Machine Learning I - Sheet 1

20.01.2020

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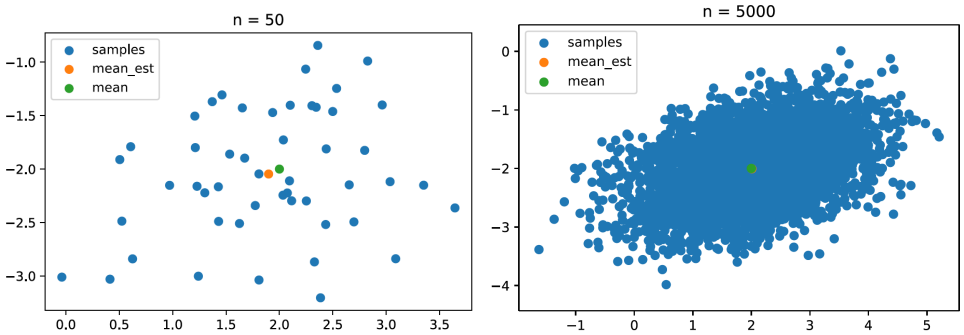
1 Maximum Likelihood

The Maximum Likelihood (ML) estimates for mean and covariance is found for different data sets. In the second part of this task, ML estimates are used to perform a simple classification.

1.1 ML Estimates

The Gaussian distribution with mean $\mu = [2, -2]^T$ and covariance $c = \begin{pmatrix} 0.9 & 0.2 \\ 0.2 & 0.3 \end{pmatrix}$ is used to generate two datasets with 50 and 5000 samples, respectively. For each data set the ML estimates for mean and covariance are calculated. The following plots illustrate the difference in the quality of the estimates for the mean:



The difference in quality can also be observed in the l2-distance to the exact mean of the distribution:

$$\mu_{50} = [1.899, -2.046]^T \quad \mu_{5000} = [2.008, -2.005]^T$$

	50	5000
l2-distance	0.1112	0.0092

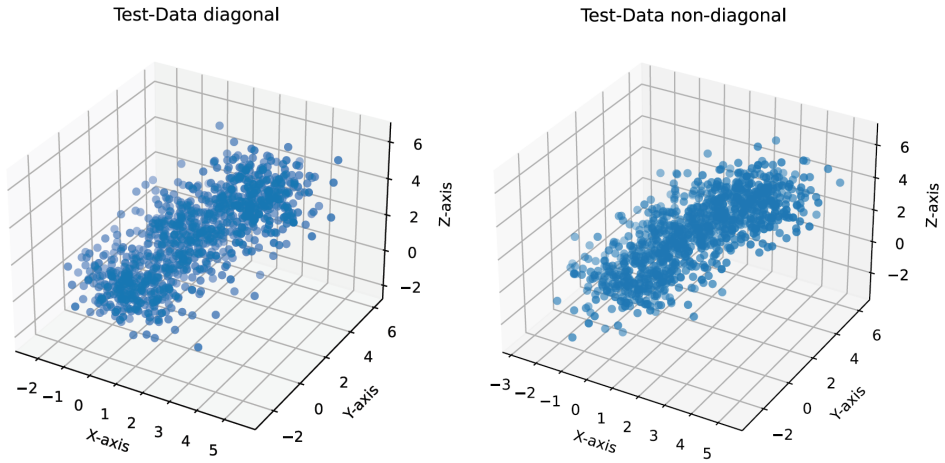
While the estimates of the mean improve by two orders of magnitudes with respect to the l2-distance, the estimate of the covariance matrix improves by one order of magnitude with respect to the frobenius-distance, when increasing the number of samples by two orders of magnitude from 50 to 5000:

$$c_{50} = \begin{pmatrix} 0.6539 & 0.1084 \\ 0.1084 & 0.3500 \end{pmatrix} \quad c_{5000} = \begin{pmatrix} 0.8855 & 0.1972 \\ 0.1972 & 0.2946 \end{pmatrix}$$

