# Assignment\_1\_Solution

March 6, 2019

# 1 Assignment 1

#### 1.1 Task 1

Suppose we have a two-class classification problem, where we denote the two classes with +1 and -1. Further assume that the joint distribution of x and y, p(x,y), is known and that the distributions of the two classes do not overlap, i.e.

$$\min\{p(\mathbf{x}|y=+1), p(\mathbf{x}|y=-1)\}=0.$$

Determine an optimal classification function g and compute the generalization error using the zero-one loss function.

#### 1.1.1 Solution

Since  $p(\mathbf{x}, y)$ , is known, we can also compute the marginals  $p(\mathbf{x})$  and p(y) by averaging (i.e. integrating or summing) over variables (e.g. if we are interested in  $p(\mathbf{x})$ , we need to average over y). Then, by definition of conditional probability, we can also compute  $p(\mathbf{x}|y) = \frac{p(\mathbf{x},y)}{p(y)}$  and similarly  $p(y|\mathbf{x})$ . According to the slides, we can therefore find an optimal classifier, given by any function g that satisfies

$$g(\mathbf{x}) \begin{cases} > 0 & \text{for} \quad p(y=1 \mid \mathbf{x}) > p(y=-1 \mid \mathbf{x}) \\ < 0 & \text{for} \quad p(y=-1 \mid \mathbf{x}) > p(y=1 \mid \mathbf{x}) \end{cases}$$

Moreover,by the definition of conditional probability(or also known as Bayes' Theorem), we have

$$p(y|\mathbf{x}) = \frac{p(\mathbf{x}|y)p(y)}{p(\mathbf{x})}.$$

This immediately leads to the conclusion that  $\min\{p(\mathbf{x}|y=+1), p(\mathbf{x}|y=-1)\}=0$  if and only if  $\min\{p(y=+1\mid\mathbf{x}), p(y=-1\mid\mathbf{x})\}=0$ . We also know from the slides how the minimal risk can be computed and therefore conclude:

$$R_{\min} = \int_X \min\{p(y = -1 \mid \mathbf{x}), p(y = 1 \mid \mathbf{x})\} p(\mathbf{x}) d\mathbf{x} = 0.$$

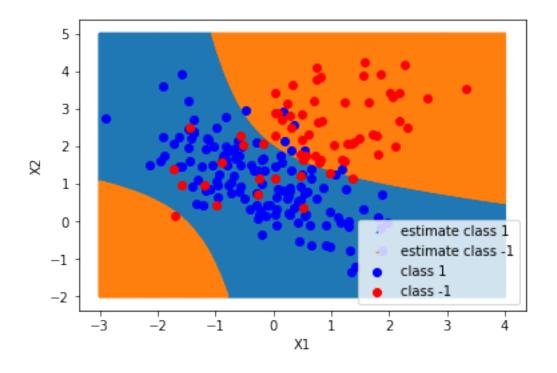
# 1.2 Task 2

Assume that the two classes in data set DataSet6 are distributed according to multivariate normal distributions. Estimate the means and covariance matrices as well as p(y=+1) and p(y=-1) from the data (you may use all 200 samples), compute an optimal classification function (see slide 9) and visualize it graphically(a two-dimensional plot suffices).

#### 1.2.1 Solution

```
In [5]: import sklearn
        import numpy as np
        import matplotlib.pyplot as plt
        #read data, split into X(features) and y(labels)
        Z = np.genfromtxt('DataSet6.csv', delimiter=',')
        X, y = Z[:,:-1], Z[:,-1]
        #further split features according to labels
        Xpos=X[y==1]
        Xneg=X[y==-1]
        #compute mean and covariance
        meanXpos=np.mean(Xpos.T,axis=1)
        meanXneg=np.mean(Xneg.T,axis=1)
        covXpos=np.cov(Xpos.T)
        covXneg=np.cov(Xneg.T)
        #compute distribution p(y=+-1)
        p_ypos=Xpos.T.shape[1]/X.shape[0]
        p_yneg=1-p_ypos
        #print corresponding values
        print("Cov of positve class=", covXpos)
        print("Cov of negative class=", covXneg)
        print("Mean of positve class=", meanXpos)
        print("Mean of negative class=", meanXneg)
        print("P(y=1)=", p_ypos)
        print("P(y=-1)=", p_yneg)
        #compute inverses of covariance matrices
        covXpos_inv = np.linalg.inv(covXpos)
        covXneg_inv = np.linalg.inv(covXneg)
        #compute parameters of discrimination function q according to slide 9
        A= covXpos_inv - covXneg_inv
        w = np.matmul(covXpos_inv, meanXpos) - np.matmul(covXneg_inv, meanXneg)
        b = (-1/2*np.matmul(np.matmul(meanXpos, covXpos_inv), meanXpos)
        + 1/2*np.matmul(np.matmul(meanXneg, covXneg_inv), meanXneg)
        - 1/2 * np.log(np.linalg.det(covXpos)) + 1/2 * np.log(np.linalg.det(covXneg))
        + np.log(p_ypos) - np.log(p_yneg) )
        #Print corresponding values
        print("Value of A=", A)
        print("Value of w=", w)
        print("Value of b=", b)
```

```
#Create grid and use it to evaluate g
        X1, X2 = np.mgrid[-3:4:500j, -2:5:500j]
        points = np.c_[X1.ravel(), X2.ravel()]
        g = -1/2*np.sum(np.dot(points, A) * points, axis=1) + np.dot(points, w) + b
        #Create Plot
        plt.scatter(X1,X2, g > 0, label="estimate class 1")
        plt.scatter(X1,X2, g < 0, label="estimate class -1")</pre>
        plt.scatter(Xpos[:,0],Xpos[:,1], color= 'blue', label="class 1")
        plt.scatter(Xneg[:,0],Xneg[:,1], color='red', label="class -1")
        plt.xlabel("X1")
        plt.ylabel("X2")
        plt.legend(loc=4)
        plt.show()
Cov of positve class= [[ 0.97776086 -0.63279722]
 [-0.63279722 0.96214327]]
Cov of negative class= [[1.18686086 0.55312556]
 [0.55312556 0.98948508]]
Mean of positve class= [-0.23072318 1.0973771]
Mean of negative class= [0.72081505 2.3428049 ]
P(y=1) = 0.6716417910447762
P(y=-1) = 0.32835820895522383
Value of A= [[0.64132121 1.80808986]
 [1.80808986 0.44294787]]
Value of w= [ 1.54525683 -1.02711802]
Value of b= 3.083468597232569
```



