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Homework 9

Part 1: Losses

Problem 1.1

Use the formulas in the class notes to write functions with headers

```
def regressionLoss(z, delta):
def hingeLoss(y, delta):
```

that take a label $z \in Z = \{0,1\}$ or $y \in Y = \{-1,1\}$ and a value δ for the signed distance from the separating hyperplane and compute the logistic-regression loss and the hinge loss as functions of label and δ . Keep in mind that the logistic-regression loss is a composition of logistic function and cross-entropy loss.

Also write a function with header

```
def plotLosses(y):
```

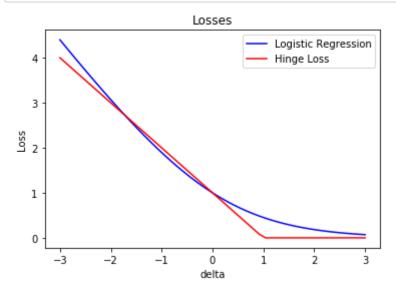
that takes a label $y \in Y = \{-1, 1\}$ and plots the two loss functions for $-3 \le \delta \le 3$. Show your code and the two plots that result from calling plotLosses, first with argument 1 and then with argument -1.

```
In [1]: import numpy as np
        import matplotlib.pyplot as plt
        %matplotlib inline
        def regressionLoss(z, delta):
            This takes label z in \{0,1\} and a value delta for the signed distanc
        e from the separating
            hyperplane and computes the logistic-regression loss of label and de
        1ta
                                                      # logistic function p
            p = 1 / (1 + np.exp(-delta))
            x_{loss} = -z*np.log2(p) - (1-z)*np.log2(1-p) # cross entropy loss, z
         is the label {0,1}, p is the logistic function
            return x loss
        def hingeLoss(y, delta):
            this takes label y in {-1,1} and a value delta for the signed distan
        ce from the separating
            hyperplane and compute the hinge loss as functions of label and delt
        a.
            loss = max([0, 1 - y*delta])
            return loss
        def plotLosses(y):
            This function takes a label y in {-1, 1} and plots the two loss func
        tions
            for -3 <= delta <= 3
            # converting y's to z's for logistic regression function which takes
         z in {0, 1}
            if y == -1: # this takes y = -1 and makes z = 0
                z = 0
            if y == 1: # this takes y = 1 and makes z = 1
            deltas = np.linspace(-3, 3, 50) # generates a list of 50 deltas (sig
        ned distnaces) between -3 and 3
            reg losses = []
                                            # initialize list of logistic regres
        sion losses
                                             # initialize list of hinge losses
            hinge losses = []
            for delta in deltas:
                reg loss = regressionLoss(z, delta)
                reg losses.append(reg loss)
                hinge_loss = hingeLoss(y, delta)
                hinge losses.append(hinge loss)
            plt.plot(deltas, reg losses, label = "Logistic Regression", color =
```

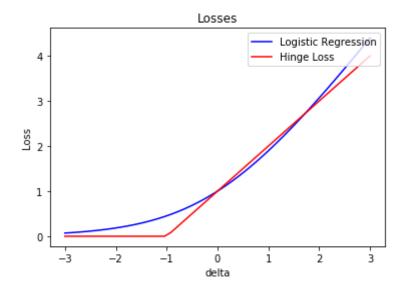
```
"b")
    plt.plot(deltas, hinge_losses, label = "Hinge Loss", color = "r")
    plt.title("Losses")
    plt.xlabel("delta")
    plt.ylabel("Loss")
    plt.legend(loc = "upper right")

    return

# y = 1
test_y = 1
plotLosses(test_y)
```



```
In [2]: # y = -1
    test_y = -1
    plotLosses(test_y)
```



Problem 1.2

Suppose that there is a single data outlier that is misclassified by a very large (negative) margin. Referring to the plots in Problem 1.1, which of the two losses is more sensitive to that outlier, and why?

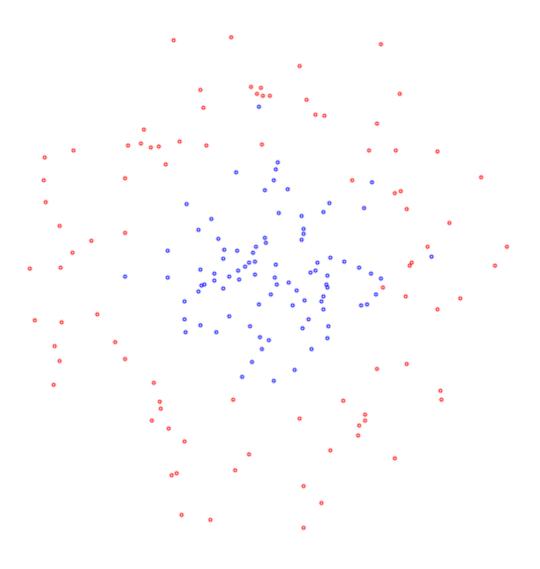
Answer

The logistic regression loss is more sensitive to a very large (negative) margin because the loss is exponential decreasing or increasing and the hinge loss is either 0 or linear. On both graphs, the logistic regression loss is greater than the hinge loss.

