







Proble	n 8.5	5									
(a)	0.06	8									
	0.05										
	0.012										
	0.38.										
	0.08										
	0.065										
(g)	0.015	5									
J											

```
# Import Necessary Modules
import glob
import matplotlib.pyplot as plt
from skimage import io
import numpy as np
import math
import scipy.stats as stats
%matplotlib inline
#This function takes in the column vectors of guesses for H0 and H1
#data and outputs the probability of error, assuming H0 and H1 are
#equally likely.
def proberror(testquess0, testquess1):
    #Calculate sizes.
    n0test = testquess0.shape[0]
    n1test = testquess1.shape[0]
    #The number of false alarms is the total number of ones in
testquess0.
    num false alarm =
np.sum(np.not_equal(testguess0,np.zeros((n0test,1))))
    #Divide by total number of guesses to estimate the probability of
false alarm.
    P FA = num false alarm / n0test
    #The number of missed detections is the total number of zeros in
testquess1.
    num missed detection =
np.sum(np.not equal(testquess1,np.ones((n1test,1))))
    #Divide by total number of guesses to estimate the probability of
missed detection.
    P MD = num missed detection / nltest
    Pe = P_FA * 0.5 + P MD * 0.5
    return Pe
#Read in synthetic data for 8.5(a), 8.5(b), 8.5(c)
dataset0 = np.genfromtxt("syntheticH0.csv", delimiter = ",")
dataset1 = np.genfromtxt("syntheticH1.csv", delimiter = ",")
#Read in pet classification data for 8.4(h)
# dataset0, dataset1 = read cats dogs()
#Determine number of samples and dimension.
n0,d0 = dataset0.shape
n1,d1 = dataset1.shape
if (d0 == d1):
    d = d0
else:
    raise Exception("dataset0 and dataset1 do not have the same number
of columns.")
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#Split dataset into training and test data.
train0 = dataset0[0:math.floor(n0/2),:]
test0 = dataset0[math.floor(n0/2):n0,:]
train1 = dataset1[0:math.floor(n1/2),:]
test1 = dataset1[math.floor(n1/2):n1,:]
n0train = train0.shape[0]
nltrain = train1.shape[0]
n0test = test0.shape[0]
n1test = test1.shape[0]
#Estimate mean vectors and covariance matrices from training data.
mu0 = np.mean(train0, axis=0)
mu1 = np.mean(train1, axis=0)
sigma0 = np.cov(train0, rowvar=False)
sigma1 = np.cov(train1, rowvar=False)
#This function takes in a row vector currentdata as well as two mean
#vectors mu0 and m1. It outputs 1 if currentdata is closer to mu1
#than mu0, and 0 if mu0 is closer. In the case of a tie, it outputs 1.
def closest average(currentdata,mu0,mu1):
    #Calculate distances.
    distance mu0 = np.math.sqrt(np.sum((currentdata-mu0)**2))
    distance mu1 = np.math.sqrt(np.sum((currentdata-mu1)**2))
    #Decide based on smaller distance.
    if (distance mu0 < distance mu1):</pre>
        quess = 0
    elif (distance mu0 >= distance mu1):
        quess = 1
    return guess
#8.5(a) Apply Gaussian ML rule for identity covariance matrix by
plugging into the PDFs.
H0guesses idcov = np.zeros((n0test, 1))
H1guesses idcov = np.zeros((n1test,1))
for i in range(n0test):
    currentdata = test0[i,:]
    h0 = stats.multivariate_normal.pdf(currentdata, mu0,
np.identity(d0))
    h1 = stats.multivariate normal.pdf(currentdata, mul,
np.identity(d0))
    if h1>= h0:
        H0guesses\ idcov[i] = 1
for i in range(n1test):
    currentdata = test1[i,:]
    h0 = stats.multivariate normal.pdf(currentdata, mu0,
np.identity(d1))
    h1 = stats.multivariate normal.pdf(currentdata, mul,
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```
np.identity(d1))
         if h1>= h0:
                  H1guesses idcov[i] = 1
Pe idcov = proberror(H0quesses idcov, H1quesses idcov)
print("Probability of error for identity covariance matrix is " +
str(Pe idcov) + ".")
Probability of error for identity covariance matrix is 0.068.
#8.5 (b) Apply Gaussian ML rule for the same covariance matrix by
plugging into the PDFs.
