1. Fitting Gaussian Mixture Models

This is an exercise in fitting mixtures of Gaussians to data. I will have you fit standard mixture models and mixture models with *spherically symmetric* covariance matrices (as you explored in the last homework for single Gaussians. You will need to write code for the EM algorithm. I encourage you to write, test, and debug the code in groups. But please run your own experiments and produce your own reports.

On the Blackboard page is a dataset *toydata1.txt* consisting of 1500, 2-d vectors. One vector per row.

(a) Make a scatter plot of the data and fit models with $2,3,4,5,\ldots$ up to 10-20 Gaussian components. Fit the model on the first 500 data vectors using the EM procedure.

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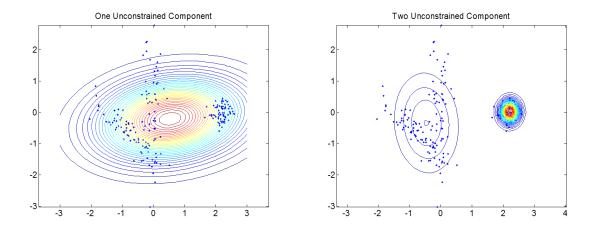
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on the traini llection of all the model parameters α_k , μ_k , Σ_k , k umber of mixture components. Since the Wals fritting finds equal to the model of the model of the position of each model. (For each model, keep the position) with the highest \mathcal{L} on the training set, as that's the best fit.)

Also make plots of the log-likelihood per data point on the holdout set (the remaining 1000 points in the data set) for each model trained. Contrast how the log-likelihood on the training set behaves as the number of components is increased with how the log-likelihood on the holdout set behaves. For several of your solutions, compare a scatter plot of the data with either a contour plot, or a surface plot of the model density. (I've included three examples below.) Comment on the results.

(b) Repeat the exercise for a model built with spherical Gaussian mixture components — i.e. components with covariance matrices $\sigma_k^2 I$, $k=1\ldots L$. As before, make plots of the log-likelihood per data point on the training set vs the number of mixture components. Repeat for the log-likelihood per data point on the holdout set. Contrast how the log-likelihood on the training set behaves as the number of components is increased with how the log-likelihood on the holdout set behaves. And contrast the results from this part with those for the unconstrained model in part (a). For several of your solutions, compare a scatter plot of the data with either a contour plot, or a surface plot of the model density. Comment on the results.



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Here's pseudocode (nearly full code!) for the EM procedure to fit the mixture of Gaussian models. It follows the procedure given in the lecture notes (but includes a regularizer)— so re-read those too.

```
Load training data D = \{ x_a, a=1...N \} % N samples, each of dimension dim
Set Number of Components L
Intialize:
       for i=1...L
          alpha_i = 1/L
         mu_i = choose randomly from data points,
         Sigma_i = IdentityMatrix
       end; i
logLike = log p(D|{alpha_i, mu_i, Sigma_i, i=1...L} )
oldLogLike = logLike-1
While ( | oldLogLike - logLike | > 0.00001 ) AND ( iterations < 500 )
    % E-Step
       for a=1 Assignment Project Exam Help
         for i=1...L
              h_{ia} = alpha_{i} * p(x_{i})
              % This regularity regu
               h_{ia} = (h_{ia} + epsilon) / (1 + L*epsilon)
          end % for i
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       end % for a
    % M-Step
     for i = 1 \dots L
       alpha_i = 1/N *sum (h_ia, a=1...N)
       pointsInI = sum( h_ia, a=1...N )
       mu_i = sum( h_ia * x_a, a=1...N ) / pointsInI
       if UnConstrained Covariance
         Sigma_i = sum( h_ia * ( x_a - m_i )*Transpose(x_a - m_i), a=1...N ) / pointsInI
         % Note (x_a - m_i) is a COLUMN vector
       else % Sigma_i = sigma2_i*IdentityMatrix
            sigma2_i = sum ( h_ia * Transpose(x_a-m_i)*(x_a-m_i), a=1...N ) / (pointsInI*dim)
              % Note this is 1/dim*Trace(SigmaUnconstrained) and you can compute it this way
            Sigma_i = sigma2_i * IndentityMatrix
     end % for i
oldLogLike = logLike
logLike = log p(D| { alpha_i, mu_i, Sigma_i, i=1...L } )
end % While
```