

HomeWork 2#

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Q1:

(a)

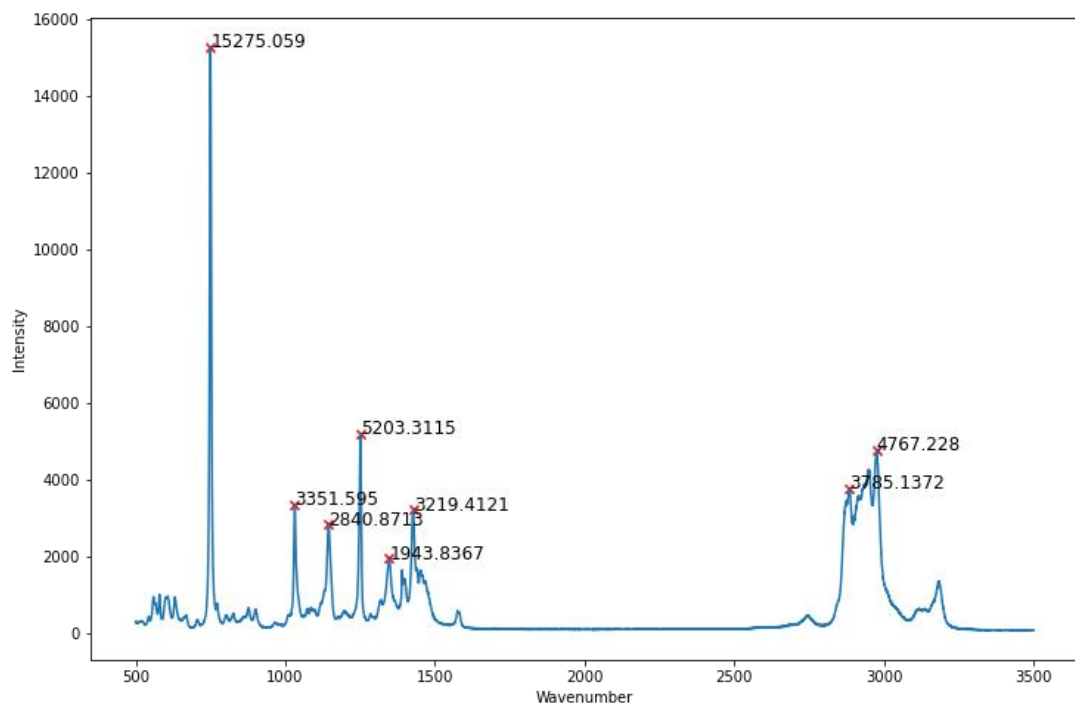
Wavenumber estimates for the eight largest spectral peak:

(750.42657, 1250.886, 2975.9155, 2884.5649, 1031.9596, 1427.287, 1145.3602, 1346.5677)

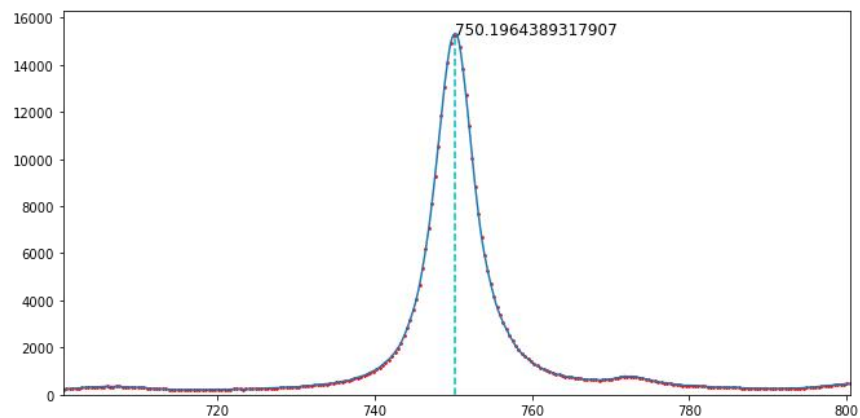
Peaks' definition:

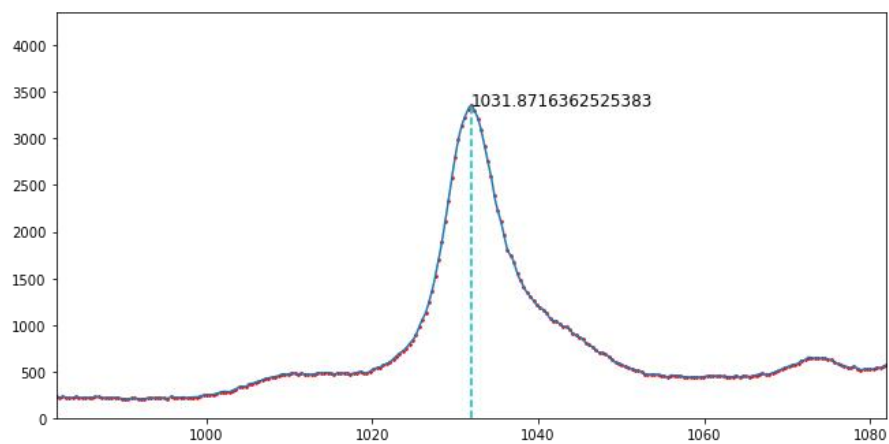
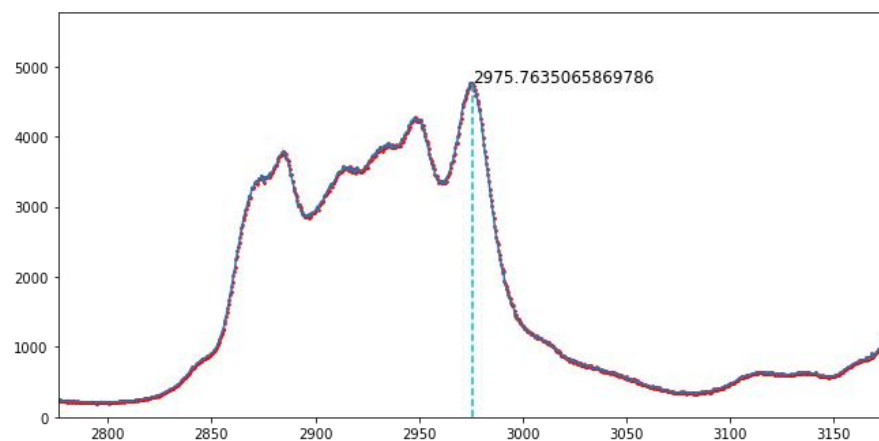
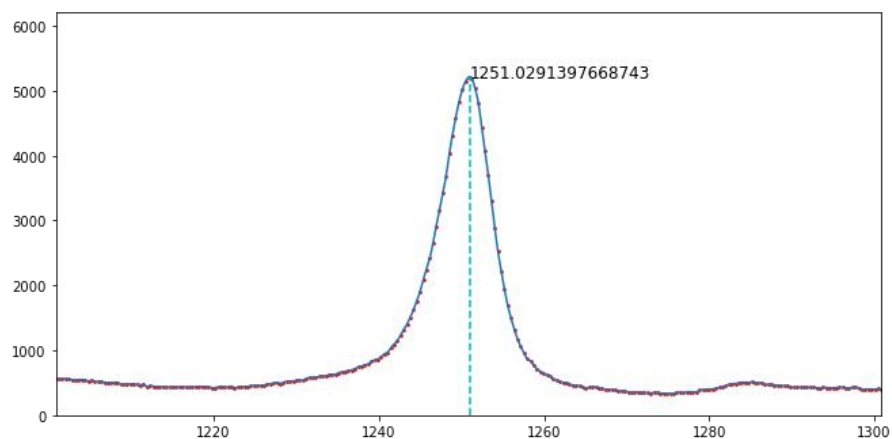
Peaks' intensity height > 1000, distance between 2 peaks more than 200 sample

(b)



(c)

