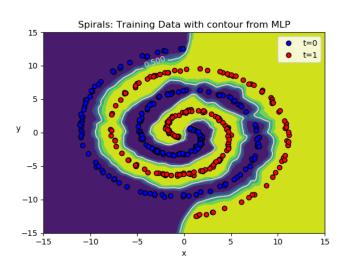
## Selected Topics in Machine Learning - Assignment 2

```
#!/bin/python3.6
import numpy as np
import tensorflow as tf
from tqdm import trange
import matplotlib.pyplot as plt
# I credit these resources with good help for this assignment (in addition to tensorflow docs of course!):
# https://developers.google.com/machine-learning/crash-course/introduction-to-neural-networks/anatomy
# https://www.jessicayung.com/explaining-tensorflow-code-for-a-multilayer-perceptron/
# https://towardsdatascience.com/sigmoid-activation-and-binary-crossentropy-a-less-than-perfect-match-
b801e130e31
# https://rdipietro.github.io/friendly-intro-to-cross-entropy-loss/
BATCH SIZE = 50
NUM BATCHES = 10000
LEARNING RATE = 0.1
NUM SAMP = 200
class Data(object):
   def __init__(self):
       np.random.seed(31415)
       sigma = 0.1
       self.index = np.arange(2*NUM SAMP)
       # spirals made to replicate given graph
       self.t1 = np.random.uniform(np.pi/4, 4*np.pi, NUM_SAMP)
       self.x1 = np.array([ self.t1*np.sin(self.t1) + np.random.normal(0, sigma, NUM SAMP),
                             self.t1*np.cos(self.t1)+ np.random.normal(0, sigma, NUM SAMP) ]).T
       \# rotate the previously generated spiral by 180 degrees to produce second spiral
       self.t2 = np.random.uniform(np.pi/4, 4*np.pi, NUM_SAMP)
       self.x2 = np.array([ self.t2*np.sin(self.t2 + np.pi) + np.random.normal(0, sigma, NUM_SAMP),
                             self.t2*np.cos(self.t2 + np.pi)+np.random.normal(0, sigma, NUM SAMP) ]).T
       self.coordinates = np.concatenate((self.x1, self.x2))
       self.labels = np.concatenate((np.zeros(NUM_SAMP), np.ones(NUM_SAMP)))
   def get batch(self):
       choices = np.random.choice(self.index, size = BATCH SIZE)
       return self.coordinates[choices,:], self.labels[choices].flatten()
class Model(tf.Module):
   def init (self, dimensions x = 2, dimensions layer 1 = 40, dimensions layer 2 = 40):
        self.weights = {
            'w1': tf.Variable(tf.random.normal(shape=[dimensions_x, dimensions_layer_1]), dtype= tf.float32),
            'w2': tf.Variable(tf.random.normal(shape=[dimensions_layer_1, dimensions_layer_2]), dtype=
tf.float32),
            'out': tf.Variable(tf.random.normal(shape=[dimensions layer 2, 1]), dtype= tf.float32)
       self.biases = {
            'b1': tf.Variable(tf.random.normal(shape=[dimensions layer 1]), dtype= tf.float32),
            'b2': tf.Variable(tf.random.normal(shape=[dimensions_layer_2]), dtype= tf.float32),
            'out': tf.Variable(tf.zeros(shape=[]), dtype= tf.float32)
   def f(self, x coords):
        layer one = tf.nn.relu6(tf.add( tf.matmul(x coords, self.weights['w1']), self.biases['b1']))
        layer two = tf.nn.relu6(tf.add( tf.matmul(layer one, self.weights['w2']), self.biases['b2'] ))
       output = tf.add( tf.matmul(layer_two, self.weights['out']), self.biases['out'])
       return tf.squeeze(output)
   def loss(self, coords, labs):
        \# Sigmoid can be used for interpreting the output of f:V 	ext{ } 	ext{-} 	ext{>} 	ext{ } 	ext{R} function as a probability,
        \# i.e using it's squashing properties to go R -> [0,1]. You can use the sigmoid to compute the
conditional density.
        # logits are unnormalized probabilities (i.e they exist in R not [0,1]) and it internally computes
        the sigmoid #for numerical reasons
        loss_p1 = tf.reduce_mean(tf.nn.sigmoid_cross_entropy_with_logits(labels = labs, logits =
self.f(coords)))
```

```
# standard linear regression problem: y \text{ hat} = w^t x + b, loss function = 0.5*(y - w^t x - b)
       # L2 penalty --> modify the loss function to 0.5*(y - w^t x - b) + lambda*norm(w)**2
         norm --> the L2 norm
         lambda --> hyperparamter where 0 removes regularization (training is then minimizing loss, risk of
overfitting)
       \#12 loss function --> output = sum(t ** 2) / 2
       loss_p2 = (.001) * tf.reduce_sum([tf.nn.12_loss(var) for var in model.weights.variables])
       loss = loss p1 + loss p2
       return loss
if __name__ == "__main__":
   data = Data()
   model = Model()
   optimizer
   bar = trange(NUM_BATCHES) # creates bar visual to show progress completion status
   for i in bar:
       with tf.GradientTape() as tape: # GradientTape= Record operations for automatic differentiation.
           coords, labs = data.get_batch()
           loss fromBatch = model.loss(coords.astype(np.float32), labs.astype(np.float32))
           grads = tape.gradient(loss fromBatch, model.variables)
           optimizer.apply gradients(zip(grads, model.variables))
   xAxis = np.linspace(-15, 15, 150)
   yAxis = np.linspace(-15, 15, 150)
   X, Y = np.meshgrid(xAxis, yAxis)
   Z = np.zeros((150, 150))
   for i in range (150):
       for j in range(150):
           Zarr = np.array([X[i,j], Y[i,j]]).reshape(1,2)
           Zval = model.f(Zarr.astype(np.float32))
           Z[i,j] = tf.nn.sigmoid(Zval)
   plt.figure()
   plt.plot(data.x1[:,0], data.x1[:,1], 'ob', markeredgecolor="black")
   plt.plot(data.x2[:,0], data.x2[:,1], 'or', markeredgecolor="black")
   point_five_line = plt.contour(X, Y, Z, levels = [0.5], colors = "pink")
   plt.clabel(point five line, inline = 1, fontsize = 10, colors='pink')
   plt.legend(["t=0", "t=1"])
                                                                 To find the functional form of f I did the computations as seen
   plt.contourf(X, Y, Z)
   plt.title("Spirals: Training Data with contour from MLP")
   plt.xlabel('x')
```



plt.ylabel('y',rotation=0)

plt.show()

in the method f in class Model. Essentially there are three layers total, in which the coordinates are fed into and a final output is produced. For each layer the output of the previous layer (or the coordinates for the first layer) is multiplied by the weights of that layer and added to the bias of that layer.

Relu6 was a sufficient activation function, and the number of layers used was appropriate for the data we have. If more layers were introduced there would be overfitting.

In the code, a pdf was essentially generated to determine the probability of it being red, and this is what was used to classify the results.

In working on the problem, the lambda was important in determining the quality of the contour produced, it tended to produce a better classification when the lambda was near

There was flexibility in the batch size and number of samples, but the number of samples I selected to resemble the sample

The number of features chosen for each layer at 40 produced adequate results.

The pink line on the graph demonstrates where the line between equal probability lies.