	50	5000
frob-distance	0.2826	0.0160

1.2 ML Classification

Using Maximum Likelihood, a classification task with three different classes was performed. All the samples had the same probability of $p(x \in k) = \frac{1}{3}$ to be drawn from class $k = 1, 2, 3$. The samples of the three classes are drawn from Gaussian distributions with different means, but a common covariance matrix. A training ($Train$) and a testing ($Test$) data set with 1000 samples each was created.
The samples in $Train$ were used to calculate the ML estimates for the mean and covariance. Then, the points in $Test$ were classified by calculating the l2-distance to each mean μ_k and assigning it to the class with the smallest distance. In order to quantify the quality of the classification, the fraction of correct assignments was determined. All of this was done for a diagonal and non-diagonal covariance matrix.



It can be observed in the plots that the classes in the data generated with the diagonal covariance are slightly more separated than in the non-diagonal case. This difference is also reflected in the ability of the ML-classifiers to assign the samples to the correct class. The accuracy differs by 4%:

	diagonal	non-diagonal
fraction of correct assignment	0.9439	0.9039

2 Expectation Maximization

In this task, the EM algorithm is applied and studied for different Gaussian mixture distributions:

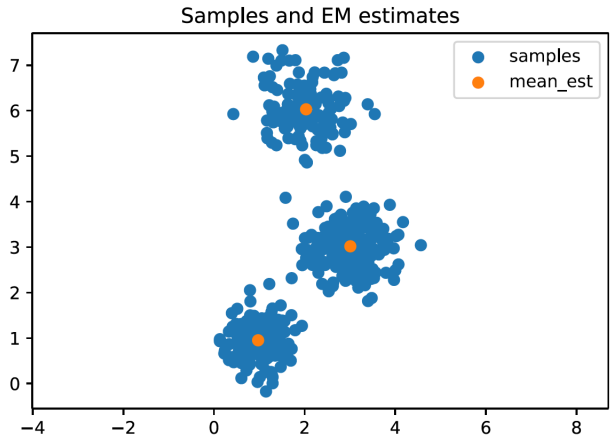
$$p(x) = \sum_{k=1}^{K} P_k \cdot p(x|\Phi_k)$$

2.1 Estimating Parameters

600 samples were drawn from the a Gaussian mixture model with the following parameters:

$$K = 3, \quad \mu_1 = [1,1]^T, \quad \mu_2 = [3,3]^T, \quad \mu_3 = [2,6]^T, \quad c_1 = 0.1 \cdot \mathbb{1},$$
$$c_2 = 0.2 \cdot \mathbb{1}, \quad c_3 = 0.3 \cdot \mathbb{1}, \quad P_1 = 0.4, \quad P_2 = 0.4, \quad P_3 = 0.2$$

This results in a distribution of samples shown below, where the mean estimates were determined with the EM algorithm using kmeans as the initialization method:



The following table shows the errors on the different parameters. For the weights and covariances, the error was calculated as the absolute difference between the true value and the estimate (covariance: factor before identity-matrix). The error on the means is given by the l2-distance. All parameters where estimated with a reasonable high precision and there were no significant differences between the three classes:

	class 1	class 2	class 3
weight_error	4.30e-05	7.83e-04	7.40e-04
mean_error	5.59e-02	1.94e-02	4.42e-02
covariance_error	1.68e-03	1.09e-03	4.56e-03

