```
import numpy as np
import pandas as pd
import scipy as sp
import matplotlib.pyplot as plt
```

Problem 2 Code

2 Linear Regression with Regularization [10pts]

In class we derived and discussed linear regression in detail. Find the result of minimize the loss of sum of the squared errors; however, add in a penalty for an L_2 penalty on the weights. More formally,

$$\arg\min_{\mathbf{w}} \left\{ \sum_{i} (\mathbf{w}^{\mathsf{T}} \mathbf{x}_{i} - y_{i})^{2} + \lambda \|\mathbf{w}\|_{2}^{2} \right\}$$

How does this change the solution to the original linear regression solution? What is the impact of adding in this penalty?

Write your own implementation of logistic regression and implement your model on either real-world (see Github data sets: https://github.com/gditzler/UA-ECE-523-Sp2018/tree/master/data), or synthetic data. If you simply use Scikit-learn's implementation of the logistic regression classifier, then you'll receive zero points. A full 10/10 will be awarded to those that implement logistic regression using the optimization of cross-entropy using stochastic gradient descent.

Problem 2 functions

```
def split_dataset(dataset,train_percent):
    Input: Dataset : dataframe train_percent: percentage (0.0 - 1.0)
    Output: 4x dataframes: x-testing,x-training,y-testing,y-training
    #dataset = shuffle dataset(dataset)
    y = dataset.iloc[:,-1]
    x = dataset.iloc[:, 0:-1]
    test size = int(train percent * y.size)
    #create testing
    yts = y.iloc[:test size,].reset index(drop=True)
    xts = x.iloc[:test_size,].reset_index(drop=True)
    #create training
    ytr = y.iloc[test_size:,].reset_index(drop=True)
    xtr = x.iloc[test_size:,].reset_index(drop=True)
    return xts,xtr,yts,ytr
def shuffle dataset(xt,yt):
    Shuffles the data collectively
            xt : feature data
                                yt : class data
    Input:
              Separate feature and class vector in appropriate order
    Output:
    xy = pd.concat([xt, yt], axis=1)
```

```
xy2 = xy.sample(frac=1).reset index(drop=True)
    y = xy2.iloc[:,-1]
    x = xy2.iloc[:, 0:-1]
    return x,y
def logistic function(x,w):
    Logistic Function 1/(1+exp(-wTx))
    wTx = np.dot(x, w)
    temp = 1 / (1 + np.exp(-wTx))
    return temp
def cross entropy(x, y, w):
    Cross Entropy: log loss measure the converges
    Only works with binary class at the moment
    temp = - np.sum (y * np.log(logistic function(x,w)) + (1-y)*
np.log(1- logistic function(x,w)))
    return temp
def gradient(x, y, w, eta):
    Gradient f(x)f()
    temp = np.dot(x.T, logistic function(x,w) - y)
    return eta * temp