Part 2: SVMs

In [3]: import numpy as np import matplotlib.pyplot as plt import sklearn.datasets as ds from sklearn.model_selection import train_test_split %matplotlib inline ns = 300 $data = \{\}$ data['x'], data['y'] = ds.make circles(n samples=ns, noise=0.2, factor=0.3, random_state=1) testFraction = 0.6 $T, S = \{\}, \{\}$ T['x'], S['x'], T['y'], S['y'] = train_test_split(data['x'], data['y'], test_size=testFraction, random_state=0) p, n = S['x'][S['y']==1], S['x'][S['y']!=1]plt.figure(figsize=(10,10)) plt.plot(p[:, 0], p[:, 1], '.b', fillstyle='none') plt.plot(n[:, 0], n[:, 1], '.r', fillstyle='none') plt.axis('equal') plt.axis('off')

Out[3]: (-1.409118932486272, 1.4707802536886163, -1.509294151919646, 1.4493604864382867)



Problem 2.1

The data in T and S is obviously not linearly separable, so a linear classifier should not be expected to do well. To verify this, train sklearn.svm.svc with arguments kernel='linear', C=1 on T, show its zero-one training and test error rates (on S) as percentages with two decimal digits after the period, and plot the data in T and decision regions. Warning: Most points will be support vectors for this first plot. This is OK.

```
In [4]: from sklearn.svm import SVC
        from matplotlib import colors
        import math
        # train sklearn.svm.SVC with arguments kernel = 'linear', C = 1 on T
        # show its zero-one training and test error rates (on S) as percentages
         with two decimal places
        clf = SVC(C = 1, kernel = 'linear') # training a linear Support Vector C
        lassification
        clf.fit(T['x'], T['y'])
                                           # fit the classifier to the training
         data
        zero_one_training_rate = 1 - clf.score(T['x'], T['y'])
        test error rate = 1 - clf.score(S['x'], S['y'])
                                                                   # test the cla
        ssifier on the test set and get a score
        print("Zero-one training error rate: {:.2f}".format(zero_one_training_ra
        te*100), "%")
        print("Test error rate: {:.2f}".format(test_error_rate*100), "%")
        # plotting the data in T and the decision regions
        # find the maximum and minimum x and y value (4 numbers)
        max_y = -math.inf
        max x = -math.inf
        min y = math.inf
        min x = math.inf
        for i in range(len(p)):
            # min and max x
            if T['x'][i][0] < min x: # check if x value in T is less than our cu</pre>
        rrent lowest x value
                min_x = T['x'][i][0]
            if T['x'][i][0] > max x: # check if x value in T is greater than our
         current highest x value
                \max_{x} = T['x'][i][0]
            if T['x'][i][1] < min y: # check if y value in T is less than our cu</pre>
        rrent lowest y value
                min_y = T['x'][i][1]
            if T['x'][i][1] > max y: # check if y value in T is greater than our
         current highest y value
                max_y = T['x'][i][1]
        \max x = \max x + 0.5 \# add 0.5 to the \max x value
        min_x = min_x - 0.5 \# subtract 0.5 to the min x value
        max y = max y + 0.5 \# subtract 0.5 to the max y value
        min y = min y - 0.5 # subtract 0.5 from the min y value
        x range = np.linspace(min x, max x, (max x - min x) / 0.02) # create a 1
        ist of the x range
        y_range = np.linspace(min_y, max_y, (max_y - min_y) / 0.02) # create a 1
        ist of the y range
        X, Y = np.meshgrid(x range, y range) # create the meshgrid of X, Y
```

