 1: ")
print(beta)
print("Train Error for p = 1: " + str(train error))
print("Test Error for p = 1: " + str(test error))
# p = 2
train sample count = 300
train data = features_18_2[:train_sample_count, : y - 1]
train labels = data[:train sample count, y - 1]
test data = features 18 2[train sample count : x + 1, : y - 1]
test labels = data[train_sample_count: x + 1, y - 1]
beta
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
train predictions = train data.dot(beta)
train_error = (np.square(train labels - train predictions)).mean(axis = None)
test predictions = test data.dot(beta)
test error = (np.square(test labels - test predictions)).mean(axis = None)
print("Coefficients of Beta for p = 2: ")
print(beta)
print("Train Error for p = 2: " + str(train error))
print("Test Error for p = 2: " + str(test error))
# p = 3
train sample count = 300
train data = features 18 3[:train sample count, : y - 1]
train labels = data[:train sample count, y - 1]
test data = features 18 3[train sample count : x + 1, : y - 1]
test labels = data[train sample count: x + 1, y - 1]
beta
inv(np.transpose(train data).dot(train data)).dot(np.transpose(train data)).do
t(train labels)
train predictions = train data.dot(beta)
train_error = (np.square(train_labels - train_predictions)).mean(axis = None)
test predictions = test data.dot(beta)
test error = (np.square(test labels - test predictions)).mean(axis = None)
print("Coefficients of Beta for p = 3: ")
print(beta)
print("Train Error for p = 3: " + str(train error))
print("Test Error for p = 3: " + str(test error))
```

Question 2

import numpy as np

```
import matplotlib.pyplot as plt
import sys
def predict(w0, weights, sample):
    tmp = 1 / (1 + np.exp(w0 + np.sum(np.multiply(weights, sample))))
    return 0 if tmp > 0.5 else 1
     gradientAscent(lr,
                            iteration count, feature count, sample count,
train data, train labels):
    w0 = 0
    weights = np.zeros((feature count))
    for it in range(iteration count):
             y minus prediction = np.asarray([train labels[i] - predict(w0,
weights, train data[i]) for i in range(sample count)])
        w0 = w0 + lr * np.sum(y minus prediction)
        tmp = train data * y minus prediction[:, np.newaxis]
        tmp = tmp.sum(axis = 0)
       weights = weights + lr * tmp
    return (w0, weights)
def forwardSelection(kf train data, kf train labels, ic, lr):
    indices = np.asarray([0])
    previous score = float(sys.maxsize)
    (t1, t2) = kf train data[0].shape
    for j in range(1, t2): # for each feature
       print("Feature at index " + str(j) + "...")
       new indices = np.append(indices, j)
       mse = np.zeros(len(kf train data))
        for l in range(len(kf train data)): # k-fold
            new data = kf train data[l][:, new indices]
            new test data = kf test data[l][:, new indices]
            (x train, y train) = new data.shape
            (w0, weights) = gradientAscent(lr, ic, y train, x train, new data,
kf train labels[l])
            predictions = np.vectorize(lambda x: 0 if x > 0.5 else 1)(1 / (1 +
np.exp(np.multiply(new test data, weights).sum(axis = 1) + w0)))
            mse[l] = np.sum(np.power(kf test labels[l] - predictions, 2))
        avg_mse = np.sum(mse) * float(1 / len(mse))
       print("avg mse: " + str(avg mse))
       print("previous score: " + str(previous score))
        if avg mse < previous score:
            print("Feature at index " + str(j) + " is accepted.")
            indices = new indices
            previous score = avg mse
        else:
            print("Feature at index " + str(j) + " is rejected.")
    return indices
```