HOguesses samecov = np.zeros((nOtest, 1))
H1quesses samecov = np.zeros((n1test,1))
sigmas = (1/(n0train+n1train-2))*((n0train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sigma0+(n1train-1)*sig
1)*sigma1)
for i in range(n0test):
         currentdata = test0[i,:]
         h0 = stats.multivariate normal.pdf(currentdata, mu0, sigmas)
         h1 = stats.multivariate normal.pdf(currentdata, mul, sigmas)
         if h1>=h0:
                  HOguesses samecov[i] = 1
for i in range(nltest):
         currentdata = test1[i.:]
         h0 = stats.multivariate normal.pdf(currentdata, mu0, sigmas)
         h1 = stats.multivariate normal.pdf(currentdata, mul, sigmas)
         if h1>=h0:
                  H1guesses samecov[i] = 1
Pe_samecov = proberror(H0guesses_samecov,H1guesses_samecov)
print("Probability of error for the same covariance matrices is " +
str(Pe samecov) + ".")
Probability of error for the same covariance matrices is 0.05.
#8.5 c Apply Gaussian ML rule for different covariance matrices by
plugging into the PDFs.
HOguesses diffcov = np.zeros((n0test, 1))
H1guesses diffcov = np.zeros((n1test,1))
for i in range(n0test):
         currentdata = test0[i,:]
         h0 = stats.multivariate normal.pdf(currentdata, mu0, sigma0)
         h1 = stats.multivariate_normal.pdf(currentdata, mul, sigmal)
         if h1>=h0:
                  HOguesses samecov[i] = 1
for i in range(nltest):
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currentdata = test1[i,:]
    h0 = stats.multivariate normal.pdf(currentdata, mu0, sigma0)
    h1 = stats.multivariate normal.pdf(currentdata, mul, sigmal)
    if h1>=h0:
       H1guesses diffcov[i] = 1
Pe diffcov = proberror(H0guesses diffcov, H1guesses diffcov)
print("Probability of error for different covariance matrices is " +
str(Pe diffcov) + ".")
Probability of error for different covariance matrices is 0.012.
## For problem 8.5(d), load a new data set:
# Read in breast cancer data
dataset0 = np.genfromtxt("benignfull.csv", delimiter = ",")
dataset1 = np.genfromtxt("malignantfull.csv", delimiter = ",")
# compute the statistics
n0,d0 = dataset0.shape
n1,d1 = dataset1.shape
if (d0 == d1):
    0b = b
else:
    raise Exception("dataset0 and dataset1 do not have the same number
of columns.")
#Split dataset into training and test data.
train0 = dataset0[0:math.floor(n0/2),:]
test0 = dataset0[math.floor(n0/2):n0,:]
train1 = dataset1[0:math.floor(n1/2),:]
test1 = dataset1[math.floor(n1/2):n1,:]
n0train = train0.shape[0]
nltrain = train1.shape[0]
n0test = test0.shape[0]
n1test = test1.shape[0]
#Estimate mean vectors and covariance matrices from training data.
mu0 = np.mean(train0, axis=0)
mu1 = np.mean(train1, axis=0)
sigma0 = np.cov(train0, rowvar=False)
sigma1 = np.cov(train1, rowvar=False)
#8.5(d) Apply Gaussian ML rule for identity covariance matrix by
plugging into the PDFs for the cancer data
# Note: you shoud reuse the code from 8.5(a)
H0guesses idcov = np.zeros((n0test,1))
H1guesses idcov = np.zeros((n1test,1))
for i in range(n0test):
    currentdata = test0[i,:]
```

```
h0 = stats.multivariate normal.pdf(currentdata, mu0,
np.identity(d0))
   h1 = stats.multivariate normal.pdf(currentdata, mu1,
np.identitv(d0))
   if h1>=h0:
        H0quesses idcov[i] = 1
for i in range(nltest):
    currentdata = test1[i,:]
   h0 = stats.multivariate normal.pdf(currentdata, mu0,
np.identity(d1))
   h1 = stats.multivariate normal.pdf(currentdata, mul,
np.identity(d1))
   if h1>=h0:
       H1guesses idcov[i] = 1
Pe idcov = proberror(H0guesses idcov, H1guesses idcov)
print("Probability of error for identity covariance matrix is " +
str(Pe idcov) + ".")
Probability of error for identity covariance matrix is 0.385.