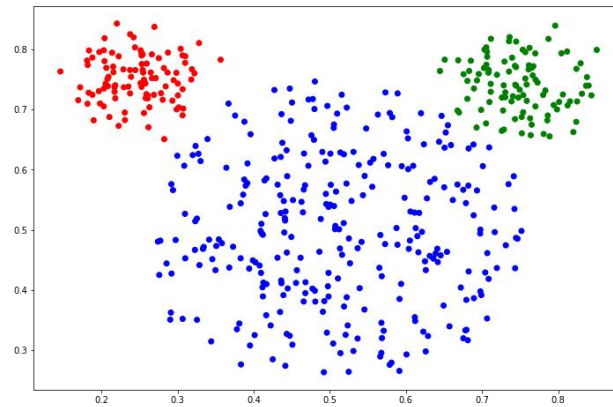




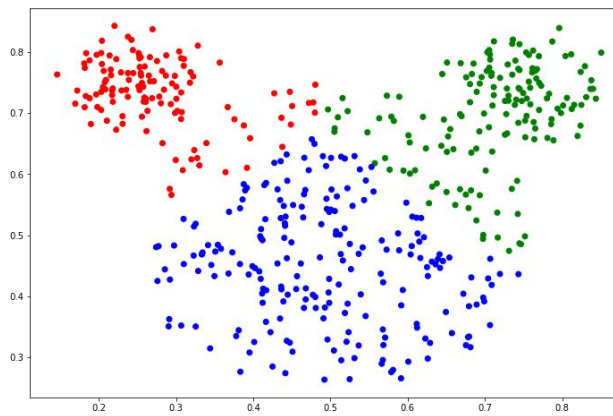
Q2

(a)

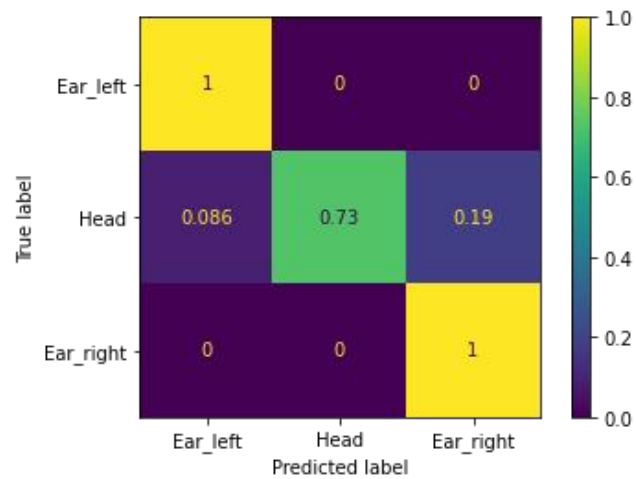
True



Predict:

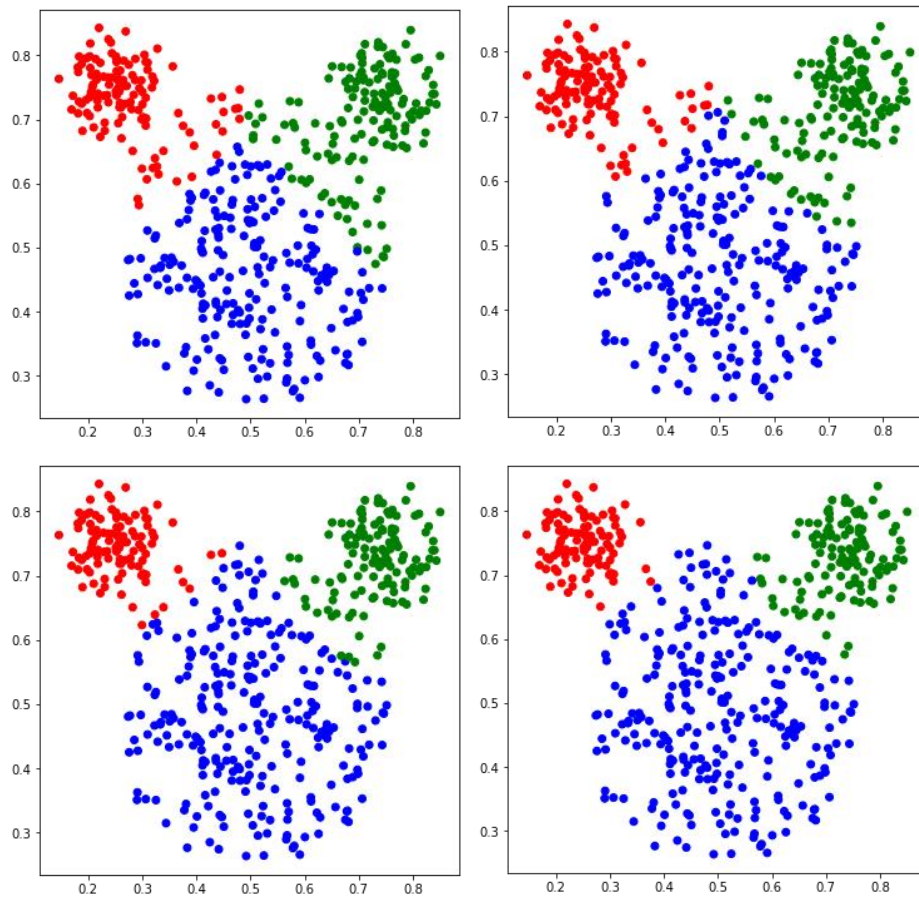


Confusion matrix

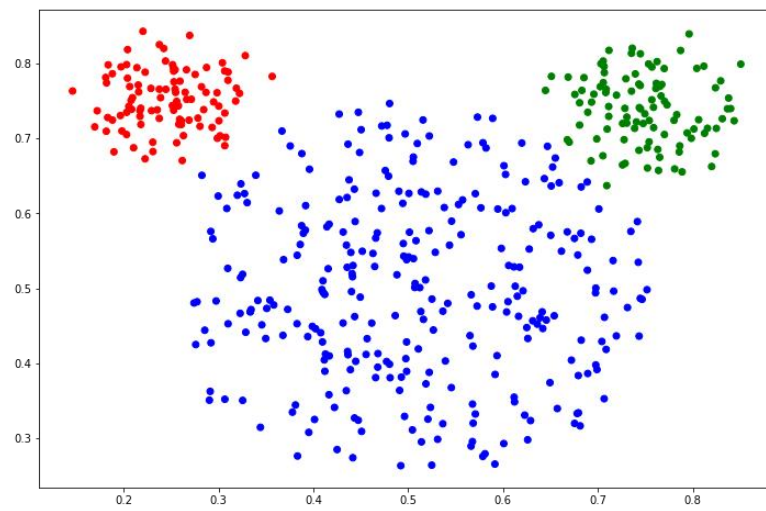


(b)

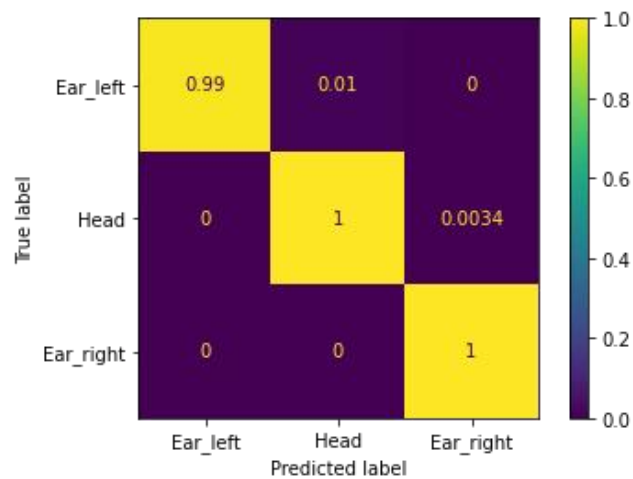
Firs 4 iterations:



Predict after 28 iterations:



Confusion matrix:



Clustering using GMM model behaves much better than K-means method.

It has more precision on bordering area while K-means only has 73% accuracy on head area.