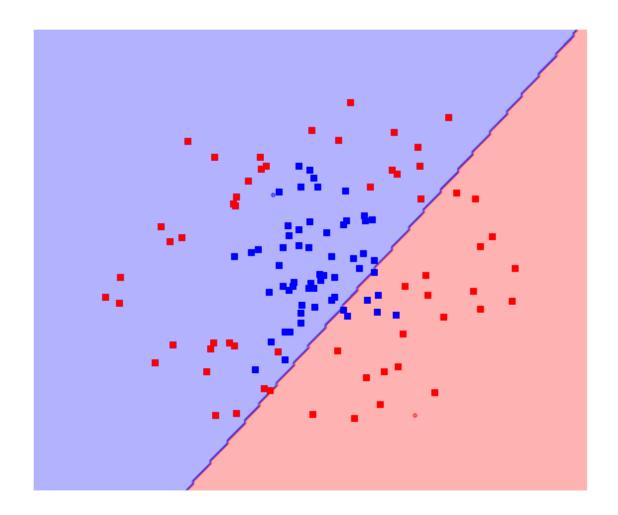
```
#flatten
X_flat = np.concatenate(X).ravel()
Y flat = np.concatenate(Y).ravel()
#stack x and y
XY = np.stack((X_flat, Y_flat), axis=-1)
predicted = clf.predict(XY).reshape((160, 193))
predicted_labels = clf.predict(T['x'])
a, b = T['x'][T['y']==1], T['x'][T['y']!=1]
# initialize the lists
true_pos_x = []
true_pos_y = []
true_neg_x = []
true_neg_y = []
false_pos_x = []
false_pos_y = []
false_neg_x = []
false_neg_y = []
svm_pos_x = []
svm_pos_y = []
svm neg x = []
svm_neg_y = []
svm indices = clf.support # indices for the support vectors
size\_of\_T = len(T['x'])
for sample index in range(size of T):
    sample = T['x'][sample_index]
                                    # sample coordinate
    true_label = T['y'][sample_index] # true label
    predicted label = predicted labels[sample index] # predicted label
    # check svm indices
    if sample index in svm indices:
        # current sample is an svm if this goes through
        if true label == 1:
            svm pos x.append(sample[0])
            svm pos y.append(sample[1])
        if true label != 1:
            svm neg x.append(sample[0])
            svm_neg_y.append(sample[1])
    # true positive
    if true label == 1 and predicted label == 1:
        true pos x.append(sample[0])
        true pos y.append(sample[1])
    # true negative
    if true_label != 1 and predicted_label != 1:
        true neg x.append(sample[0])
```

```
true neg y.append(sample[1])
    # false positive
    if true_label != 1 and predicted_label == 1:
        false pos x.append(sample[0])
        false pos y.append(sample[1])
    # false negatives
    if true label == 1 and predicted label != 1:
        false neg x.append(sample[0])
        false_neg_y.append(sample[1])
plt.figure(figsize=(10,10))
plt.plot(true_pos_x, true_pos_y, '.b', fillstyle ='none')
plt.plot(true_neg_x, true_neg_y, '.r', fillstyle ='none')
plt.plot(false_pos_x, false_pos_y, 'xr', fillstyle = 'none')
plt.plot(false_neg_x, false_neg_y, 'xb', fillstyle ='none')
plt.plot(svm_pos_x, svm_pos_y, 'sb', fillstyle = None)
plt.plot(svm_neg_x, svm_neg_y, 'sr', fillstyle = None)
plt.axis('equal')
plt.axis('off')
h = plt.contourf(X,Y,predicted, alpha=0.3, cmap = colors.ListedColormap
(['r','b'])) #check colors
```

Zero-one training error rate: 37.50 %
Test error rate: 41.11 %

/Users/joycechoi/anaconda3/lib/python3.6/site-packages/ipykernel_launch er.py:43: DeprecationWarning: object of type <class 'numpy.float64'> ca nnot be safely interpreted as an integer.

/Users/joycechoi/anaconda3/lib/python3.6/site-packages/ipykernel_launch er.py:44: DeprecationWarning: object of type <class 'numpy.float64'> ca nnot be safely interpreted as an integer.