def sqd(xtr,ytr, T,eta):
    SGD: Logistic Regression using Stochastic Gradiant Descent
    Input: xtr:training feature data, ytr:training class data
           T: iterations, eta: learning rate
    Output: wnew: parameter vector, k: occurences cross enthropy
worked
   wnew = np.zeros((xtr.shape[1],1)) #initialize parameter vector
    n = len(ytr.iloc[:,]) # size of the training dataset
    ce = 1 #initialize cross enthropy val
    k = 0
    for i in range(T):
        xtr,ytr = shuffle dataset(xtr,ytr) #shuffle data each
iteration
        for j in range(n):
            wold = wnew
            x i = np.array([xtr.iloc[j]]) #set xtrain row to nparray
```

```
y i = np.array(ytr.iloc[j]) # set ytrain nparray
            wnew = wold - gradient(x i,y i,wold,eta) #get updated
parameters
            temp ce = cross entropy(x i, y i, wnew) #perform cross
etrophy
            ce dif = abs(temp ce - ce) #compare previous ce with new
ce
            if (ce dif < 0.00001): #break if minimum is found less</pre>
then 0.00001
                ce = 1
                k=k+1
                                #count occurences of ce
                break
            else:
                ce = temp ce #progress ce
    return wnew, k
Problem 2 Test Program
path = "/content/gdrive/MyDrive/Git/ECE523/HW2/acute_inflamation.csv"
dataset = pd.read_csv(path, header=None)
#split dataset into testing and traing
xtest,xtrain,ytest,ytrain = split dataset(dataset,.6)
#train w parameters
wtr,k = sgd(xtrain,ytrain,1000,0.02)
#test w parameters against xtest feature data
vhat = logistic function(xtest,wtr)
for i in range(\overline{l}en(yhat)):
    if yhat[i] >=0.5:
        vhat[i] = 1
    else:
        yhat[i] = 0
print("w parameters: \n", wtr)
print("k occurences using Cross Entropy: ",k)
dif y = yhat[:,0] - ytest[:]
perc correct = (1.0 - (float(np.count nonzero(dif y))/len(dif y)))*100
print("Precent Accuracy with test data: ", perc correct)
print("\nModify ytest[2] value (1 out of %s) to confirm SGD
working..." % ytest.size)
if ytest[2]==1:
    ytest[2]=0
else:
    ytest[2]=1
```

```
dif_y = yhat[:,0] - ytest[:]
perc correct = (1.0 - (float(np.count nonzero(dif y))/len(dif y)))*100
print("Precent Accuracy with altered test data: ", perc correct)
print(ytest.size)
w parameters:
 [[-2.78368288]
 [ 1.99715886]
 [-2.0301456]
 [ 5.46981692]
 [ 3.22064441]
 [-0.13217954]]
k occurences using Cross Entropy:
                                  610
Precent Accuracy with test data:
                                  100.0
Modify ytest[2] value (1 out of 72) to confirm SGD working...
Precent Accuracy with altered test data: 98.61111111111111
72
```