After 28 iterations, GMM can have pretty good accuracy.

Code

Q1:

```
import matplotlib.pyplot as plt
```

```
import scipy.signal
```

```
import numpy as np
```

```
from scipy import interpolate as it
```

```
raman = "raman.txt"
```

```
wavenum, intensity = [], []
```

```
with open(raman, 'r') as f:
```

```
    lines = f.readlines()
```

```
    for data in lines:
```

```
        value = [float(s) for s in data.split()]
```

```
        wavenum.append(value[0])
```

```
        intensity.append(value[1])
```

```

peaks, _ = scipy.signal.find_peaks(intensity, height = 1000, distance = 200)
peaks_wav = [wavenum[i] for i in peaks]
peaks_intensity = [intensity[i] for i in peaks]
Z = zip(peaks_wav, peaks_intensity)
Z = sorted(Z, key = lambda x: x[1], reverse = True)
peaks_wav, peaks_intensity = zip(*Z)
# Peak definition:
#     peaks' intensity height > 1000, distance between 2 peaks more than 200 samples
print(peaks_wav[0:8])
print(peaks_intensity[0:8])
fig, ax = plt.subplots(figsize=(12,8))
plt.plot(wavenum, intensity)
plt.scatter(peaks_wav[0:8], peaks_intensity[0:8], marker = "x", color = "red")
for i in range(8):
    ax.text(peaks_wav[i], peaks_intensity[i], peaks_intensity[i], size=12)
plt.xlabel("Wavenumber")
plt.ylabel("Intensity")
plt.show()

tck = it.splrep(wavenum, intensity, s=0)
xfit = np.arange(500, 3500, np.pi/50)
yfit = it.splev(xfit, tck, der=0)

deriv = it.splev(xfit, tck, der = 1)

zero_cross = []
for i in range(len(deriv) - 1):
    if deriv[i] * deriv[i+1] < 0 and yfit[i] > 1000:
        if xfit[i] > 740 and xfit[i] < 760:
            zero_cross.append(i)
        elif xfit[i] > 1240 and xfit[i] < 1260:
            zero_cross.append(i)
        elif xfit[i] > 2950 and xfit[i] < 3000 and yfit[i] > 4600:
            zero_cross.append(i)
        elif xfit[i] > 1020 and xfit[i] < 1040:
            zero_cross.append(i)

fig = plt.figure(figsize = (24,12))
ax1 = fig.add_subplot(221)
ax1.scatter(wavenum, intensity, s = 3,color = 'red')
ax1.plot(xfit, yfit)
ax1.vlines(xfit[zero_cross[0]], 0, peaks_intensity[0], colors = "c", linestyle = "dashed")
ax1.text(xfit[zero_cross[0]], yfit[zero_cross[0]], xfit[zero_cross[0]], size=12)

```

```

ax1.set_xlim(peaks_wav[0] - 50, peaks_wav[0] + 50)
ax1.set_ylim(0, peaks_intensity[0] + 1000)

fig = plt.figure(figsize = (24,12))
ax2 = fig.add_subplot(222)
ax2.scatter(wavenum, intensity, s = 3,color = 'red')
ax2.plot(xfit, yfit)
ax2.vlines(xfit[zero_cross[2]], 0, peaks_intensity[1], colors = "c", linestyle = "dashed")
ax2.text(xfit[zero_cross[2]], yfit[zero_cross[2]], xfit[zero_cross[2]], size=12)
ax2.set_xlim(peaks_wav[1] - 50, peaks_wav[1] + 50)
ax2.set_ylim(0, peaks_intensity[1] + 1000)

fig = plt.figure(figsize = (24,12))
ax3 = fig.add_subplot(223)
ax3.scatter(wavenum, intensity, s = 3,color = 'red')
ax3.plot(xfit, yfit)
ax3.vlines(xfit[zero_cross[5]], 0, peaks_intensity[2], colors = "c", linestyle = "dashed")
ax3.text(xfit[zero_cross[5]], yfit[zero_cross[5]], xfit[zero_cross[5]], size=12)
ax3.set_xlim(peaks_wav[2] - 200, peaks_wav[2] + 200)
ax3.set_ylim(0, peaks_intensity[2] + 1000)

fig = plt.figure(figsize = (24,12))
ax4 = fig.add_subplot(224)
ax4.scatter(wavenum, intensity, s = 3,color = 'red')
ax4.plot(xfit, yfit)
ax4.vlines(xfit[zero_cross[1]], 0, peaks_intensity[4], colors = "c", linestyle = "dashed")
ax4.text(xfit[zero_cross[1]], yfit[zero_cross[1]], xfit[zero_cross[1]], size=12)
ax4.set_xlim(peaks_wav[4] - 50, peaks_wav[4] + 50)
ax4.set_ylim(0, peaks_intensity[4] + 1000)