Problem 2.2

Do the same with kernel='rbf' (same value for C as before).

```
In [5]: from sklearn.svm import SVC
        from matplotlib import colors
        import math
        # train sklearn.svm.SVC with arguments kernel = 'rbf', C = 1 on T
        # show its zero-one training and test error rates (on S) as percentages
         with two decimal places
        clf = SVC(C = 1, kernel = 'rbf') # training a linear Support Vector Clas
        sification
        clf.fit(T['x'], T['y'])
                                            # fit the classifier to the training
         data
        zero_one_training_rate = 1 - clf.score(T['x'], T['y'])
        test error rate = 1 - clf.score(S['x'], S['y'])
                                                                   # test the cla
        ssifier on the test set and get a score
        print("Zero-one training error rate: {:.2f}".format(zero_one_training_ra
        te*100), "%")
        print("Test error rate: {:.2f}".format(test_error_rate*100), "%")
        # plotting the data in T and the decision regions
        # find the maximum and minimum x and y value (4 numbers)
        max_y = -math.inf
        max x = -math.inf
        min y = math.inf
        min x = math.inf
        for i in range(len(p)):
            # min and max x
            if T['x'][i][0] < min x: # check if x value in T is less than our cu</pre>
        rrent lowest x value
                min_x = T['x'][i][0]
            if T['x'][i][0] > max x: # check if x value in T is greater than our
         current highest x value
                \max_{x} = T['x'][i][0]
            if T['x'][i][1] < min y: # check if y value in T is less than our cu</pre>
        rrent lowest y value
                min_y = T['x'][i][1]
            if T['x'][i][1] > max y: # check if y value in T is greater than our
         current highest y value
                max_y = T['x'][i][1]
        \max x = \max x + 0.5 \# add 0.5 to the \max x value
        min_x = min_x - 0.5 \# subtract 0.5 to the min x value
        max y = max y + 0.5 \# subtract 0.5 to the max y value
        min y = min y - 0.5 # subtract 0.5 from the min y value
        x range = np.linspace(min x, max x, (max x - min x) / 0.02) # create a 1
        ist of the x range
        y_range = np.linspace(min_y, max_y, (max_y - min_y) / 0.02) # create a 1
        ist of the y range
        X, Y = np.meshgrid(x range, y range) # create the meshgrid of X, Y
```

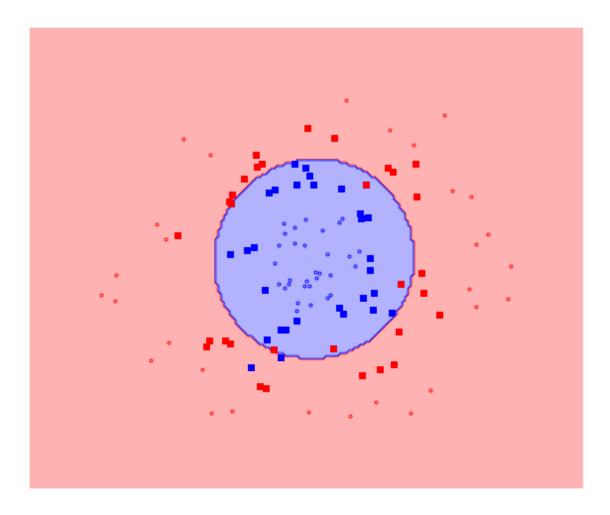
```
#flatten
X_flat = np.concatenate(X).ravel()
Y flat = np.concatenate(Y).ravel()
#stack x and y
XY = np.stack((X_flat, Y_flat), axis=-1)
predicted = clf.predict(XY).reshape((160, 193))
predicted_labels = clf.predict(T['x'])
a, b = T['x'][T['y']==1], T['x'][T['y']!=1]
# initialize the lists
true_pos_x = []
true_pos_y = []
true_neg_x = []
true_neg_y = []
false_pos_x = []
false_pos_y = []
false_neg_x = []
false_neg_y = []
svm_pos_x = []
svm_pos_y = []
svm neg x = []
svm_neg_y = []
svm indices = clf.support # indices for the support vectors
size\_of\_T = len(T['x'])
for sample index in range(size of T):
    sample = T['x'][sample_index]
                                    # sample coordinate
    true_label = T['y'][sample_index] # true label
    predicted label = predicted labels[sample index] # predicted label
    # check svm indices
    if sample index in svm indices:
        # current sample is an svm if this goes through
        if true label == 1:
            svm pos x.append(sample[0])
            svm pos y.append(sample[1])
        if true label != 1:
            svm neg x.append(sample[0])
            svm_neg_y.append(sample[1])
    # true positive
    if true label == 1 and predicted label == 1:
        true pos x.append(sample[0])
        true pos y.append(sample[1])
    # true negative
    if true label != 1 and predicted label != 1:
        true neg x.append(sample[0])
```

```
true neg y.append(sample[1])
    # false positive
    if true_label != 1 and predicted_label == 1:
        false pos x.append(sample[0])
        false pos y.append(sample[1])
    # false negatives
    if true label == 1 and predicted label != 1:
        false neg x.append(sample[0])
        false_neg_y.append(sample[1])
plt.figure(figsize=(10,10))
plt.plot(true_pos_x, true_pos_y, '.b', fillstyle ='none')
plt.plot(true_neg_x, true_neg_y, '.r', fillstyle ='none')
plt.plot(false_pos_x, false_pos_y, 'xr', fillstyle = 'none')
plt.plot(false_neg_x, false_neg_y, 'xb', fillstyle ='none')
plt.plot(svm_pos_x, svm_pos_y, 'sb', fillstyle = None)
plt.plot(svm_neg_x, svm_neg_y, 'sr', fillstyle = None)
plt.axis('equal')
plt.axis('off')
h = plt.contourf(X,Y,predicted, alpha=0.3, cmap = colors.ListedColormap
(['r','b'])) #check colors
```

Zero-one training error rate: 5.00 %
Test error rate: 3.33 %

/Users/joycechoi/anaconda3/lib/python3.6/site-packages/ipykernel_launch er.py:43: DeprecationWarning: object of type <class 'numpy.float64'> ca nnot be safely interpreted as an integer.