#Now we start testing with the log-likelihood tests
# Problem 8.5(e) Apply Gaussian ML rule for identity covariance
matrix by using the closest average classifier.
# Recall that function was defined above.
H0guesses idcov = np.zeros((n0test, 1))
H1quesses idcov = np.zeros((n1test,1))
for i in range(n0test):
    currentdata = test0[i,:]
   H0guesses idcov[i] = closest average(currentdata, mu0, mu1)
for i in range(n1test):
    currentdata = test1[i,:]
   H1guesses idcov[i] = closest average(currentdata,mu0,mu1)
Pe idcov = proberror(H0quesses idcov,H1quesses idcov)
print("Probability of error for identity covariance matrix is " +
str(Pe idcov) + ".")
Probability of error for identity covariance matrix is
/var/folders/pf/619m vfi3w5cwc52r3vxbb 40000gn/T/
ipykernel 46189/403505849.py:7: DeprecationWarning: `np.math` is a
deprecated alias for the standard library `math` module (Deprecated
Numpy 1.25). Replace usages of `np.math` with `math`
  distance mu0 = np.math.sqrt(np.sum((currentdata-mu0)**2))
```

```
/var/folders/pf/619m yfj3w5cwc52r3yxbb 40000gn/T/ipykernel 46189/40350
5849.py:8: DeprecationWarning: `np.math` is a deprecated alias for the
standard library `math` module (Deprecated Numpy 1.25). Replace usages
of `np.math` with `math`
  distance mu1 = np.math.sqrt(np.sum((currentdata-mu1)**2))
#8.5 (f) Apply Gaussian ML rule for the same covariance matrix by
using the log-likelihood ratio
# using poled covariances
HOguesses samecov = np.zeros((nOtest, 1))
H1guesses samecov = np.zeros((n1test, 1))
sigmas = (1/(n0train+n1train-2))*((n0train-1)*sigma0+(n1train-1)*
1)*sigma1)
sigmainv = np.linalg.pinv(sigmas)
for i in range(n0test):
    currentdata = test0[i,:]
    value0 = np.matmul(np.matmul((currentdata-
mu0),sigmainv),np.transpose(currentdata-mu0))
    value1 = np.matmul(np.matmul((currentdata-
mul), sigmainv), np.transpose(currentdata-mul))
    if value1<=value0:</pre>
        HOguesses samecov[i] = 1
for i in range(n1test):
    currentdata = test1[i,:]
    value0 = np.matmul(np.matmul((currentdata-
mu0),sigmainv),np.transpose(currentdata-mu0))
    value1 = np.matmul(np.matmul((currentdata-
mul), sigmainv), np.transpose(currentdata-mul))
    if value1<=value0:</pre>
        H1guesses samecov[i] = 1
Pe samecov = proberror(H0quesses samecov, H1quesses samecov)
print("Probability of error for the same covariance matrices is " +
str(Pe samecov) + ".")
Probability of error for the same covariance matrices is 0.065.
#8.5(q) Apply Gaussian ML rule for different covariance matrices by
using the log-likelihood ratio.
HOguesses diffcov = np.zeros((n0test, 1))
Hlquesses diffcov = np.zeros((nltest,1))
sigma0inv = np.linalg.pinv(sigma0)
sigmalinv = np.linalg.pinv(sigmal)
offset = math.log(np.linalg.det(sigma0)) -
math.log(np.linalg.det(sigma1))
```

```
for i in range(n0test):
    currentdata = test0[i,:]
    value0 = np.matmul(np.matmul((currentdata-
mu0),sigma0inv),np.transpose(currentdata-mu0))
    value1 = np.matmul(np.matmul((currentdata-
mul), sigmalinv), np.transpose(currentdata-mul))
    if value1<=(value0+offset):</pre>
        H0guesses diffcov[i] = 1
for i in range(n1test):
    currentdata = test1[i,:]
    value0 = np.matmul(np.matmul((currentdata-
mu0),sigma0inv),np.transpose(currentdata-mu0))
    value1 = np.matmul(np.matmul((currentdata-
mul),sigmalinv),np.transpose(currentdata-mul))
    if value1<=(value0+offset):</pre>
        Hlguesses diffcov[i] = 1
Pe diffcov = proberror(H0guesses diffcov,H1guesses diffcov)
print("Probability of error for different covariance matrices is " +
str(Pe diffcov) + ".")
Probability of error for different covariance matrices is 0.015.
```