```
def backwardElimination(kf train data, kf train labels, ic, lr):
    indices = np.arange(len(kf train data[0][0]))
   previous score = -sys.maxsize
    (t1, t2) = kf train data[0].shape
    for i in range(t2):
       print("Feature at index " + str(i) + "...")
        new indices = np.delete(indices, i)
       mse = np.zeros(len(kf train data))
        for l in range(len(kf train data)): # k-fold
            new data = kf train data[l][:, new indices]
            new test data = kf test data[l][:, new indices]
            (x train, y train) = new data.shape
            (w0, weights) = gradientAscent(lr, ic, y train, x train, new data,
kf train labels[l])
           predictions = np.vectorize(lambda x: 0 if x > 0.5 else 1)(1 / (1 +
np.exp(np.multiply(new test data, weights).sum(axis = 1) + w0)))
           mse[l] = np.sum(np.power(kf test labels[l] - predictions, 2))
        avg_mse = np.sum(mse) * float(1 / len(mse))
       print("avg mse: " + str(avg mse))
       print("previous_score: " + str(previous_score))
        if avg mse > previous score:
           print("Feature at index " + str(i) + " is accepted.")
           print("Feature at index " + str(i) + " is rejected.")
            indices = new indices
           previous score = avg mse
    return indices
data = np.loadtxt("../data/ovariancancer.csv", dtype = "float", delimiter=',')
labels = np.loadtxt("../data/ovariancancer labels.csv", dtype = "float",
delimiter=',')
test data = np.row stack((data[:20,:], data[121:141,:]))
test labels = np.concatenate((labels[:20], labels[121:141]))
train data = np.row stack((data[20:121,:], data[141:,:]))
train labels = np.concatenate((labels[20:121], labels[141:]))
iteration_count = np.array([500, 1000, 1500, 2000, 2500, 3000, 3500, 4000,
4500, 5000])
learning rate = np.array([0.001, 0.002, 0.005, 0.01, 0.015, 0.02, 0.025,
kf test data
                         np.asarray((train data[:35],
                                                        train data[35:70],
train data[70:105], train_data[105:140], train_data[140:]))
kf train data = np.asarray((train data[35:], np.concatenate((train data[:35],
train data[70:])),
                     np.concatenate((train data[:70], train data[105:])),
np.concatenate((train_data[:105], train_data[140:])), train_data[:140]))
kf_test_labels = np.asarray((train_labels[:35], train_labels[35:70],
```

```
train labels[70:105], train labels[105:140], train labels[140:]))
kf train labels
                                                np.asarray((train labels[35:],
np.concatenate((train labels[:35],
                                                         train labels[70:])),
np.concatenate((train labels[:70],
                                                         train labels[105:])),
np.concatenate((train labels[:105], train labels[140:])), train labels[:140]))
kf error ic = np.zeros(len(iteration count))
kf error lr = np.zeros(len(learning rate))
index = 0
for ic in iteration count:
    print("Iteration Count = " + str(ic))
    mse = np.zeros(len(kf train data))
    for i in range(len(kf train data)):
       print("k-Fold iteration " + str(i) + "...")
        (x train, y train) = kf train data[i].shape
        (x test, y test) = kf test data[i].shape
        (w0, weights) = gradientAscent(learning rate[0], ic, y train, x train,
kf train data[i], kf train labels[i])
              predictions = 1 / (1 + np.exp(np.multiply(kf_test_data[i],
weights).sum(axis = 1) + w0))
        for k in range(len(predictions)):
           predictions[k] = 0 if predictions[k] > 0.5 else 1
            # if predictions[k] != kf train labels[i][k]:
           mse[i] = mse[i] + (kf test labels[i][k] - predictions[k]) ** 2
    kf error ic[index] = np.sum(mse) * 0.2
    index = index + 1
for lr in learning rate:
   print("Learning Rate = " + str(lr))
   mse = np.zeros(len(kf train data))
    for i in range(len(kf train_data)):
       print("k-Fold iteration " + str(i) + "...")
        (x train, y train) = kf train data[i].shape
        (x test, y test) = kf test data[i].shape
            (w0, weights) = gradientAscent(lr, iteration count[0], y train,
x train, kf train data[i], kf train labels[i])
              predictions = 1 / (1 + np.exp(np.multiply(kf test data[i],
weights).sum(axis = 1) + w0))
        for k in range(len(predictions)):
            predictions[k] = 0 if predictions[k] > 0.5 else 1
            # if predictions[k] != kf train labels[i][k]:
           mse[i] = mse[i] + (kf test labels[i][k] - predictions[k]) ** 2
    kf error lr[index] = np.sum(mse) * 0.2
    index = index + 1
# k-Fold Results
kfold_ic = iteration_count[np.argmin(kf_error_ic)]
```