/Users/joycechoi/anaconda3/lib/python3.6/site-packages/ipykernel_launch er.py:44: DeprecationWarning: object of type <class 'numpy.float64'> ca nnot be safely interpreted as an integer.



Problem 2.3

Each data point in T and S has two features, x_1 and x_2 . Augment these by adding the following redundant features

$$x_3 = x_1^2$$

$$x_4 = x_2^2$$

$$x_5 = x_1 x_2$$

Then repeat the experiment above with SVC(kernel='linear', C=1) on the augmented data. Of course, plot just x_1 and x_2 , not the other features, unless you have a 5D printer handy.

```
In [6]: # train sklearn.svm.SVC with arguments kernel = 'linear', C = 1 on aug T
        _x (Augmented training set T )
        # show its zero-one training and test error rates (on S) as percentages
         with two decimal places
        # finding the max and min along each dimension for plotting the contour
        x1 dimension = T['x'][:, 0] # get all the values of T['x'] along the x1
         dimension
        x2_dimension = T['x'][:, 1] # get all the values of T['x'] along the x2
         dimension
        \max x1 = \max(x1 \text{ dimension}) \# \max value of x1
        min x1 = min(x1 dimension) # min value of x1
        \max x^2 = \max(x^2 \text{ dimension}) \# \max value \ of \ x^2
        min_x2 = min(x2_dimension) # min_value of x2
        # add or subtract from each max or min respectively
        \max x1 = \max x1 + 0.5 \# add 0.5 to the \max x1 value
        min x1 = min x1 - 0.5 # subtract 0.5 to the min x1 value
        \max x2 = \max x2 + 0.5 \# add 0.5 to the \max x2 value
        min_x2 = min_x2 - 0.5 \# subtract 0.5 from the min_x2 value
        x1 range = np.linspace(min x1, max x1, (max x1 - min x1) / 0.02) # creat
        e a list of the x1 range
        x2_range = np.linspace(min_x2, max_x2, (max_x2 - min_x2) / 0.02) # creat
        e a list of the x2 range
        X1, X2 = np.meshgrid(x1 range, x2 range)
                                                                           # creat
        e the meshgrid of X1, X2
        #flatten X1 and X2 to make it useable in np.stack
        X1 flat = np.concatenate(X1).ravel()
        X2 flat = np.concatenate(X2).ravel()
        #stack X1 and X2. Creates an array an (dim(X1) x dim(X2), 2) array
        stack = np.stack((X1 flat, X2 flat), axis=-1)
        # augment the stack with x3, x4, and x5
        stack length = len(stack)
                                                             # stack length or nu
        mber of entries in stack
        augmented_training_set = np.zeros((len(T['x']), 5)) # the same length as
         T['x'] but 5-d as opposed to 2-d
        augmented_test_set = np.zeros((len(S['x']), 5)) # the same length as
         S['x'] but 5-d as opposed to 2-d
        augmented T length = len(augmented training set) # length of augmente
        d T
        augmented S length = len(augmented test set)
                                                            # length of augmente
        augmented stack = np.zeros((stack length, 5)) # initialize the aug
        mented stack
        \# loop through the stack to create x3, x4, and x5 and put those values i
        nto the augmented stack
        for sample index in range(stack length):
            x1 = stack[sample index][0]
            x2 = stack[sample index][1]
```

```
x3 = x1**2
    x4 = x2**2
    x5 = x1*x2
    augmented stack[sample index][0] = x1
    augmented stack[sample index][1] = x2
    augmented_stack[sample_index][2] = x3
    augmented stack[sample index][3] = x4
    augmented stack[sample index][4] = x5
for sample index in range(augmented T length):
    sample = T['x'][sample index]
    x1 = sample[0] # looks at the first coordinate x1 of the sample at s
ample index in T['x']
    x2 = sample[1] # looks at the second coordinate x2 of the sample at
 sample index in T['x']
   x3 = x1**2
    x4 = x2**2
    x5 = x1*x2
    augmented_training_set[sample_index][0] = x1
    augmented training set[sample index][1] = x2
    augmented training set[sample index][2] = x3
    augmented training set[sample index][3] = x4
    augmented training set[sample index][4] = x5
for sample_index in range(augmented_S_length):
    sample = S['x'][sample index]
    x1 = sample[0] # looks at the first coordinate x1 of the sample at s
ample index in S['x']
    x2 = sample[1] # looks at the second coordinate x2 of the sample at
 sample index in S['x']
    x3 = x1**2
   x4 = x2**2
    x5 = x1*x2
    augmented test set[sample index][0] = x1
    augmented_test_set[sample index][1] = x2
    augmented test set[sample index][2] = x3
    augmented_test_set[sample_index][3] = x4
    augmented test set[sample index][4] = x5
clf = SVC(C = 1, kernel = 'linear')
  # training a linear Support Vector Classification
clf.fit(augmented training set, T['y'])
  # fit the classifier to the training data
zero one training rate = 1 - clf.score(augmented training set, T['y'])
  # train the classifier on augmented training set aug_T_x and get error
test error rate = 1 - clf.score(augmented test set, S['y'])
  # test the classifier on the augmented test set aug S x and get error
 rate
print("Zero-one training error rate: {:.2f}".format(zero one training ra
te*100), "%")
print("Test error rate: {:.2f}".format(test error rate*100), "%")
predicted = clf.predict(augmented stack).reshape(X1.shape) # train the c
lassifier on the augmented stack
predicted_labels = clf.predict(augmented_training_set)
                                                          # get the pre
```