```

Q2:

```

import numpy as np
import matplotlib.pyplot as plt
import re
import math
from scipy.cluster.vq import kmeans, vq, whiten
from sklearn import metrics
from scipy.stats import multivariate_normal as mn
cluster = "cluster.txt"

```

#Read data

```

x, y, pairs, cls = [], [], [], []
with open(cluster, 'r') as f:

```

```

lines = f.readlines()
for line in lines:
    if re.match('[1-9]\d*.\d*|0.\d*[1-9]\d*[1-9]\d*.\d*|0.\d*[1-9]\d*[A-Z]\w+', line, flags
= 0):
        data = line.split()
        x.append(float(data[0]))
        y.append(float(data[1]))
        pairs.append([float(data[0]), float(data[1])])
        if data[2] == 'Head':
            cls.append(1)
        elif data[2] == "Ear_left":
            cls.append(0)
        elif data[2] == "Ear_right":
            cls.append(2)

```

#Q2(a) predict

```
centroids,_ = kmeans(pairs,3)
```

```
centx = [centroids[i][0] for i in range(3)]
```

```
Z = zip(centroids, centx)
```

```
Z = sorted(Z, key = lambda x: x[1])
```

```
centroids, centx = zip(*Z)
```

```
km_res,_ = vq(pairs,centroids)
```

```
colors = ['red', 'blue', 'green']
```

```
col = []
```

```
for i in range(len(km_res)):
```

```
    col.append(colors[km_res[i]])
```

```
fig, ax = plt.subplots(figsize=(12,8))
```

```
plt.scatter(x, y, c = col)
```

```
plt.show()
```

#Q2(a) true

```
col2 = []
```

```
for i in range(len(cls)):
```

```
    if(cls[i] == 1):
```

```
        col2.append('blue')
```

```
    elif(cls[i] == 0):
```

```
        col2.append('red')
```

```
    elif(cls[i] == 2):
```

```
        col2.append('green')
```

```
fig, ax = plt.subplots(figsize=(12,8))
```

```
plt.scatter(x, y, c = col2)
```

```
plt.show()
```



```
#Confusion matrix
metrics.ConfusionMatrixDisplay.from_predictions(cls, km_res, normalize = 'true', display_labels =
['Ear_left', 'Head', 'Ear_right'])
plt.show()
```

#M-step

```
def update_mu(pairs, gamma, n_sample, n_class):
    mu = np.zeros((n_class,2))
    pairs = np.array(pairs)
    for i in range(n_class):
        top = 0
        bot = 0
        for j in range(n_sample):
            top += gamma[j][i] * pairs[j]
            bot += gamma[j][i]
        mu[i] = top / bot
    return mu
```

```
def update_covar(pairs, gamma, mu, n_sample, n_class):
    covar = np.zeros((n_class,2))
    pairs = np.array(pairs)
    for i in range(n_class):
        top = 0
        bot = 0
        for j in range(n_sample):
            top += gamma[j][i] * ((pairs[i] - mu[i])**2)
            bot += gamma[j][i]
        covar[i] = top / bot
    return covar
```

```
def update_covar1(pairs, gamma, mu, n_sample, n_class):
    covar = np.zeros((n_class,2))
    for i in range(n_class):
        covar[i] = np.average((pairs - mu[i])**2, axis = 0, weights = gamma[:,i])
    return covar
```

```
def update_w(gamma):
    w = gamma.sum(axis = 0) / gamma.sum()
    return w
```

