

# **Machine Learning**

Lecture. 11.

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Fundamental ML problem



- Fundamental ML problem
- Parametric Probability Density Estimation



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- Maximum Likelihood Estimation



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- Mixture Density Models



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- EM Algorithm



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- Given data and labels require estimation of each set of  $oldsymbol{ heta}_k$
- Employ likelihood function and estimate parameters which maximise the likelihood



•  $N_k$  examples from class k, assume D features are distributed as Multivariate Gaussian. Likelihood is  $\mathcal{L}_k$ 



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$$\prod_{n=1}^{N_k} p(\mathbf{x}_n | \boldsymbol{\theta}_k) = \prod_{n=1}^{N_k} p(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$$

$$= \prod_{n=1}^{N_k} \frac{1}{\sqrt{(2\pi)^D |\boldsymbol{\Sigma}_k|}} \exp\left\{-\frac{1}{2}(\mathbf{x}_n - \boldsymbol{\mu}_k)^\mathsf{T} \boldsymbol{\Sigma}_k^{-1} (\mathbf{x}_n - \boldsymbol{\mu}_k)\right\}$$



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Work with logarithm of likelihood  $\log \mathcal{L}_k$  and drop constant  $-\frac{N_k D}{2} \log 2\pi$ 

$$-\frac{N}{2}\log|\mathbf{\Sigma}_k| - \frac{1}{2}\sum_{n=1}^{N_k}(\mathbf{x}_n - \boldsymbol{\mu}_k)^\mathsf{T}\mathbf{\Sigma}_k^{-1}(\mathbf{x}_n - \boldsymbol{\mu}_k)$$



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- Expand the quadratic term and drop all terms independent of  $\mu_k$  then

$$\frac{\partial}{\partial \boldsymbol{\mu}_k} \log \mathcal{L}_k = \frac{\partial}{\partial \boldsymbol{\mu}_k} \left( \frac{1}{2} \sum_{n=1}^{N_k} \left\{ 2 \boldsymbol{\mu}_k^\mathsf{T} \boldsymbol{\Sigma}_k^{-1} \mathbf{x}_n - \boldsymbol{\mu}_k^\mathsf{T} \boldsymbol{\Sigma}_k^{-1} \boldsymbol{\mu}_k \right\} \right)$$



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Vector derivatives obtain

$$\frac{\partial}{\partial \boldsymbol{\mu}_k} \log \mathcal{L}_k = \sum_{n=1}^{N_k} \left\{ \boldsymbol{\Sigma}_k^{-1} \mathbf{x}_n - \boldsymbol{\Sigma}_k^{-1} \boldsymbol{\mu}_k \right\}$$



• Setting the gradient to zero we then obtain

$$\sum_{n=1}^{N_k} \mathbf{\Sigma}_k^{-1} \mathbf{x}_n = \sum_{n=1}^{N_k} \mathbf{\Sigma}_k^{-1} \boldsymbol{\mu}_k = N_k \mathbf{\Sigma}_k^{-1} \boldsymbol{\mu}_k$$



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• Now we can multiply both sides by the matrix  $\Sigma_k$  to obtain  $\sum_{n=1}^{N_k} \mathbf{x}_n = N_k \boldsymbol{\mu}_k$ 



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- Now we can multiply both sides by the matrix  $\Sigma_k$  to obtain  $\sum_{n=1}^{N_k} \mathbf{x}_n = N_k \boldsymbol{\mu}_k$
- Maximum-Likelihood estimate for the mean of the class-conditional Multivariate Gaussian as

$$\widehat{\boldsymbol{\mu}}_k = \frac{1}{N_k} \sum_{n=1}^{N_k} \mathbf{x}_n$$



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So

$$\frac{\partial}{\partial \Sigma_k} \frac{N_k}{2} \log |\Sigma_k| = \frac{N_k}{2|\Sigma_k|} |\Sigma_k| (\Sigma_k)^{-1} = \frac{N_k}{2} \Sigma_k^{-1}$$



• Cookbook, Section 2.2 expression third from bottom, shows that  $\frac{\partial}{\partial \mathbf{X}} \mathbf{a}^\mathsf{T} \mathbf{X}^{-1} \mathbf{b} = -\mathbf{X}^{-1} \mathbf{a} \mathbf{b}^\mathsf{T} \mathbf{X}^{-1}$ 



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$$-\sum_{n=1}^{N_k} \boldsymbol{\Sigma}_k^{-1} (\mathbf{x}_n - \boldsymbol{\mu}_k) (\mathbf{x}_n - \boldsymbol{\mu}_k)^\mathsf{T} \boldsymbol{\Sigma}_k^{-1}$$

Plugging everything together then we obtain

$$\frac{\partial}{\partial \Sigma_k} \log \mathcal{L}_k = -\frac{N_k}{2} \Sigma_k^{-1} + \frac{1}{2} \sum_{n=1}^{N_k} \Sigma_k^{-1} (\mathbf{x}_n - \boldsymbol{\mu}_k) (\mathbf{x}_n - \boldsymbol{\mu}_k)^\mathsf{T} \Sigma_k^{-1}$$



 Setting gradient to zero, replacing mean vectors with their ML estimates, after a little manipulation the estimate for the class-conditioned covariance is, as we would expect

$$\widehat{\Sigma}_k = \frac{1}{N_k} \sum_{n=1}^{N_k} (\mathbf{x}_n - \widehat{\boldsymbol{\mu}}_k) (\mathbf{x}_n - \widehat{\boldsymbol{\mu}}_k)^\mathsf{T}$$



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 ML estimation method can be adopted for any parametric form of probability density or distribution function. Of course we can also adopt a Bayesian approach by setting appropriate priors for the mean and covariance terms - we will resist this temptation for the time being

## **Illustrative Examples**



 Matlab script gauss\_density\_est.m generates random sample drawn from 2D Gaussian with parameters

$$\boldsymbol{\mu} = \begin{bmatrix} 1.0 \\ 3.0 \end{bmatrix} \quad \mathbf{C} = \begin{bmatrix} 1.5 & 0.6 \\ 0.6 & 0.4 \end{bmatrix}$$

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• Use sample to obtain estimates for the required parameters clearly sample size  $N \to \infty$  then estimates converge to true values

### **Illustrative Examples**



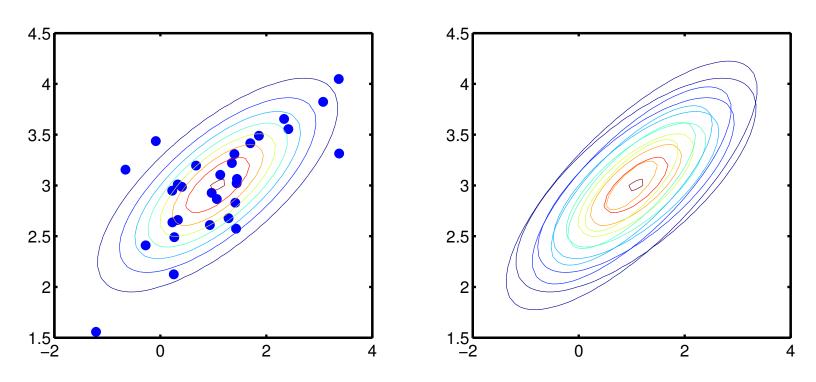


Figure 1: The left plot shows a random sample of 30 points drawn from a 2D Gaussian, the iso-contours of estimated probability density are superimposed on the plot. The iso-contours of probability density for the Gaussian with the actual parameter values are given on the right hand plot superimposed upon the iso-contours of estimated density.



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- The density is a mixture of two Gaussians with mean and covariances of

$$\mu_1 = \begin{bmatrix} 0.5 \\ 2.0 \end{bmatrix}$$
  $\mathbf{C}_1 = \begin{bmatrix} 1.0 & 0.0 \\ 0.0 & 1.0 \end{bmatrix}$ 

$$\mu_2 = \begin{bmatrix} 3.0 \\ 4.0 \end{bmatrix}$$
  $\mathbf{C}_2 = \begin{bmatrix} 1.0 & 0.0 \\ 0.0 & 1.0 \end{bmatrix}$ 



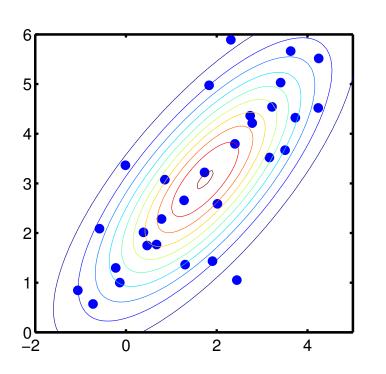
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 Data considered as coming from two sub-populations, or there are two distinct generating processes each responsible for producing the data we observe.





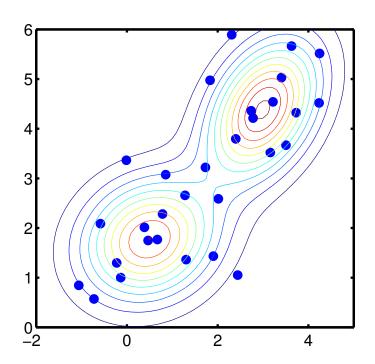


Figure 2: The left plot shows a random sample of 30 points drawn from 2 distinct 2D Gaussians, the iso-contours of estimated probability density under the assumption that the density is a single 2D Gaussian, are superimposed on the plot. The right plot shows the same random sample with the iso-contours of estimated probability density where the true functional form has been employed i.e. two 2D Gaussians.



• Average likelihood of points spread uniformly across the regions shown in the figures assuming a single Gaussian is -3.261.



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- This is higher than that achieved when assuming a single Gaussian and so provides a superior predictive generative model of the data.

# **Density Estimation**



- Average likelihood of points spread uniformly across the regions shown in the figures assuming a single Gaussian is -3.261.
- Average likelihood of points spread uniformly across the regions shown in the figures assuming a mixture of two Gaussians is -3.123.
- This is higher than that achieved when assuming a single Gaussian and so provides a superior predictive generative model of the data.
- A Matlab script mix\_gauss\_density.m is available on the course website to allow you to replicate these results.



 The probability density function for the case of two Gaussians can be represented as

$$p(\mathbf{x}|\boldsymbol{\theta}) = \pi p(\mathbf{x}|\boldsymbol{\theta}_1) + (1-\pi)p(\mathbf{x}|\boldsymbol{\theta}_2)$$
$$= \pi \mathcal{N}_{\mathbf{x}}(\boldsymbol{\mu}_1, \mathbf{C}_1) + (1-\pi)\mathcal{N}_{\mathbf{x}}(\boldsymbol{\mu}_2, \mathbf{C}_2)$$



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• where  $\theta=\{\pi, \theta_1, \theta_2\}$  and each set of parameters is defined by  $\theta_1=\{\mu_1, \mathbf{C}_1\}$  and  $\theta_2=\{\mu_2, \mathbf{C}_2\}$ 



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- where  $\theta=\{\pi, \theta_1, \theta_2\}$  and each set of parameters is defined by  $\theta_1=\{\mu_1, \mathbf{C}_1\}$  and  $\theta_2=\{\mu_2, \mathbf{C}_2\}$
- The parameter  $\pi$  is the probability that a point  $\mathbf{x}$  will be generated from  $p(\mathbf{x}|\boldsymbol{\theta}_1)$  and so the probability that the point will be generated from  $p(\mathbf{x}|\boldsymbol{\theta}_2)$  is  $1-\pi$



• In the more general case where there are  ${\cal M}$  components, of arbitrary parametric form the probability density will be expressed as

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• where the whole parameter set is defined as  $\boldsymbol{\theta} = \{\pi_1 \cdots \pi_M, \boldsymbol{\theta}_1 \cdots \boldsymbol{\theta}_M\}$  and  $\sum_{m=1}^M \pi_m = 1$  as each  $\pi_m$  is the probability that the mth component of the mixture will produce a data point so it must sum to one to be a valid probability over the M selection events



• Given data  $\mathcal{D} = \{\mathbf{x}_1 \cdots \mathbf{x}_N\}$  assuming mixture model  $p(\mathbf{x}|\boldsymbol{\theta}) = \sum_{m=1}^{M} \pi_m p(\mathbf{x}|\boldsymbol{\theta}_m)$  estimate parameters



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- Require estimates of each  $\pi_m$ , probability of data being generated by each m, just count how many points from  $\mathcal{D}$  coming from each of M components then normalise by N. Count  $N_m$  points in  $\mathcal{D}$  drawn from component m then

$$\widehat{\pi}_m = \frac{N_m}{N}$$

where each  $N_m$  can be obtained from  $N_m = \sum_{n=1}^N z_{mn}$  where each  $z_{mn} = 1$  if the nth point was drawn from component m and  $z_{mn} = 0$  otherwise.



• What of the specific parameters of each of the components  $\theta_m$ ? This is also easy as all we need to do is obtain the estimates  $\widehat{\theta}_m$  which maximise the likelihood of the data points which were drawn from component m under the parametric form  $p(\mathbf{x}|\boldsymbol{\theta}_m)$ .



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- For example if the mixture components were Gaussians then the Maximum-Likelihood estimate for the component mean vectors would simply be

$$\widehat{\boldsymbol{\mu}}_m = \frac{\sum_{n=1}^N z_{nm} \mathbf{x}_n}{\sum_{n=1}^N z_{nm}} = \frac{1}{N_m} \sum_{n \in m} \mathbf{x}_n$$



 The expression for the covariance matrices for each component would follow simply as

$$\widehat{\Sigma}_m = \frac{1}{N_m} \sum_{n=1}^{N} z_{mn} (\mathbf{x}_n - \widehat{\boldsymbol{\mu}}_m) (\mathbf{x}_n - \widehat{\boldsymbol{\mu}}_m)^{\mathsf{T}}$$

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- There is one small difficulty which we have overlooked, we do not have values for the indictor variables  $z_{mn}$  on which we have relied.
- This is a major difficulty as the fact that the variables  $z_{mn}$  are hidden or latent then our ML estimates cannot follow in the straightforward manner we had anticipated.



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- Need the joint likelihood of data  $\mathbf{X} = \{\mathbf{x}_1 \cdots \mathbf{x}_N\}$  and indicator variables  $\mathbf{Z} = \{\mathbf{z}_1 \cdots \mathbf{z}_N\}$  where each  $\mathbf{z}_n = \{z_{1n} \cdots z_{Mn}\}$



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- Given  $\theta = \{\theta_1 \cdots \theta_M\}$  we can marginalise over all possible component allocations

$$p(\mathbf{X}|\boldsymbol{\theta}) = \sum_{\mathbf{Z}} p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta})$$

where the summation is over all possible values which **Z** may take on.



$$\log p(\mathbf{X}|\boldsymbol{\theta}) = \log \sum_{\mathbf{Z}} p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta})$$
$$= \log \sum_{\mathbf{Z}} P(\mathbf{Z}|\mathbf{X}) \frac{p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta})}{P(\mathbf{Z}|\mathbf{X})}$$



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Use inequality  $\log E\{f(X)\} \ge E\{\log f(X)\}$  so can write



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$$\log \sum_{\mathbf{Z}} P(\mathbf{Z}|\mathbf{X}) \frac{p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta})}{P(\mathbf{Z}|\mathbf{X})} \geq \sum_{\mathbf{Z}} P(\mathbf{Z}|\mathbf{X}) \log \frac{p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta})}{P(\mathbf{Z}|\mathbf{X})}$$

$$= \sum_{\mathbf{Z}} P(\mathbf{Z}|\mathbf{X}) \log p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta})$$

$$- \sum_{\mathbf{Z}} P(\mathbf{Z}|\mathbf{X}) \log P(\mathbf{Z}|\mathbf{X})$$
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As  $\mathbf{x}_n$  drawn iid from m exclusively then summation over all  $\mathbf{Z}$  equals a summation over all n and m i.e.  $\mathcal{L}_B$  equals

$$\sum_{\mathbf{Z}} P(\mathbf{Z}|\mathbf{X}) \log \frac{p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta})}{P(\mathbf{Z}|\mathbf{X})} = \sum_{m,n}^{M,N} P(m|\mathbf{x}_n) \log \frac{p(\mathbf{x}_n|\boldsymbol{\theta}_m)P(m|\mathbf{x}_n)}{P(m|\mathbf{x}_n)}$$

$$= \sum_{m=1}^{N} \sum_{n=1}^{N} P(m|\mathbf{x}_n) \log p(\mathbf{x}_n|\boldsymbol{\theta}_m) P(m)$$

$$-\sum_{m=1}^{M}\sum_{n=1}^{N}P(m|\mathbf{x}_n)\log P(m|\mathbf{x}_n)$$

where now  $P(m|\mathbf{x}_n)$  is the probability that  $z_{mn}=1$  and P(m) is the probability that  $z_{mn}=1$  for any n.



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- The Expectation Maximisation (EM) algorithm is a general purpose method to *Maximise* the likelihood of the complete data ( $\mathbf{X} \& \mathbf{Z}$ ) so as to obtain estimates of the component parameters  $\boldsymbol{\theta}_m$ .
- Before performing the *Maximisiation* step we require to obtain the *Expected* values of a set of hidden binary allocation variables  $z_{mn}$ .
- Once we have obtained the *Expected* values of the latent variables we then perform the *Maximisation* step to obtain our current parameter estimates.
- This EM interleaving is continued until some convergence criterion is achieved.



• Taking functional derivatives of the lower-bound with respect to  $P(m|\mathbf{x}_n)$  then

$$\frac{\partial \mathcal{L}_B}{\partial P(m|\mathbf{x}_n)} = \log P(m|\mathbf{x}_n) - \log p(\mathbf{x}_n|\boldsymbol{\theta}_m)P(m) - 1$$



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• Setting to zero we see that  $P(m|\mathbf{x}_n) \propto p(\mathbf{x}_n|\boldsymbol{\theta}_m)P(m)$  and normalising appropriately yields the distribution of the form

$$P(m|\mathbf{x}_n) = \frac{p(\mathbf{x}_n|\boldsymbol{\theta}_m)P(m)}{\sum_{m'=1}^{M} p(\mathbf{x}_n|\boldsymbol{\theta}_{m'})P(m')}$$



• You should now be able to see that this is the posterior distribution over the mixture components m which generated  $\mathbf{x}_n$ , or the expected value of the binary variable  $z_{mn}$ .



- You should now be able to see that this is the posterior distribution over the mixture components m which generated  $\mathbf{x}_n$ , or the expected value of the binary variable  $z_{mn}$ .
- Now that we have maximised the bound with respect to the *Expected* value of the indicator variable we need to *Maximise* the bound with respect to the parameter values.



• The only terms in the bound  $\mathcal{L}_B$  which are dependent on the component parameters are

$$\sum_{m=1}^{M} \sum_{n=1}^{N} P(m|\mathbf{x}_n) \log p(\mathbf{x}_n|\boldsymbol{\theta}_m) P(m)$$

in which case we maximise the above with respect to each  $\theta_m$ .



• As an example assume that each  $p(\mathbf{x}_n|\boldsymbol{\theta}_m)$  is a multivariate Gaussian, then expanding and retaining the elements dependent on the parameters we obtain

$$- \frac{1}{2} \sum_{m=1}^{M} \sum_{n=1}^{N} P(m|\mathbf{x}_n) \log |\mathbf{\Sigma}_k|$$

$$- \frac{1}{2} \sum_{m=1}^{M} \sum_{n=1}^{N} P(m|\mathbf{x}_n) (\mathbf{x}_n - \boldsymbol{\mu}_m)^{\mathsf{T}} \mathbf{\Sigma}_m^{-1} (\mathbf{x}_n - \boldsymbol{\mu}_m)$$

$$+ \sum_{m=1}^{M} \sum_{n=1}^{N} P(m|\mathbf{x}_n) \log P(m)$$



• Taking derivatives wrt  $oldsymbol{\mu}_m$  and solving yields

$$\widehat{\boldsymbol{\mu}}_m = \frac{\sum_{n=1}^N P(m|\mathbf{x}_n)\mathbf{x}_n}{\sum_{n=1}^N P(m|\mathbf{x}_n)}$$



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• Nice result, compare with the estimator when we have perfect knowledge of the allocation variables  $z_{mn}$  i.e.

$$\widehat{\boldsymbol{\mu}}_m = \frac{\sum_{n=1}^N z_{mn} \mathbf{x}_n}{\sum_{n=1}^N z_{mn}}$$

so in the absence of the values  $z_{mn}$  we employ the expected values, or the posterior probabilities  $P(m|\mathbf{x}_n)$  which are obtained in the *Expectation* step



 Leaving you to have some fun with the derivation of the estimator for the covariance matrices we obtain

$$\widehat{\Sigma}_m = \frac{\sum_{n=1}^N P(m|\mathbf{x}_n)(\mathbf{x}_n - \widehat{\boldsymbol{\mu}}_m)(\mathbf{x}_n - \widehat{\boldsymbol{\mu}}_m)^{\mathsf{T}}}{\sum_{n=1}^N P(m|\mathbf{x}_n)}$$

again we can see that we have replaced perfect knowledge of the allocation variables with our current estimates of the posteriors  $P(m|\mathbf{x}_n)$ 



• Finally we need an estimate for P(m) taking derivatives then we observe that

$$P(m) \propto \sum_{n=1}^{N} P(m|\mathbf{x}_n)$$

This needs to be properly normalised and so

$$P(m) = \frac{1}{N} \sum_{n=1}^{N} P(m|\mathbf{x}_n)$$

### **EM Algorithm**



E Step

$$P(m|\mathbf{x}_n) = \frac{p(\mathbf{x}_n|\boldsymbol{\theta}_m)P(m)}{\sum_{m'=1}^{M} p(\mathbf{x}_n|\boldsymbol{\theta}_{m'})P(m')}$$

M Step

$$\widehat{\boldsymbol{\mu}}_{m} = \frac{\sum_{n=1}^{N} P(m|\mathbf{x}_{n}) \mathbf{x}_{n}}{\sum_{n=1}^{N} P(m|\mathbf{x}_{n})}$$

$$\widehat{\boldsymbol{\Sigma}}_{m} = \frac{\sum_{n=1}^{N} P(m|\mathbf{x}_{n}) (\mathbf{x}_{n} - \widehat{\boldsymbol{\mu}}_{m}) (\mathbf{x}_{n} - \widehat{\boldsymbol{\mu}}_{m})^{\mathsf{T}}}{\sum_{n=1}^{N} P(m|\mathbf{x}_{n})}$$

$$P(m) = \frac{1}{N} \sum_{n=1}^{N} P(m|\mathbf{x}_{n})$$



• Data with equal probability from three 2D Gaussians with a common unit variance i.e. I means of [0,0],[3,3],[-3,3] see Matlab file Gauss\_Mix\_Data.mat



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- There is also a  $1500 \times 2$  dimensional data set drawn from the same distribution which can be used to obtain values of likelihood on independent test data



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- There is also a  $1500 \times 2$  dimensional data set drawn from the same distribution which can be used to obtain values of likelihood on independent test data
- Wish to estimate probability density for data. There is, as always, one slight snag, our EM algorithm requires the number of components in the mixture. For now lets assume that we have a good idea what this value is



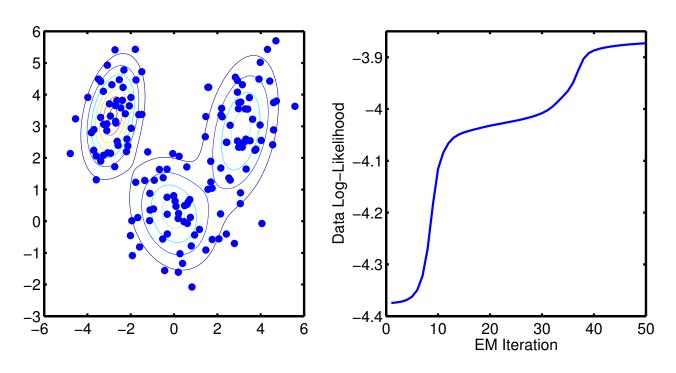


Figure 3: The left plot shows a random sample of 150 points drawn from 3 distinct 2D Gaussians, the iso-contours of estimated probability density under the assumption that the density is a mixture of three 2D Gaussians, are superimposed on the plot. The right plot shows the data likelihood under the mixture model at each EM step, it is clear that the likelihood does not decrease at each step.



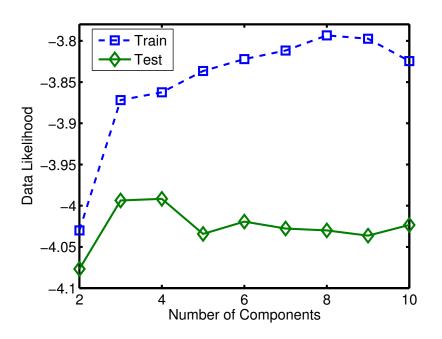


Figure 4: The left plot shows a random sample of 150 points drawn from 3 distinct 2D Gaussians, the iso-contours of estimated probability density under the assumption that the density is a mixture of three 2D Gaussians, are superimposed on the plot. The right plot shows the data likelihood under the mixture model at each EM step, it is clear that the likelihood does not decrease at each step.