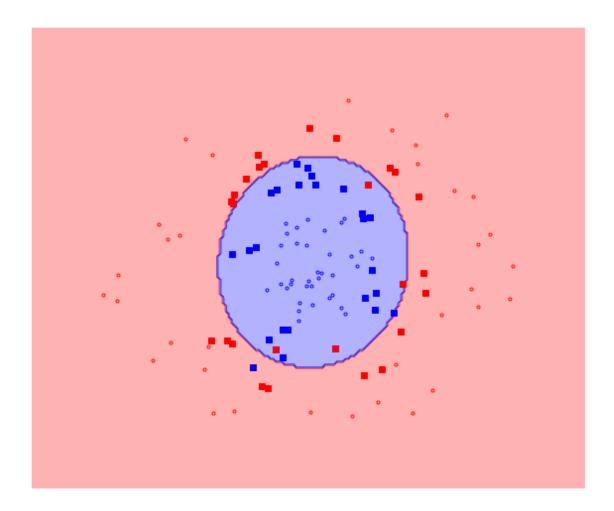
```
dictions of the classifier on the augmented training set
# initialize the different categories of data points
true pos x = []
true_pos_y = []
true_neg_x = []
true_neg_y = []
false_pos_x = []
false_pos_y = []
false_neg_x = []
false_neg_y = []
svm pos x = []
svm_pos_y = []
svm neg x = []
svm neg y = []
svm_indices = clf.support # indices for the support vectors
# correctly place each data point into one of the 6 categories:
# true positive, true negative, false positive, false negatives, svm pos
itive, svm negative
size of T = len(augmented training set)
for sample index in range(size of T):
    sample = augmented training set[sample index]
                                                                     # sa
mple coordinate
    true label = T['y'][sample index]
                                                      # true label
    predicted label = predicted labels[sample index] # predicted label
    # check svm indices
    if sample index in svm indices:
        # current sample is an svm if this goes through
        if true label == 1:
                                          # put the sample into the posit
ive svm list
            svm pos x.append(sample[0])
            svm pos y.append(sample[1])
        if true label != 1:
                                         # put the sample into the negat
ive svm list
            svm neg x.append(sample[0])
            svm neg y.append(sample[1])
    # true positive
    if true label == 1 and predicted label == 1:
        true_pos_x.append(sample[0])
        true_pos_y.append(sample[1])
    # true negative
    if true label != 1 and predicted label != 1:
        true neg x.append(sample[0])
        true neg y.append(sample[1])
    # false positive
    if true label != 1 and predicted label == 1:
        false pos x.append(sample[0])
```

```
false_pos_y.append(sample[1])
    # false negatives
    if true label == 1 and predicted label != 1:
        false neg x.append(sample[0])
        false_neg_y.append(sample[1])
plt.figure(figsize=(10,10))
# plot the data points from the 6 categories
plt.plot(true_pos_x, true_pos_y, '.b', fillstyle ='none')
plt.plot(true_neg_x, true_neg_y, '.r', fillstyle ='none')
plt.plot(false_pos_x, false_pos_y, 'xr', fillstyle ='none')
plt.plot(false_neg_x, false_neg_y, 'xb', fillstyle = 'none')
plt.plot(svm_pos_x, svm_pos_y, 'sb', fillstyle = None)
plt.plot(svm_neg_x, svm_neg_y, 'sr', fillstyle = None)
# specifiy some plotting things
plt.axis('equal')
plt.axis('off')
# plot the decision boundary
h = plt.contourf(X1,X2,predicted, alpha=0.3, cmap = colors.ListedColorma
p(['r','b'])) #check colors
```