2.2 Varying Initial Parameters

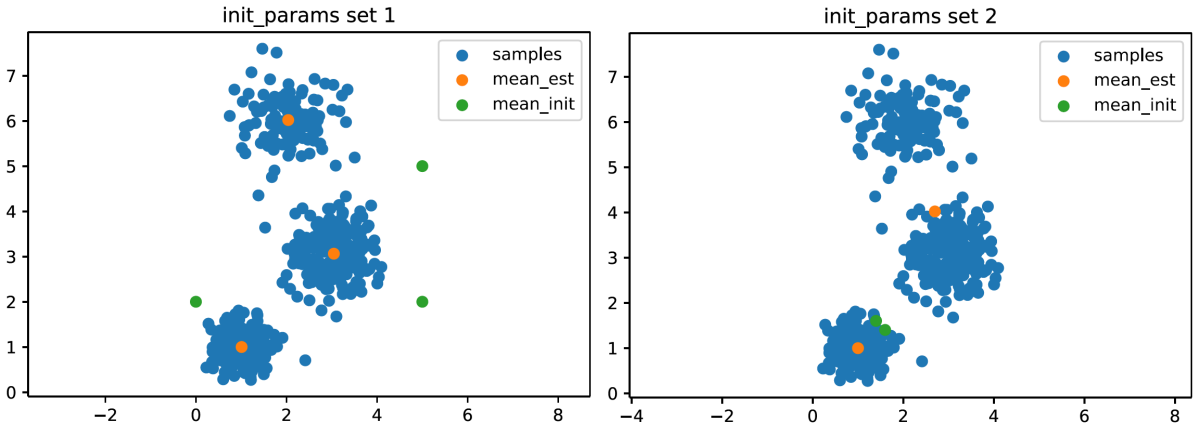
Then, the performance of EM was tested by using two different sets of initial parameters and observing the results. The first set of parameters was:

$$K = 3, \quad \mu_1 = [0, 2]^T, \quad \mu_2 = [5, 2]^T, \quad \mu_3 = [5, 5]^T,$$
$$c_1 = 0.15 \cdot \mathbb{1}, \quad c_2 = 0.27 \cdot \mathbb{1}, \quad c_3 = 0.4 \cdot \mathbb{1}, \quad P_1 = P_2 = P_3 = 0.33$$

And the second set:

$$K = 2, \quad \mu_1 = [1.6, 1.4]^T, \quad \mu_2 = [1.4, 1.6]^T,$$
$$c_1 = 0.2 \cdot \mathbb{1}, \quad c_2 = 0.4 \cdot \mathbb{1}, \quad P_1 = P_2 = 0.5$$

Below, the results for the two sets of initial parameters are shown.



With the first set of parameters, the algorithm converges to the correct results with similar errors as before with kmeans used as an initialization method.
Of course, EM had no chance to determine good estimates with the second set of initial parameters, as it tried to find estimates for two instead of three classes. However, the mean of class 1 was estimated quite precisely and the second mean was estimated to lie between classes 2 and 3. Furthermore, it is interesting to note that the estimated weights and covariances seem to indicate that classes 2 and 3 where taken to be one class by the EM algorithm.
After further experimentation with the initial parameters, the following can be summarized for the EM algorithm: if the number of classes is not set correctly initially, the results will

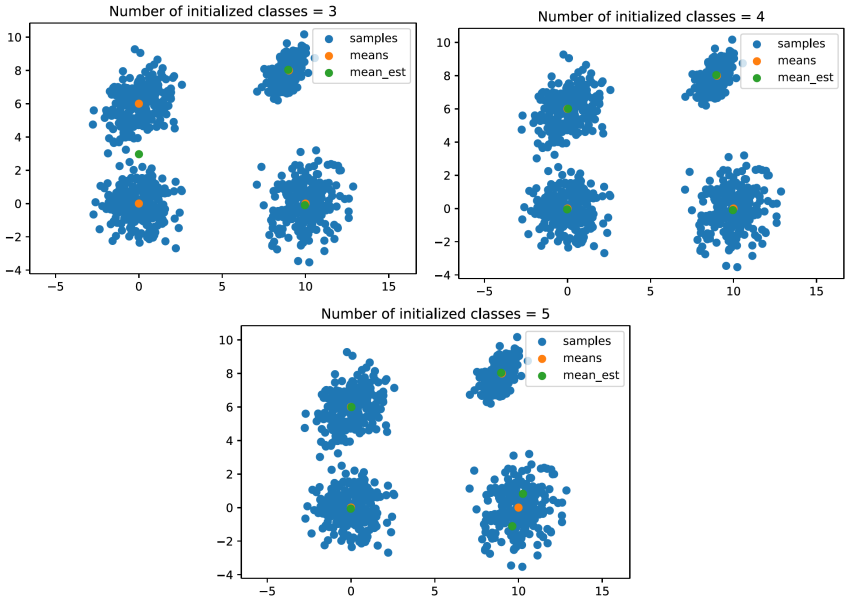
deviate greatly from the exact parameters; the other initial parameters have to be roughly in the right ballpark, whilst the EM seems to be most sensitive to the initialization of the means.