#E-step

```
def gaussian_pdf(pairs, mu, covar, num, cls, dim):
    f = np.exp(-(1/2) * np.transpose(pairs[num] - mu[cls]) * np.linalg.inv(cov) * (pairs[num] -
```

```
mu[cls])) / (np.sqrt(np.linalg.det(covar)((2*math.pi)**dim)))
    return f
```

```
def update_gamma(pairs, mu, covar, w, n_sample, n_class):
    pdf = np.zeros((n_sample, n_class))
    for i in range(n_class):
        pdf[:,i] = w[i] * mn.pdf(pairs, mean = mu[i], cov = np.diag(covar[i]))
        gamma = pdf / pdf.sum(axis = 1).reshape(-1, 1)
    return gamma
```

```
def neg_log_like(pairs, mu, covar, w, n_sample, n_class):
    pdf = np.zeros((n_sample, n_class))
    for i in range(n_class):
        pdf[:,i] = w[i] * mn.pdf(pairs, mean = mu[i], cov = np.diag(covar[i]))
    l = np.log(pdf.sum(axis = 1))
    return np.mean(l)
```

#Initialize

```
gamma = []
w = np.zeros(3)
for i in range(len(km_res)):
    if(km_res[i] == 0):
        temp = [1, 0, 0]
        w[0] += 1
        gamma.append(temp)
    if(km_res[i] == 1):
        temp = [0, 1, 0]
        w[1] += 1
        gamma.append(temp)
    if(km_res[i] == 2):
        temp = [0, 0, 1]
        w[2] += 1
        gamma.append(temp)
w = w / 490
gamma = np.array(gamma)
mu = update_mu(pairs, gamma, 490, 3)
covar = update_covar1(pairs, gamma, mu, 490, 3)
```

#First 4 iterations:

```
gmm_res = []
for i in range(490):
    gmm_res.append(np.argmax(gamma[i]))
col2 = []
for i in range(len(gmm_res)):
```

```

        col2.append(colors[gmm_res[i]])
fig, ax = plt.subplots(figsize = (6, 6))
plt.scatter(x, y, c = col2)
plt.show()
for n in range(3):
    gamma = update_gamma(pairs, mu, covar, w, 490, 3)
    w = update_w(gamma)
    mu = update_mu(pairs, gamma, 490, 3)
    covar = update_covar1(pairs, gamma, mu, 490, 3)
    gmm_res = []
    for i in range(490):
        gmm_res.append(np.argmax(gamma[i]))
    col2 = []
    for i in range(len(gmm_res)):
        col2.append(colors[gmm_res[i]])
    fig, ax = plt.subplots(figsize = (6, 6))
    plt.scatter(x, y, c = col2)
    plt.show()

```

#Iteration

```

prev_l = 100
count = 0
while True:
    count += 1
    gamma = update_gamma(pairs, mu, covar, w, 490, 3)
    w = update_w(gamma)
    mu = update_mu(pairs, gamma, 490, 3)
    covar = update_covar1(pairs, gamma, mu, 490, 3)
    l = neg_log_like(pairs, mu, covar, w, 490, 3)
    if(abs(prev_l - l) < 10**-15):
        print("Iteration : ", count)
        break
    else:
        prev_l = l

```

#Plot

```

gmm_res = []
for i in range(490):
    gmm_res.append(np.argmax(gamma[i]))
col2 = []
for i in range(len(gmm_res)):
    col2.append(colors[gmm_res[i]])
fig, ax = plt.subplots(figsize=(12,8))
plt.scatter(x, y, c = col2)

```

```
plt.show()
```

```
#Confusion matrix
```

```
metrics.ConfusionMatrixDisplay.from_predictions(cls, gmm_res, normalize = 'true', display_labels  
= ['Ear_left', 'Head', 'Ear_right'])  
plt.show()
```