/Users/joycechoi/anaconda3/lib/python3.6/site-packages/ipykernel_launch er.py:20: DeprecationWarning: object of type <class 'numpy.float64'> ca nnot be safely interpreted as an integer.

/Users/joycechoi/anaconda3/lib/python3.6/site-packages/ipykernel_launch er.py:21: DeprecationWarning: object of type <class 'numpy.float64'> ca nnot be safely interpreted as an integer.

Zero-one training error rate: 4.17 %
Test error rate: 3.89 %



Problem 2.4

Explain carefully why this type of data augmentation works well for this specific data set.

When we change the dimensions of the data to 5 dimensions, the points are still plotted in the same locations but now the linear classifier has 3 more dimensions worth of information which is why it's able to more closely follow the data and make a circular curve.

In this case, we augment with values x_1^2 , x_2^2 , $(x_1)*(x_2)$, which are the transformations that give an ellipse (as $x_1^2 + x_2^2 + (x_1)*(x_2) = r$ is the equation for an ellipse). Therefore, augmenting the original points allows us to get a distribution of points that are separable in a higher dimension, which is why augmenting the data works to generate the SVM in this particular data set.

Problem 2.5

Try the experiment in Problem 2.3 with sklearn.linear_model.LogisticRegression. Use parameters C=1e5, solver='lbfgs', multi_class='multinomial', random_state=0.

```
In [7]: from sklearn.linear_model import LogisticRegression
        # train sklearn.svm.SVC with arguments kernel = 'linear', C = 1 on aug T
         x (Augmented training set T )
        # show its zero-one training and test error rates (on S) as percentages
         with two decimal places
        # finding the max and min along each dimension for plotting the contour
        x1_dimension = T['x'][:, 0] # get all the values of T['x'] along the x1
         dimension
        x^2 dimension = T['x'][:, 1] # get all the values of T['x'] along the x^2
         dimension
        \max x1 = \max(x1 \text{ dimension}) \# \max value of x1
        min_x1 = min(x1_dimension) # min value of x1
        \max x^2 = \max(x^2 \text{ dimension}) \# \max value \text{ of } x^2
        min_x2 = min(x2_dimension) # min value of x2
        # add or subtract from each max or min respectively
        \max x1 = \max x1 + 0.5 \# add 0.5 to the \max x1 value
        min_x1 = min_x1 - 0.5 \# subtract 0.5 to the min_x1 value
        max_x^2 = max_x^2 + 0.5 \# add 0.5 to the max x^2 value
        min x2 = min x2 - 0.5 # subtract 0.5 from the min x2 value
        x1_range = np.linspace(min_x1, max_x1, (max_x1 - min_x1) / 0.02) # creat
        e a list of the x1 range
        x2 range = np.linspace(min x2, max x2, (max x2 - min x2) / 0.02) # creat
        e a list of the x2 range
        X1, X2 = np.meshgrid(x1 range, x2 range)
                                                                           # creat
        e the meshgrid of X1, X2
        #flatten X1 and X2 to make it useable in np.stack
        X1 flat = np.concatenate(X1).ravel()
        X2 flat = np.concatenate(X2).ravel()
        #stack X1 and X2. Creates an array an (dim(X1) \times dim(X2), 2) array
        stack = np.stack((X1 flat, X2 flat), axis=-1)
        # augment the stack with x3, x4, and x5
        stack length = len(stack)
                                                             # stack length or nu
        mber of entries in stack
        augmented training set = np.zeros((len(T['x']), 5)) # the same length as
         T['x'] but 5-d as opposed to 2-d
        augmented test set = np.zeros((len(S['x']), 5)) # the same length as
         S['x'] but 5-d as opposed to 2-d
        augmented_T_length = len(augmented_training_set) # length of augmente
        augmented S length = len(augmented test set) # length of augmente
        augmented_stack = np.zeros((stack_length, 5))  # initialize the aug
        mented stack
        \# loop through the stack to create x3, x4, and x5 and put those values i
        nto the augmented stack
        for sample index in range(stack length):
```

```
x1 = stack[sample index][0]
    x2 = stack[sample index][1]
    x3 = x1**2
    x4 = x2**2
    x5 = x1*x2
    augmented_stack[sample_index][0] = x1
    augmented stack[sample index][1] = x2
    augmented stack[sample index][2] = x3
