PGM HOMEWORK2 REPORT

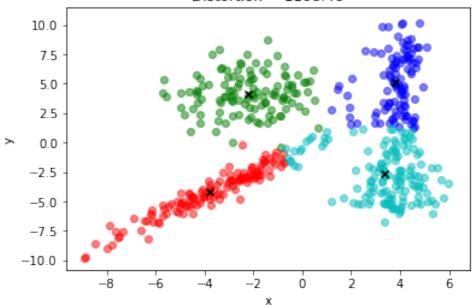
Please, see the scans in the end of the report for theoretical questions.

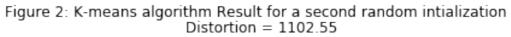
4 Implementation - Gaussian mixtures :

a) For the implementation of the first question: $K_means(data,k)$ which is initialized randomly by k different points from data and returns a list containing in the first element the clusters found and in the second their centers. The function distortion(kmeans) takes the k_means results as arguments and return the distortion of the classes found.

To plot the clusters and theirs centers, I implemented the function plt_kmeans(cluster,center,color) which plot the cluster with color color and its center in black.

Figure 1: K-means algorithm Result for a random intialization Distortion = 1108.46





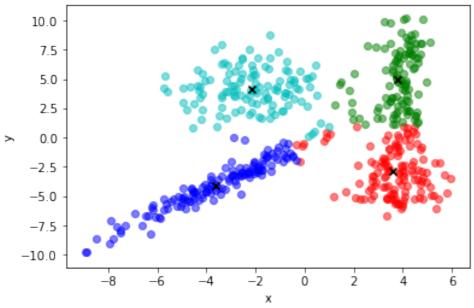


Table 1: Comparison between different results of k-means algorithms on train data

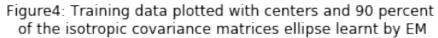
Clusters' centers	Distortion	Clusters' centers	Distortion
[[3.6 -2.89] [-2.16 4.11] [-3.64 -4.05] [3.79 5.]]	1102.54	[[-2.14 3.97] [-3.72 -4.18] [3.79 5.] [3.57 -2.88]]	1103.92
[[-3.78 -4.22] [-2.24 4.16] [3.8 5.1] [3.36 -2.66]]	1108.46	[[-2.24 4.13] [-3.82 -4.27] [3.8 5.1] [3.34 -2.64]]	1109.42
[[3.8 5.1] [-3.66 -4.07] [3.48 -2.7] [-2.24 4.24]]	1105.84	[[-3.78 -4.22] [3.36 -2.71] [-2.24 4.16] [3.8 5.03]]	1107.88

From the different results, we could conclude that k-means algorithm get trapped in local minimas and we have to run it several time with random initialization and preserve the best clustering which has the minimal distortion. However, we can notice that for the results in the table 1 the distortion value is in the range [1102,1110] which gives an error of 8/1106 = 0.7% which is not a large precision error. We can notice also that the centers are approximately the same (differ by the first value after comma).

b) For the theoretical question, please see the scans in the end of the end of the report. For the implementation of the EM algorithm for Gaussian Mixture when the covariance matrices are isotropic (proportional to the identity matrix), I used the functions: initializationStep(data,k): Which returns initialized means of gaussians and weights with K means algorithm and coefficients of covariances with a strict positive random value, stepE(data, means, cov, weights, k) which returns the responsibilities (posterior probabilities) and the current log-likelihood, stepM(data, resp, k) which returns the updated means, variances and weights given the responsabilities. Then implemented the EM algorithm with function algorithmEM(data,k) which returns the means, covariances, weights and responsibilities when the log-likelihood converges.

The graphical Representation of the training data, the centers, as well as 90% percentage of covariance ellipses learnt by EM algorithm is shown in Figure 4.

For the representation of the latent variables for all data points with the parameters learned by EM, I implemented the function clusters(data,k) which assigns each point to the cluster with the highest value of responsibility. The Figure 5 shows the clusters with different colors.



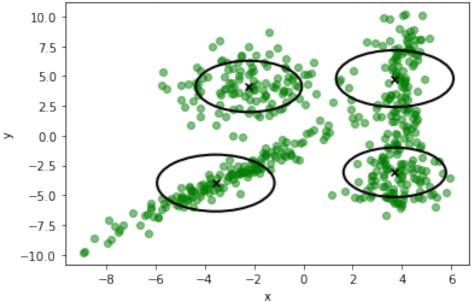
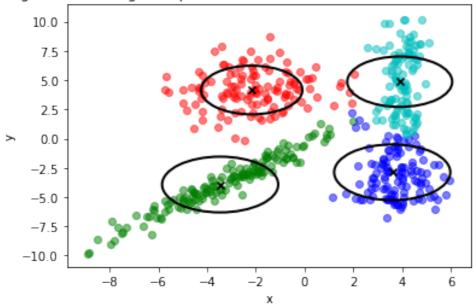


Figure 5: Training data plotted with different colors for latent variables



c). For the implementation of the EM algorithm for Gaussian Mixture when the covariance matrices are in general form, I used the functions: intializationStepGeneralCase(data,k) Which returns initialized means of gaussians, weights and covariances with K means algorithm. stepE(data, means, cov, weights, k)

the same one as for the isotropic case and stepMGeneralCase(data,resp,k) which returns the updated means, covariance matrices and weights given the responsabilities. Then implemented the EM algorithm with function algorithmEMgeneralCase(data,k) which returns the means, covariances, weights and responsibilities when the log-likelihood converges.

The graphical Representation of the training data, the centers, as well as 90% percentage of covariance ellipses learnt by EM algorithm is shown in Figure 6.

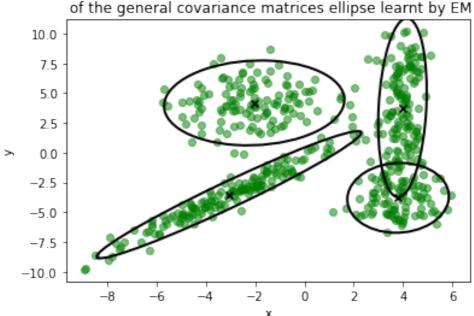


Figure 6: Training data plotted with centers and 90 percent of the general covariance matrices ellipse learnt by EM

For the representation of the latent variables for all data points with the parameters learned by EM, I implemented the function <code>clustersEMgeneralCase(data,k)</code> which assigns each point to the cluster with the highest value of responsibility. The Figure 7 shows the clusters with different colors.

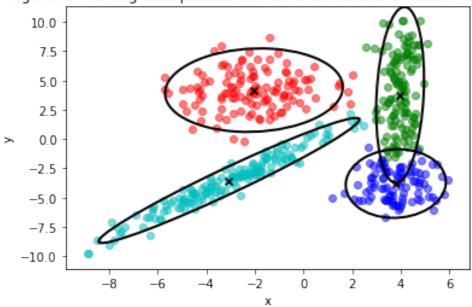


Figure 7: Training data plotted with different colors for latent variables

d) The comparison of maximum log-likelihood between different models and data (train and test) is show in the following table:

Table 2: Comparison	- C N A - ' I	19 19	C T	
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	Maximum Log- Likelihood on Train data	Maximum Log-Likelihood on Test data
Isotropic covariance matrices	-4469.33	-4341.51
General covariance matrices	-3358.18	-3404.92

We can notice that the maximum log-likelihood when using general covariance matrices is larger than the one using isotropic covariance matrices, but if we compare the elapsed times while convergence of the two models (34.39 seconds on test-data with isotropic model and 48.83 seconds on test-data with general model), we find that the second one converges rapidly than the first, this was expected because the isotropic covariances has only 1 parameter to learn while in the general case we have d*(d+1)/2 with *d: the dimension of data.* Moreover, by analyzing the figures 4, 5, 6 and 7, we can say that the model of isotropic covariances isn't suitable for our data.

RHOULAM

HOME WORK2 - PGM

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2) Let 6 be a directed free and 6'it's corresponding underected free First let show that the maximal diques in 6' are composed of only two nodes if IKI > 2 Let enpose that there exists a chique of 3 nodes 12,8,0 in 6° then it takes then sin 6 6' is the und rected free corresponding to 6, there exists an orientation of plus lique front is m &. We can see know all the 32 = 9 partible overtation gives a v stricture; but the directed tree G doesn't contains v. structures then it's an absund. Then: | Chiques (6'1) <2 if G'= { Lood S, d) than cliques (G')= 1. and the har exists a nade that

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3- Entropy and Mutual Information:
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