3 Clustering

A data set $Train$ was created that contains $N = 1000$ samples, grouped into four equally sized classes. Each class is described by a Gaussian distribution with the following parameters:

$$\mu_1 = [0,0]^T, \quad \mu_2 = [10,0]^T, \quad \mu_3 = [0,6]^T, \quad \mu_4 = [9,8]^T,$$
$$c_1 = \mathbb{1}, \quad c_2 = \begin{pmatrix} 1 & 0.2 \\ 0.2 & 1.5 \end{pmatrix}, \quad c_3 = \begin{pmatrix} 1 & 0.4 \\ 0.4 & 1.1 \end{pmatrix}, \quad c_4 = \begin{pmatrix} 0.3 & 0.2 \\ 0.2 & 0.5 \end{pmatrix}$$

Now the K-means algorithm is applied to $Train$ with different initial parameters. First, only the number of initialized classes was varied.

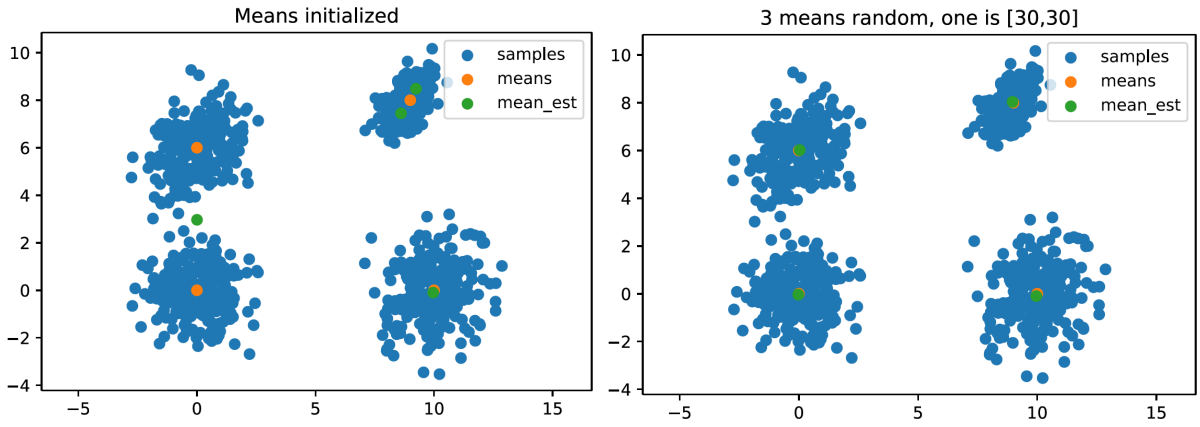


Obviously, the algorithm performed best in the case where the number of initialized classes matches the actual number of classes. In the case with three initialized classes, K-Means takes the classes 1 and 3 to be one, probably because they are closest together. In the last case, the algorithm splits class 2 in half, seemingly since it is spread the most (see covariance matrices).

Furthermore, the algorithm was run two more times, with the correct number of classes initialized. Once, with the means initialized in the following way:

$$\mu_1 = [-2, -2]^T, \quad \mu_2 = [-2.1, -2.1]^T, \quad \mu_3 = [-2, -2.2]^T, \quad \mu_4 = [-2.1, -2.2]^T$$

And another time with the first three means initialized randomly and the fourth as $\mu_4 = [30, 30]^T$.



This shows that even when the number of classes is initialized correctly, the results depend highly on the initial means. On the other hand, it is interesting that K-means can sometimes handle one mean to be initialized completely wrong. However, in some runs of the algorithm (with different random means and the same fourth mean $\mu_4 = [30, 30]^T$), it didn't converge correctly, supporting the statement that the results are highly dependent on the initial conditions.

Maximum_Likelihood

January 20, 2021

[ ]:
```
import numpy as np
import matplotlib.pyplot as plt
from mpl_toolkits.mplot3d import Axes3D
```

ML Estimates

[ ]:
```
#Defining mean and covariance of the distribution
mean =[2,-2]
cov = [[0.9,0.2],[0.2,0.3]]

for n_samples in [50, 5000]:
    #generate samples and calculate ML-estimates for the mean and covariance
    X = np.random.multivariate_normal(mean,cov,n_samples)
    mean_est = np.mean(X,axis=0)
    cov_est = np.cov(X.T)

    #print mean and covariance estimates and quality metrics
    print('mean_' + str(n_samples) + ' :')
    print(mean_est)
    print('\nl2_distance_' + str(n_samples) + ' :')
    print(np.linalg.norm(mean-mean_est))

    print('\ncovariance_' + str(n_samples) + ' :')
    print(cov_est)

    print('\nfrob_distance_' + str(n_samples) + ' :')
    print(np.linalg.norm(np.subtract(cov,cov_est)))

    #plot the samples and the true and estimated means
    plt.scatter(X[:,0], X[:,1], label='samples')
    plt.scatter(mean_est[0], mean_est[1], label='mean_est')
    plt.scatter(mean[0], mean[1], label='mean')
    plt.axis('equal')
    plt.title('n = ' + str(n_samples))
    plt.legend()
    plt.show()
```

ML Classification

[ ]:
```
mean1 = [0,0,0]
mean2 = [1,2,2]
mean3 = [3,3,4]
cov_diag = 0.8*np.identity(3)                                    #for the␣
 ↪first part of the exercise (diagonal covariance)
cov_non_diag = np.array([[0.8,0.2,0.1],[0.2,0.8,0.2],[0.1,0.2,0.8]])     #for␣
 ↪the second part of the exercise (non-diagonal covariance)
n_samples = 999
titles = ['Test-Data diagonal', 'Test-Data non-diagonal']
counter = 0
for cov in [cov_diag, cov_non_diag]:
    #Generating the train and test data for the three classes and merging them␣
 ↪into one dataset for training and one for testing
    Train1 = np.random.multivariate_normal(mean1,cov,int(n_samples/3))
    Train2 = np.random.multivariate_normal(mean2,cov,int(n_samples/3))
    Train3 = np.random.multivariate_normal(mean3,cov,int(n_samples/3))
    Train = np.concatenate((Train1,Train2,Train3), axis=0)

    Test1 = np.random.multivariate_normal(mean1,cov,int(n_samples/3))
    Test2 = np.random.multivariate_normal(mean2,cov,int(n_samples/3))
    Test3 = np.random.multivariate_normal(mean3,cov,int(n_samples/3))
    Test = np.concatenate((Test1,Test2,Test3), axis=0)

    #Plotting the Test-Data to show the difference caused by the different␣
 ↪covariances
    fig = plt.figure()
    ax = Axes3D(fig)
   # ax.view_init(30, 30)
    ax.scatter(Test[:,0], Test[:,1],Test[:,2])
    ax.set_xlabel('X-axis')
    ax.set_ylabel('Y-axis')
    ax.set_zlabel('Z-axis')
    plt.title(titles[counter])
    plt.show()

    counter = counter + 1


    #Calculating ML estimates for mean and covariance
    mean1_est = np.mean(Train1,axis=0)
    mean2_est = np.mean(Train2,axis=0)
    mean3_est = np.mean(Train3,axis=0)
    Cov1_est = np.cov(Train1.T)
    Cov2_est = np.cov(Train2.T)
    Cov3_est = np.cov(Train3.T)
    Cov_est = (Cov1_est + Cov2_est + Cov3_est)/3
```

```
    #Classifying the samples in Test according to the distance to the means of␣
↪the classes
    class_label = []
    correct_label = 0
    for x in Test:
        #calculating the distance of x to each mean
        d1 = np.linalg.norm(x-mean1_est)
        d2 = np.linalg.norm(x-mean2_est)
        d3 = np.linalg.norm(x-mean3_est)

        #finding the smallest distance and saving the according class label to␣
↪the list class_label
        m = min(d1,d2,d3)
        if m == d1:
            class_label.append(1)
        elif m == d2:
            class_label.append(2)
        elif m == d3:
            class_label.append(3)

    #Calculating how many samples are labeled correctly (the implementation␣
↪works because the samples in Test are ordered according to classes)
    for i in range(len(class_label)):
        if i < 333:
            if class_label[i] == 1:
                correct_label +=1
        elif i >= 333 and  i < 666:
            if class_label[i] == 2:
                correct_label +=1
        elif i >= 666 and  i < 999:
            if class_label[i] == 3:
                correct_label +=1

    correct_frac = correct_label/len(Test)
    print('fraction of correct assignment: ' + str(correct_frac) + '\n')
```

[ ]:

Expectation_Maximization

January 20, 2021

[ ]:
```
import numpy as np
import matplotlib.pyplot as plt
from sklearn import mixture
```

Estimating Parameters

[ ]:
```
#Define means and covariances of the three classes
mean1 = [1,1]
mean2 = [3,3]
mean3 = [2,6]
mean = [mean1,mean2, mean3]         #used for error calculation later
cov1 = 0.1*np.identity(2)
cov2 = 0.2*np.identity(2)
cov3 = 0.3*np.identity(2)
cov = [0.1, 0.2, 0.3]
weights = [0.4, 0.4, 0.2]           #used for error calculation later
n_samples = 600

#Generate the training data
Train1 = np.random.multivariate_normal(mean1,cov1,int(n_samples*0.4))
Train2 = np.random.multivariate_normal(mean2,cov2,int(n_samples*0.4))
Train3 = np.random.multivariate_normal(mean3,cov3,int(n_samples*0.2))
Train = np.concatenate((Train1,Train2,Train3), axis=0)
```

[ ]:
```
#Use EM algorithm to determine the parameters of the gaussian mixture␣
 ↪distribution generated above
gmm = mixture.GaussianMixture(n_components=3,␣
 ↪covariance_type='spherical',random_state=5)
gmm.fit(Train)

#plot the data and the estimates for the means
plt.scatter(Train[:,0], Train[:,1], label='samples')
plt.scatter(gmm.means_[:,0], gmm.means_[:,1], label='mean_est')
plt.title('Samples and EM estimates')
plt.axis('equal')
plt.legend()
plt.show()
```

```
#calculate and print the errors for the different parameters in each class
for k in [0,1,2]:
    weight_error = abs(weights[k] - gmm.weights_[k])
    mean_error = np.linalg.norm(mean[k] - gmm.means_[k])
 ↪    #error in l2 norm
    covariance_error = abs(cov[k] - gmm.covariances_[k])

    print('\nResults for class ' + str(k+1) + ':')
    print('weight_error = ' + str(weight_error))
    print('mean_error = ' + str(mean_error))
    print('covariance_error = ' + str(covariance_error))
```

Varying Initial Parameters

[ ]:
```
#Setting the initial parameters of the two cases
K = [3,2]
weights_init = [[0.34, 0.33, 0.33], [0.5, 0.5]]
means_init = [[[0,2], [5,2], [5,5]], [[1.6,1.4], [1.4,1.6]]]
precision_init = [[6.66, 3.70, 2.50], [5, 2.5]]
titles = ['init_params set 1', 'init_params set 2']

for i in [0,1]:
    gmm = mixture.GaussianMixture(n_components=K[i],
 ↪covariance_type='spherical', weights_init=weights_init[i],
 ↪means_init=means_init[i], precisions_init=precision_init[i] ,random_state=50)
    gmm.fit(Train)

    tmp = np.array(means_init[i])   #constructing an array from the list, to
 ↪plot it more easily

    #plot the samples and the estimated means for each set
    plt.scatter(Train[:,0], Train[:,1], label='samples')
    plt.scatter(gmm.means_[:,0], gmm.means_[:,1], label='mean_est')
    plt.scatter(tmp[:,0], tmp[:,1], label='mean_init')
    plt.legend()
    plt.axis('equal')
    plt.title(titles[i])
    plt.show()

    #print all the estimated parameters
    print("weights:")
    print(gmm.weights_)
    print("means:")
    print(gmm.means_)
    print("covariances:")
    print(gmm.covariances_)
```

Classification

January 20, 2021

[ ]:
```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
```

[ ]:
```
#Defining means and covariances for the distributions
mean1 = [0,0]
mean2 = [10,0]
mean3 = [0,6]
mean4 = [9,8]
means = np.array([mean1, mean2, mean3, mean4])      #needed for the plots later
cov1 = np.identity(2)
cov2 = [[1,0.2],[0.2,1.5]]
cov3 = [[1,0.4],[0.4,1.1]]
cov4 = [[0.3,0.2],[0.2,0.5]]
n_samples = 1000

#creating the data set Train to which kmeans will be fitted later
Train1 = np.random.multivariate_normal(mean1,cov1,int(n_samples*0.25))
Train2 = np.random.multivariate_normal(mean2,cov2,int(n_samples*0.25))
Train3 = np.random.multivariate_normal(mean3,cov3,int(n_samples*0.25))
Train4 = np.random.multivariate_normal(mean4,cov4,int(n_samples*0.25))
Train = np.concatenate((Train1,Train2,Train3,Train4), axis=0)
```

[ ]:
```
#running kmeans for different numbers of initialized classes
C = [4,3,5]
for c in C:
    kmeans = KMeans(n_clusters=c, init='random', algorithm='full', n_init=1,␣
 ↪random_state=50)
    kmeans.fit(Train)

    #plotting the data and the results
    plt.scatter(Train[:,0], Train[:,1], label='samples')
    plt.scatter(means[:,0], means[:,1], label='means')
    plt.scatter(kmeans.cluster_centers_[:,0],kmeans.cluster_centers_[:,1],␣
 ↪label='mean_est')
    plt.axis('equal')
    plt.legend()
```

```
    plt.title('Number of initialized classes = ' + str(c))
    plt.show()
```

[ ]:
```
#running kmeans for two additional initializations
means_init1 = np.array([[-2,-2], [-2.1,-2.1], [-2,-2.2], [-2.1,-2.2]])
means_init2 = np.array([[np.random.randint(-5,15),np.random.randint(-4,11)],␣
 ↪[np.random.randint(-5,15),np.random.randint(-4,11)], [np.random.
 ↪randint(-5,15),np.random.randint(-4,11)], [30,30]]) #x- and y-value are␣
 ↪randomly chosen within the range these values lie in the Train data set
means_init = [means_init1, means_init2]
titles = ['Means initialized', '3 means random, one is [30,30]']

for i in range(len(means_init)):
    kmeans = KMeans(n_clusters=4, init=means_init[i], algorithm='full',␣
 ↪random_state=1)
    kmeans.fit(Train)

    plt.scatter(Train[:,0], Train[:,1], label='samples')
    plt.scatter(means[:,0], means[:,1], label='means')
    plt.scatter(kmeans.cluster_centers_[:,0],kmeans.cluster_centers_[:,1],␣
 ↪label='mean_est')
    plt.axis('equal')
    plt.legend()
    plt.title(titles[i])
    plt.show()
```