Problem 3 Code

3 Density Estimation [20pts]

The ECE523 Lecture notes has a function for generating a checkerboard data set. Generate checkerboard data from two classes and use any density estimate technique we discussed to classify new data using

$$\widehat{p}_{Y|X}(y|x) = \frac{\widehat{p}_{X|Y}(x|y)\widehat{p}_{Y}(y)}{\widehat{p}_{X}(x)}$$

where $\hat{p}_{Y|X}(y|x)$ is your estimate of the posterior given you estimates of $\hat{p}_{X|Y}(x|y)$ using a density estimator and $\hat{p}_{Y}(y)$ using a maximum likelihood estimator. You should plot $\hat{p}_{X|Y}(x|y)$ using a pseudo color plot (see https://goo.gl/2SDJPL). Note that you must model $\hat{p}_{X}(x)$, $\hat{p}_{Y}(y)$, and $\hat{p}_{X|Y}(x|y)$. Note that $\hat{p}_{X}(x)$ can be calculated using the Law of Total Probability.

Problem 3 Functions

```
lab = 2 - (s%2)
    data = d.T
    return data, lab
def algorithm k NN(k,point,dataset):
    k = k number of point to compare to
    xt point = test data point
    x = x1, x2 feature data
    v = classifiers
    #Add distance from test point to training points in datasets to
datasets
    dataset['distance'] = np.sqrt((dataset['x1'] - point[0]) ** 2 +
                                          (dataset['x2'] - point[1]) **
2)
    #Sort Distances in Dataset in ascending order
    dataset.sort values(by=['distance'],inplace=True)
    #create a set of closest K distances
    k set = dataset.iloc[:k]
    k_list = k_set['y1'].to_numpy()
    #Take largest distance for kmax and set to radius
    radius = k set['distance'].max()
    #len of training data
    y1 len = len(dataset.y1)
    \#Total\ Count\ of\ training\ where\ y\ ==\ 1,2
    n1 = (dataset.y1.values == 1).sum()
    n2 = (dataset.y1.values == 2).sum()
    #Count when K == 1,2
    k1, k2 = 0, 0
    if (1.0 in k_list) == True:
        k1 = k_set['y1'].value_counts()[1]
    if (2.0 in k list) == True:
        k2 = k set['y1'].value counts()[2]
    #calculate volume
    volume = np.pi * (radius**2)
    #p(x|y=1)
    pxy1 = k1/(n1*volume)
    #p(x|y=2)
    pxy2 = k2/(n2*volume)
    \#p(y=1) NOTE: n1 equals total 1's in y1
    py1 = n1/y1 len
    \#p(y=2) NOTE: n2 equals total 2's in y1
```

```
py2 = n2/y1 len
    \#p(x) both equate
    pxtot = ((pxy1*py1)+(pxy2*py2))
    px = k/((n1+n2)*volume)
    #p(y=1|x)
    py1x = (pxy1*py1)/px
    #p(y=2|x)
    py2x = (pxy2*py2)/px
    #compare and output
    if py1x > py2x:
        return 1, py1x, py2x
    else:
        return 2, py1x, py2x
Problem 3 Test Program
N = 1500
x,y = gen checkerboard(N, .25, 3.14159/4)
xtest, yt = gen checkerboard(N, .25, 3.14159/4)
k = int(np.sqrt(N))
#organize training dataset
dataset = pd.DataFrame(x,columns=['x1','x2'])
dataset['y1'] = y
#Create v test output set
ytest = np.zeros(len(yt))
#Go through each test point and run Knn
i=0
for xp in xtest:
    ytest[i],pnan,qnan = algorithm k NN(k,xp,dataset)
    i+=1
# Using the bounds of the test feature vector,
# generate 50x50 set for each possible point
xmin, xmax = np.min(xtest[:,0]), np.max(xtest[:,0])
ymin, ymax = np.min(xtest[:,1]),np.max(xtest[:,1])
xm, ym = np.mgrid[xmin:xmax:50j, ymin:ymax:50j]
\#Generate\ empty\ posterior\ sets\ same\ shape\ as\ xm,ym\ (50x50)
pxy1 = np.zeros(xm.shape)
pxy2 = np.zeros(xm.shape)
#Iterate through each point on grid, and collect the posterior for 1
or 2
for i in range(len(xm[0])):
    for j in range(len(xm[1])):
        ynan, pxy1[i,j], pxy2[i,j] = algorithm k NN(k,
[xm[i,j],ym[i,j]],dataset)
```

```
#Plot
plt.figure()
plt.plot(x[np.where(y==2)[0],0],x[np.where(y==2)[0],1],'s',c = 'r')
plt.plot(x[np.where(y==1)[0],0],x[np.where(y==1)[0],1],'o')
plt.xlabel("X1")
plt.ylabel("X2")
plt.title('Original Training Data')
plt.show()
plt.plot(xtest[np.where(ytest==2)[0],0],xtest[np.where(ytest==2)
[0],1],'s',c = 'r')
plt.plot(xtest[np.where(ytest==1)[0],0],xtest[np.where(ytest==1)
[0],1],'o')
plt.title('Test Training Data')
plt.xlabel("X1")
plt.ylabel("X2")
plt.show()
#Plot posterior probabilities for each point in grids
plt.figure()
plt.pcolormesh(xm,ym,pxy1,cmap='jet')
plt.colorbar()
plt.title('P(y=1|x) Mesh Data')
plt.xlabel("X1")
plt.ylabel("X2")
plt.show()
plt.figure()
plt.pcolormesh(xm,ym,pxy2,cmap='jet')
plt.colorbar()
plt.title('P(y=2|x) Mesh Data')
plt.xlabel("X1")
plt.ylabel("X2")
plt.show()
```