    augmented stack[sample index][3] = x4
    augmented stack[sample index][4] = x5
for sample_index in range(augmented_T_length):
    sample = T['x'][sample index]
    x1 = sample[0] # looks at the first coordinate x1 of the sample at s
ample index in T['x']
    x2 = sample[1] # looks at the second coordinate x2 of the sample at
 sample index in T['x']
   x3 = x1**2
    x4 = x2**2
    x5 = x1*x2
    augmented training set[sample index][0] = x1
    augmented training set[sample index][1] = x2
    augmented training set[sample index][2] = x3
    augmented training set[sample_index][3] = x4
    augmented training set[sample index][4] = x5
for sample index in range(augmented S length):
    sample = S['x'][sample index]
    x1 = sample[0] # looks at the first coordinate x1 of the sample at s
ample index in S['x']
    x2 = sample[1] # looks at the second coordinate x2 of the sample at
 sample index in S['x']
   x3 = x1**2
    x4 = x2**2
    x5 = x1*x2
    augmented test set[sample index][0] = x1
    augmented test set[sample index][1] = x2
    augmented test set[sample index][2] = x3
    augmented test set[sample index][3] = x4
    augmented test set[sample index][4] = x5
clf = LogisticRegression(C=1e5, solver='lbfgs', multi class='multinomia
1', random_state=0)
clf.fit(augmented training set, T['y'])
  # fit the classifier to the training data
zero one training rate = 1 - clf.score(augmented training set, T['y'])
  # train the classifier on augmented training set aug T x and get error
 rate
test_error_rate = 1 - clf.score(augmented_test_set, S['y'])
  # test the classifier on the augmented test set aug S x and get error
 rate
print("Zero-one training error rate: {:.2f}".format(zero one training ra
te*100), "%")
print("Test error rate: {:.2f}".format(test error rate*100), "%")
predicted = clf.predict(augmented stack).reshape(X1.shape) # train the c
```

```
lassifier on the augmented stack
predicted labels = clf.predict(augmented training set)
                                                          # get the pre
dictions of the classifier on the augmented training set
# initialize the different categories of data points
true_pos_x = []
true pos y = []
true neg x = []
true neg y = []
false_pos_x = []
false_pos_y = []
false_neg_x = []
false_neg_y = []
# correctly place each data point into one of the 4 categories:
# true positive, true negative, false positive, false negatives
size_of_T = len(augmented_training_set)
for sample index in range(size of T):
    sample = augmented_training_set[sample_index]
                                                                     # sa
mple coordinate
    true_label = T['y'][sample_index]
                                                      # true label
    predicted label = predicted labels[sample_index] # predicted label
    # true positive
    if true label == 1 and predicted label == 1:
        true_pos_x.append(sample[0])
        true pos y.append(sample[1])
    # true negative
    if true label != 1 and predicted label != 1:
        true_neg_x.append(sample[0])
        true neg y.append(sample[1])
    # false positive
    if true label != 1 and predicted label == 1:
        false pos x.append(sample[0])
        false pos y.append(sample[1])
    # false negatives
    if true label == 1 and predicted label != 1:
        false neg x.append(sample[0])
        false neg y.append(sample[1])
plt.figure(figsize=(10,10))
# plot the data points from the 6 categories
plt.plot(true_pos_x, true_pos_y, '.b', fillstyle ='none')
plt.plot(true_neg_x, true_neg_y, '.r', fillstyle ='none')
plt.plot(false_pos_x, false_pos_y, 'xr', fillstyle ='none')
plt.plot(false_neg_x, false_neg_y, 'xb', fillstyle ='none')
# specifiy some plotting things
plt.axis('equal')
plt.axis('off')
```

plot the decision boundary
h = plt.contourf(X1,X2,predicted, alpha=0.3, cmap = colors.ListedColorma
p(['r','b'])) #check colors

Zero-one training error rate: 4.17 %
Test error rate: 5.56 %

/Users/joycechoi/anaconda3/lib/python3.6/site-packages/ipykernel_launch er.py:22: DeprecationWarning: object of type <class 'numpy.float64'> ca nnot be safely interpreted as an integer.

/Users/joycechoi/anaconda3/lib/python3.6/site-packages/ipykernel_launch er.py:23: DeprecationWarning: object of type <class 'numpy.float64'> ca nnot be safely interpreted as an integer.