```
kfold lr = learning rate[np.argmin(kf error lr)]
(kfold sample count, kfold feature count) = test data.shape
(kfold w0, kfold weights) = gradientAscent(0.001, 2500, kfold feature count,
kfold sample count, test data, test labels)
kfold predictions = 1
                                               np.exp(np.multiply(test data,
                              / (1
kfold weights).sum(axis = 1) + kfold w0))
confusion matrix = {"tp": 0, "tn": 0, "fp": 0, "fn": 0}
for i in range(len(kfold predictions)):
    kfold predictions[i] = 0 if kfold predictions[i] > 0.5 else 1
    if test labels[i] == 1 and kfold predictions[i] == 1: # tp
        confusion matrix["tp"] = confusion matrix["tp"] + 1
    elif test labels[i] == 0 and kfold predictions[i] == 0: # tn
        confusion matrix["tn"] = confusion matrix["tn"] + 1
   elif test labels[i] == 0 and kfold predictions[i] == 1: # fp
        confusion matrix["fp"] = confusion matrix["fp"] + 1
    elif test labels[i] == 1 and kfold predictions[i] == 0: # fn
        confusion matrix["fn"] = confusion matrix["fn"] + 1
print("Iteration Count: " + str(kfold ic) + ", Learning Rate: " +
str(learning rate))
print("Confusion Matrix: ")
print(confusion matrix)
# Feature Selection
indices forward = forwardSelection(kf train data, kf train labels, kfold ic,
kfold lr)
print("Forward Selection...")
new test data = test data[:, indices forward]
(test sample, test feature) = new test data.shape
(w0, weights) = gradientAscent(kfold lr, kfold ic, test feature, test sample,
new test data, test labels)
predictions = 1 / (1 + np.exp(np.multiply(new test data, weights).sum(axis =
1) + w0))
confusion matrix = {"tp": 0, "tn": 0, "fp": 0, "fn": 0}
for i in range(len(predictions)):
   predictions[i] = 0 if predictions[i] > 0.5 else 1
   if test labels[i] == 1 and predictions[i] == 1: # tp
        confusion matrix["tp"] = confusion matrix["tp"] + 1
   elif test labels[i] == 0 and predictions[i] == 0: # tn
        confusion matrix["tn"] = confusion matrix["tn"] + 1
    elif test labels[i] == 0 and predictions[i] == 1: # fp
        confusion matrix["fp"] = confusion matrix["fp"] + 1
    elif test labels[i] == 1 and predictions[i] == 0: # fn
        confusion matrix["fn"] = confusion matrix["fn"] + 1
print("Confusion Matrix: ")
print(confusion matrix)
```

```
# Backward Elimination
indices backward
                        backwardElimination(kf train data,
                                                             kf train labels,
kfold ic, kfold lr)
print("Backward Elimination...")
new test data = test data[:, indices backward]
(test sample, test feature) = new test data.shape
(w0, weights) = gradientAscent(kfold lr, kfold ic, test feature, test sample,
new test data, test labels)
predictions = 1 / (1 + np.exp(np.multiply(new test data, weights).sum(axis =
1) + w0))
confusion matrix = {"tp": 0, "tn": 0, "fp": 0, "fn": 0}
for i in range(len(predictions)):
   predictions[i] = 0 if predictions[i] > 0.5 else 1
    if test labels[i] == 1 and predictions[i] == 1: # tp
        confusion matrix["tp"] = confusion matrix["tp"] + 1
    elif test labels[i] == 0 and predictions[i] == 0: # tn
        confusion matrix["tn"] = confusion matrix["tn"] + 1
    elif test labels[i] == 0 and predictions[i] == 1: # fp
        confusion matrix["fp"] = confusion matrix["fp"] + 1
    elif test labels[i] == 1 and predictions[i] == 0: # fn
        confusion matrix["fn"] = confusion matrix["fn"] + 1
print("Confusion Matrix: ")
print(confusion matrix)
Question 3
from sklearn import svm
import numpy as np
data = np.asarray(np.loadtxt("../data/UCI Breast Cancer.csv", dtype = "int32",
delimiter=','))[:, 1:]
labels = data[:, -1]
data = data[:, :-1]
# 3.3
train data = data[:500,:]
train labels = labels[:500]
test data = data[500:,:]
test labels = labels[500:]
kf test data = np.asarray((train data[:100],
                                                       train data[100:200],
train data[200:300], train data[300:400], train data[400:]))
kf train data
                                                 np.asarray((train_data[100:],
np.concatenate((train data[:100],
                                                           train data[200:])),
np.concatenate((train data[:200],
                                                           train data[300:])),
np.concatenate((train_data[:300], train_data[400:])), train_data[:400]))
kf test labels = np.asarray((train labels[:100], train labels[100:200],
train_labels[200:300], train_labels[300:400], train_labels[400:]))
                                              np.asarray((train labels[100:],
kf train labels
np.concatenate((train labels[:100],
                                                         train labels[200:])),
```