### 1.3 Task 3

#### 1.3.1 Part 1

Given the model class M of all exponential distributions with parameter  $\lambda$ . i.e.  $M = \{f_{\lambda}(x) \mid \lambda > 0\}$  with

$$f_{\lambda}(x) = \begin{cases} \lambda e^{-\lambda x} & x \ge 0\\ 0 & x < 0 \end{cases}$$

Derive a formula for the maximum likelihood estimator  $\lambda^*$  for the parameter  $\lambda$ .

Hint: Maximize the logarithm of the likelihood function instead of the likelihood function itself.

#### 1.3.2 Solution

Assume that  $\mathbf{x} = (x_1, ..., x_n)$ , where the  $x_i$ s are iid, according to an exponential distribution with parameter  $\lambda$ . Without restriction we can work with positive  $x_i$ 's. The liklehood can then be computed, according to the slides, as

$$\mathcal{L}(\{\mathbf{x}\};\lambda) = p(\{\mathbf{x}\};\lambda\}) = \prod_{i=1}^n p(x_i;\lambda) = \prod_{i=1}^n f_{\lambda}(x_i) = \lambda^n \prod_{i=1}^n e^{-\lambda x_i}.$$

Now we take the logarithm and obtain

$$\log(\mathcal{L}(\{\mathbf{x}\};\lambda)) = n\log(\lambda) - \lambda \sum_{i=1}^{n} x_{i}.$$

Taking the derivative wrt.  $\lambda$  yields

$$\frac{\partial \log(\mathcal{L}(\{\mathbf{x}\};\lambda))}{\partial \lambda} = \frac{n}{\lambda} - \sum_{i=1}^{n} x_i.$$

Setting the above expression to 0, finally leads us to the following result for the optimal  $\lambda^*$ :

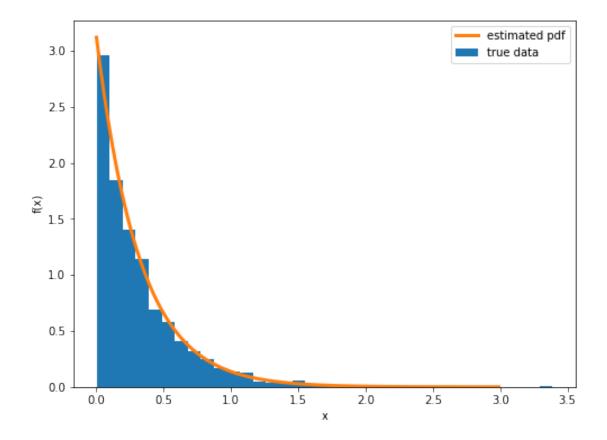
$$\lambda^* = \frac{n}{\sum_{i=1}^n x_i}.$$

## 1.3.3 Part 2

Apply the formula for  $\lambda^*$  from the previous part to the data of DataSet7. Visualize the density defined by this optimal  $\lambda^*$  and compare it to the true data distribution (e.g. by using a histogram).

#### 1.3.4 Solution

```
In [6]: # load the data
       data = np.genfromtxt('DataSet7.csv', delimiter=',', skip_header=1)
        # estimate the parameter according to the previously derived formula
        lambd_est = len(data)/np.sum(data)
        print('Estimated lambda = ',lambd est)
        #Define function that computes estimated pdf
        def f(x_i, lambd):
            if x_i < 0:
               return 0
            return lambd * np.exp(-lambd * x_i)
        #Evaluate f on points with distance 0.01 in interval [0,3], store values
       x = np.arange(0, 3.0, 0.01)
       y = [f(x_i, lambd_est) for x_i in x]
        #Make plot
       plt.figure(figsize = (8,6))
       plt.hist(data, 35, density = True, label = 'true data')
       plt.plot(x, y, label = 'estimated pdf', linewidth = 3)
       plt.xlabel('x')
       plt.ylabel('f(x)')
       plt.legend()
       plt.show()
Estimated lambda = 3.120304521828994
```



In []: