EXERCISE 6

1. Mixture of Gaussians

Exercise solved by Benjamin Ostendorf and Wajid Ghafoor

(1) Define the parameters of 3 Gaussian distribution functions in two dimensions.

```
In [51]: # imports needed for Exercise 6
import numpy as np
import math as m
import random
import matplotlib.pyplot as plt
from matplotlib.patches import Ellipse
In [58]: # Not Important anymore, because a more sophisticted way is implemented!
```

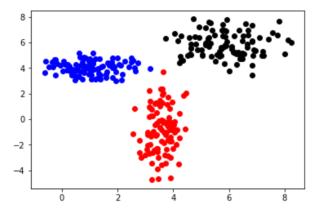
```
In [58]: # Not Important anymore, because a more sophisticted way is implemented!
def bivariate_gauss(x, mu, covarianzmatrix):
    a = (1/(m.sqrt((2* m.pi)**2)) * np.linalg.det(covarianzmatrix))
    b = (m.e ** (np.matmul(np.matmul(-0.5*np.transpose(x-mu), np.linalg.inv(covarianzmatrix)), x-mu)))
    return a * b

#bivariate_gauss(np.array([[100],[100]]), np.array([[2],[2]]), np.array([[1,0],[0,1]]))
```

```
In [5]: #create three clusters of points drawn from bivariate gaussians
gauss_1 = np.random.multivariate_normal([6,6],[[1,0],[0,1]],100)
gauss_2 = np.random.multivariate_normal([3.5,-1],[[0.1,0],[0,3]],100)
gauss_3 = np.random.multivariate_normal([1,4],[[0.9,0],[0,0.2]],100)
#concatenate the lists of points to one
gauss_com = np.concatenate((gauss_1,gauss_2, gauss_3),axis=0)
```

```
In [6]: #spliting the points into x and y components
    gauss_1_x = gauss_1[:,0]
    gauss_1_y = gauss_2[:,0]
    gauss_2_x = gauss_2[:,1]
    gauss_3_x = gauss_3[:,0]
    gauss_3_y = gauss_3[:,1]

#plotting each cluster with different color
    fig = plt.figure()
    ax = fig.add_subplot(1, 1, 1)
    ax.scatter(gauss_1_x, gauss_1_y, c="black")
    ax.scatter(gauss_2_x, gauss_2_y, c="red")
    ax.scatter(gauss_3_x, gauss_3_y, c="blue")
    plt.show()
```



```
In [7]:
        import numpy as np
        import matplotlib.pyplot as plt
        from matplotlib import cm
        from mpl toolkits.mplot3d import Axes3D
        # Our 2-dimensional distribution will be over variables X and Y
        N = 100
        X = gauss_1[:,0]
Y = gauss_1[:,1]
        X, Y = np.meshgrid(X, Y)
        # Mean vector and covariance matrix
        mu = np.array([3., 4.])
        Sigma = np.array([[ 1. , 0.], [0., 1.]])
        # Pack X and Y into a single 3-dimensional array
        pos = np.empty(X.shape + (2,))
        pos[:, :, 0] = X
        pos[:, :, 1] = Y
        def multivariate gaussian(pos, mu, Sigma):
             """Return the multivariate Gaussian distribution on array pos."""
            n = mu.shape[0]
            Sigma_det = np.linalg.det(Sigma)
            Sigma_inv = np.linalg.inv(Sigma)
            N = np.sqrt((2*np.pi)**n * Sigma_det)
            # This einsum call calculates (x-mu)T.Sigma-1.(x-mu) in a vectorized
            # way across all the input variables.
            fac = np.einsum('...k,kl,...l->...', pos-mu, Sigma_inv, pos-mu)
             return np.exp(-fac / 2) / N
        # The distribution on the variables X, Y packed into pos.
        Z = multivariate gaussian(pos, mu, Sigma)
In [8]: # Calculate the probability of a point with the following
        multivariate gaussian(np.array([3.5,-1]), np.array([3.5,-1]), np.array([[0.1]])
        ,0],[0,3]]))
Out[8]: 0.2905758415662736
In [9]: | #mue, covariance, pi
        #creates a intial distribution to use for expectation maximization
        def initial():
             return [np.random.rand(2), np.array([[1,0],[0,1]]), np.random.rand(1)]
```

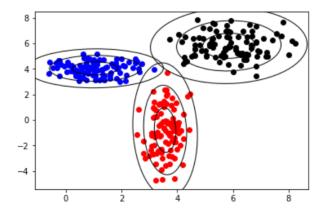
```
In [10]: #dist 1 to 3 are bivariate gaussians
         #X is a list of two dimensional points, which the gaussians should be fit to
         #returns a list of lists containing the likelyhood values of each x towards
         each gaussian
         def e_step(dist_1, dist_2, dist_3, X):
              Xs = [[dist_1[2] * multivariate_gaussian(x, dist_1[0], dist_1[1]),
                     dist_2[2] * multivariate_gaussian(x, dist_2[0], dist_2[1]),
dist_3[2] * multivariate_gaussian(x, dist_3[0], dist_3[1])] for x
         in X1
              return [[xs[0]/sum(xs), xs[1]/sum(xs), xs[2]/sum(xs)] for xs in Xs]
         #Xs is an numpy array of lists xs
         \#xs = [dist_1_l, dist_2_l, dist_3_l] is a list of 3 elements, each of which
          correspond to the likelyhood value for the corresponding point
             to be drawn from the distribution
         def m step(dist_1, dist_2, dist_3, Xs, X):
              #for each gaussian parameters have to be adjusted
              #m_c is a weight for a gaussian c
              m_c1 = sum([xs[0] for xs in Xs])
             m_c^2 = sum([xs[1] \text{ for } xs \text{ in } Xs])
             m_c3 = sum([xs[2] for xs in Xs])
             #m equals the number of trainingsdata
             m = len(Xs)
             #pi c is the normalized weight of a gaussian c
              pi c1 = m c1/m
              pi c2 = m c2/m
              pi_c3 = m_c3/m
              zipped_points = list(zip(Xs, X))
              #mue c is the adjusted mean vector for a gaussian c
              \#xsx = ([list of likelyhood values], point x)
              mue_c1 = (1/float(m_c1)*sum([xsx[0][0]*xsx[1] for xsx in zipped_points])
          )[np.newaxis]
              mue c2 = (1/float(m c2)*sum([xsx[0][1]*xsx[1] for xsx in zipped points])
          )[np.newaxis]
              mue_c3 = (1/float(m_c3)*sum([xsx[0][2]*xsx[1] for xsx in zipped_points])
          )[np.newaxis]
              #cov c is the adjusted covariance matrix of a gaussian
              \#xsx[0] is the likelyhood list for one point x
              \#xsx[1] is the point x
              #xsx[1][np.newaxis] makes the array two dimensional, for that it has the
          same shape as a mue vector
              #the np.dot(...) part calulates a covariance matrices, which in the end
         will be summed up
              cov_c1 = 1/float(m_c1)*sum([np.dot(np.transpose(xsx[0][0]*
                                                                (xsx[1][np.newaxis]-mue
         c1)),
                                                  xsx[1][np.newaxis]-mue_c1) for xsx in
          zipped points])
              cov_c2 = 1/float(m_c2)*sum([np.dot(np.transpose(xsx[0][1]*
                                                                (xsx[1][np.newaxis]-mue_
         c2)),
                                                  xsx[1][np.newaxis]-mue_c2) for xsx in
          zipped_points])
              cov c3 = 1/float(m c3)*sum([np.dot(np.transpose(xsx[0][2]*
                                                                (xsx[1][np.newaxis]-mue_
         c3)),
                                                  xsx[1][np.newaxis]-mue_c3) for xsx in
          zipped_points])
              return ([mue c1. cov c1. pi c1].[mue c2. cov c2. pi c2]. [mue c3. cov c3
```

```
In [11]: def expectation_maximization(dist_1, dist_2, dist_3, X):
    for i in range(500):
        r_ic = e_step(dist_1, dist_2, dist_3, X)
        m_step_results = m_step(dist_1, dist_2, dist_3, r_ic, X)
        dist_1 = m_step_results[0]
        dist_2 = m_step_results[1]
        dist_3 = m_step_results[2]
        return (dist_1, dist_2, dist_3)
In [19]: em = expectation_maximization(initial(), initial(), initial(), gauss_com)

In [23]: em
Out[23]: ([array([[5.85541012, 5.81626369]]), array([[0.98807032, 0.00959686], [0.00959686, 0.86785281]]), array([[0.33345344])], [array([[1.04914964, 4.04710132]]), array([[0.67043981, -0.01055142], [-0.01055142, 0.25988909]]), array([[0.3330946])], [array([[3.56756962, -0.7832351]]), array([[0.14975247, 0.06763706], [0.06763706, 3.07501472]]), array([[0.33345196])])
```

The Plot for the EM Algorithmn!

```
In [57]: | fig = plt.figure()
          ax = fig.add_subplot(1, 1, 1)
          ax.scatter(gauss_1_x, gauss_1_y, c="black")
          ax.scatter(gauss_2_x, gauss_2_y, c="red")
          ax.scatter(gauss_3_x, gauss_3_y, c="blue")
#plt.plot(centroids[:,0], centroids[:,1], 'sg', markersize=15)
          cov = em[0][1]
          lambda_, v = np.linalg.eig(cov)
lambda_ = np.sqrt(lambda_)
          for j in xrange(1, 4):
              ell = Ellipse(xy=(em[0][0][0][0], em[0][0][0][1]),
                              width=lambda_[0]*j*2, height=lambda_[1]*j*2,
                              angle=np.rad2deg(np.arccos(em[0][2][0])))
              ell.fill=False
              ax.add artist(ell)
          cov = em[1][1]
          lambda_, v = np.linalg.eig(cov)
          lambda_ = np.sqrt(lambda_)
          for j in xrange(1, 4):
              ell = Ellipse(xy=(em[1][0][0][0], em[1][0][0][1]),
                              width=lambda_[0]*j*2, height=lambda_[1]*j*2,
                              angle=np.arccos(em[1][2][0]))
              ell.fill=False
              ax.add_artist(ell)
          cov = em[2][1]
          lambda_, v = np.linalg.eig(cov)
          lambda_ = np.sqrt(lambda_)
          for j in xrange(1, 4):
              ell = Ellipse(xy=(em[2][0][0][0], em[2][0][0][1]),
                              width=lambda_[0]*j*2, height=lambda_[1]*j*2,
                              angle=np.arccos(em[2][2][0]))
              ell.fill=False
              ax.add artist(ell)
          plt.show()
```



```
In [15]: def assign and adjust(centroits, X):
             #calculate the euclidean distance for one point with each centroit and s
         elect the argmin over the distance
             #to determine the membership. The order is identical to the order of poi
         nts in X
             membership = [np.argmin(list(map(lambda centroit: np.linalg.norm(centroi
         t-x), centroits))) for x in X]
             #base values to calculate the new centroit point by averaging the assign
         ed points values
             centroit mean = [[np.array([0,0]), 1] for centroit in centroits]
             for x_index in range(len(X)):
                 member_index = membership[x_index]
                 #update the running sum and quotient
                 centroit mean[member index][0] = centroit mean[member index][0] + X[
         x index]
                 centroit mean[member index][1] += 1
             #fix the adjustment of the centroits to the mean of their assigned point
         S
             centroits = np.array(list(map(lambda centroit: centroit[0]/centroit[1],
         centroit mean)))
             return (centroits, membership)
         def k means(k, X, iterations=None):
             #our initial centroits get choosen in respect to the points we want to c
         luster
             #therefore our centroits get drawn from a gaussian with mean of the mean
         over all points
             #and the identity matrix as covariance matrix
             point mean = sum(X)/len(X)
             centroits = np.random.multivariate normal(point mean,[[1,0],[0,1]],k)
             membership = []
             #the iteration can be removed, but for this exercise issue we use it
             iterations = 100
             #Either a certain number of iterations or a convergence criterium can be
         used
             if iterations:
                 for i in range(iterations):
                     centroits, membership = assign and adjust(centroits, X)
             else:
                 while(True):
                     centroit, membership_new = assign_and_adjust(centroits, X)
                     if membership == membership new:
                         break
                     else:
                         membership = membership_new
             return (centroits, membership)
```

```
In [16]: centroids, assignments = k_means(3, gauss_com)
```

The Plot for the K-means Algorithmn!

```
In [17]: fig = plt.figure()
    ax = fig.add_subplot(1, 1, 1)
    ax.scatter(gauss_1_x, gauss_1_y, c="black")
    ax.scatter(gauss_2_x, gauss_2_y, c="red")
    ax.scatter(gauss_3_x, gauss_3_y, c="blue")
    plt.plot(centroids[:,0],centroids[:,1],'sg',markersize=15)
    plt.show()
```