```
np.concatenate((train labels[:200],
                                                         train labels[300:])),
np.concatenate((train labels[:300], train labels[400:])), train labels[:400]))
C = np.array((10**-3, 10**-2, 10**-1, 1, 10**1, 10**2, 10**3))
errors = np.zeros(len(C))
index = 0
for c in C:
    mse = np.zeros(5)
    for i in range(5): # kfold
          model = svm.LinearSVC(C=c, dual=False, fit intercept=True, loss =
"squared hinge", max iter=1000, penalty = "12", random state=0, tol=1e-05)
        model.fit(kf train data[i], kf train labels[i])
        predictions = model.predict(kf test data[i])
        mse[i] = np.sum(np.power(kf test labels[i] - predictions, 2))
    errors[index] = float(np.sum(mse) / 5)
    index = index + 1
c = C[np.argmin(errors)]
print("Linear SVM with C = " + str(c))
            svm.LinearSVC(C=c, dual=False, fit intercept=True,
"squared hinge", max iter=1000, penalty = "12", random state=0, tol=1e-05)
model.fit(train data, train labels)
score = model.score(test data, test labels)
predictions = model.predict(test data)
confusion matrix = {"tp": 0, "tn": 0, "fp": 0, "fn": 0}
for i in range(len(test labels)):
    if test labels[i] == 4 and predictions[i] == 4: # tp
        confusion matrix["tp"] = confusion matrix["tp"] + 1
    elif test labels[i] == 2 and predictions[i] == 2: # tn
        confusion matrix["tn"] = confusion matrix["tn"] + 1
    elif test labels[i] == 2 and predictions[i] == 4: # fp
        confusion matrix["fp"] = confusion matrix["fp"] + 1
    elif test labels[i] == 4 and predictions[i] == 2: # fn
        confusion matrix["fn"] = confusion matrix["fn"] + 1
print("Confusion Matrix -> ")
print(confusion matrix)
print("Accuracy: " + str(score))
# 3.4
train data = data[:500,:]
train labels = labels[:500]
test data = data[500:,:]
test labels = labels[500:]
kf_test_data = np.asarray((train data[:100], train data[100:200],
train data[200:300], train data[300:400], train data[400:]))
kf train data
                                                 np.asarray((train data[100:],
np.concatenate((train data[:100],
                                                           train data[200:])),
np.concatenate((train data[:200],
                                                           train data[300:])),
```

```
np.concatenate((train data[:300], train data[400:])), train data[:400]))
kf test labels
               = np.asarray((train labels[:100], train labels[100:200],
train labels[200:300], train labels[300:400], train labels[400:]))
kf train labels
                                               np.asarray((train labels[100:],
np.concatenate((train labels[:100],
                                                         train labels[200:])),
np.concatenate((train labels[:200],
                                                         train labels[300:])),
np.concatenate((train labels[:300], train labels[400:])), train labels[:400]))
gammA = np.array((2**-4, 2**-3, 2**-2, 2**-1, 1, 2**1, 2**2, 2**3, 2**4))
errors = np.zeros(len(gammA))
index = 0
for gamma in gammA:
   mse = np.zeros(5)
    for i in range(5): # kfold
       model = svm.SVC(kernel = "rbf", gamma = gamma)
       model.fit(kf train data[i], kf train labels[i])
       predictions = model.predict(kf test data[i])
       mse[i] = np.sum(np.power(kf test labels[i] - predictions, 2))
    errors[index] = float(np.sum(mse) / 5)
    index = index + 1
gamma = gammA[np.argmin(errors)]
print("SVM with RBF kernel, gamma = " + str(gamma))
model = svm.SVC(kernel = "rbf", gamma = gamma)
model.fit(train data, train labels)
score = model.score(test data, test labels)
predictions = model.predict(test data)
confusion matrix = {"tp": 0, "tn": 0, "fp": 0, "fn": 0}
for i in range(len(test labels)):
    if test labels[i] == 4 and predictions[i] == 4: # tp
        confusion matrix["tp"] = confusion matrix["tp"] + 1
    elif test labels[i] == 2 and predictions[i] == 2: # tn
        confusion matrix["tn"] = confusion matrix["tn"] + 1
    elif test labels[i] == 2 and predictions[i] == 4: # fp
        confusion matrix["fp"] = confusion matrix["fp"] + 1
    elif test labels[i] == 4 and predictions[i] == 2: # fn
        confusion matrix["fn"] = confusion matrix["fn"] + 1
print("Confusion Matrix -> ")
print(confusion matrix)
print("Accuracy: " + str(score))
```