

CSC 411: Lecture 13: Mixtures of Gaussians and EM

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- Mixture of Gaussians
- EM algorithm
- Latent Variables

A Generative View of Clustering

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 - ▶ It may help us decide on the number of clusters
- An obvious approach is to imagine that the data was produced by a generative model
 - ▶ Then we adjust the model parameters to maximize the probability that it would produce exactly the data we observed

Gaussian Mixture Model (GMM)

- A **Gaussian mixture model** represents a **distribution** as

$$p(\mathbf{x}) = \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x} | \mu_k, \Sigma_k)$$

with π_k the **mixing coefficients**, where:

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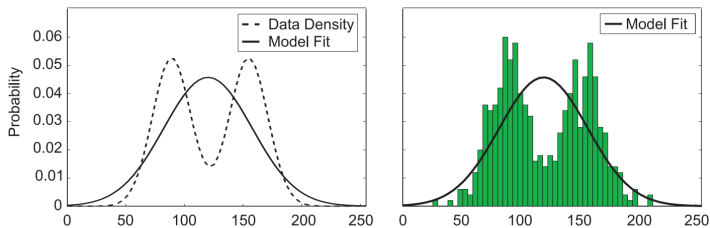
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- GMMs are **universal approximators of densities** (if you have enough Gaussians). Even diagonal GMMs are universal approximators.

Visualizing a Mixture of Gaussians – 1D Gaussians

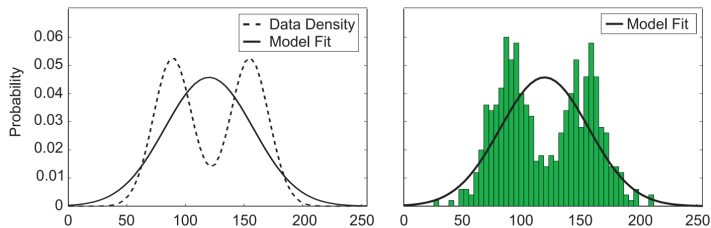
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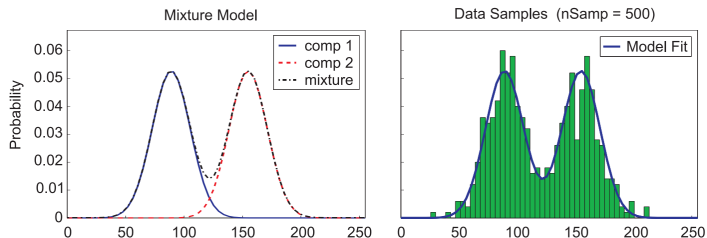
[Slide credit: K. Kutulakos]

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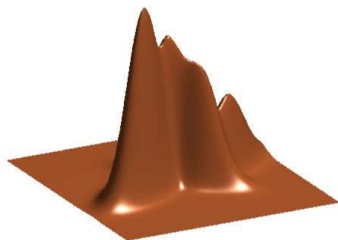
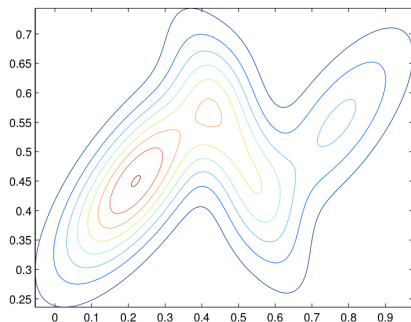
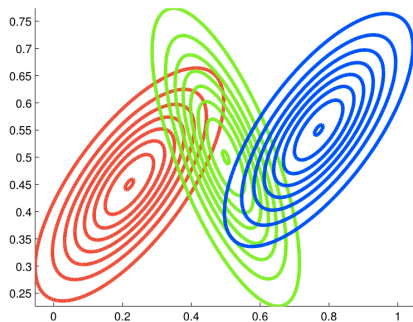


- Now, we are trying to fit a GMM (with $K = 2$ in this example):



[Slide credit: K. Kutulakos]

Visualizing a Mixture of Gaussians – 2D Gaussians



Fitting GMMs: Maximum Likelihood

- Maximum likelihood maximizes

$$\ln p(\mathbf{X}|\pi, \mu, \Sigma) = \sum_{n=1}^N \ln \left(\sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}^{(n)}|\mu_k, \Sigma_k) \right)$$

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- Don't forget to satisfy the constraints on π_k

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- Then:

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- In a **mixture model**, the identity of the component that generated a given datapoint is a latent variable

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- We would get this (check old slides):

$$\begin{aligned}\mu_k &= \frac{\sum_{n=1}^N \mathbf{1}_{[z^{(n)}=k]} \mathbf{x}^{(n)}}{\sum_{n=1}^N \mathbf{1}_{[z^{(n)}=k]}} \\ \Sigma_k &= \frac{\sum_{n=1}^N \mathbf{1}_{[z^{(n)}=k]} (\mathbf{x}^{(n)} - \mu_k)(\mathbf{x}^{(n)} - \mu_k)^T}{\sum_{n=1}^N \mathbf{1}_{[z^{(n)}=k]}} \\ \pi_k &= \frac{1}{N} \sum_{n=1}^N \mathbf{1}_{[z^{(n)}=k]}\end{aligned}$$

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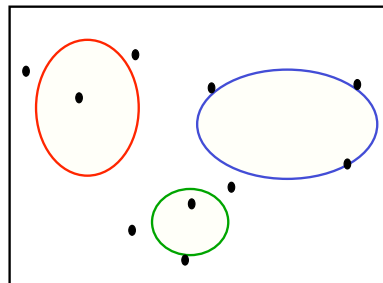
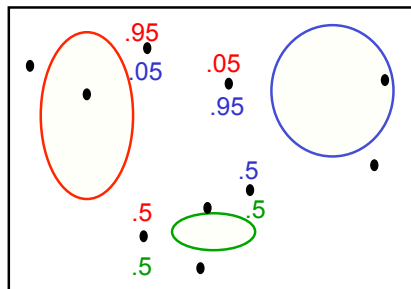
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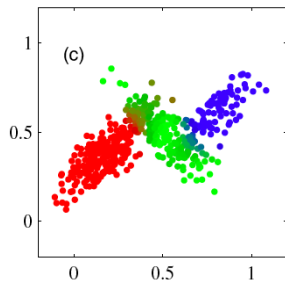
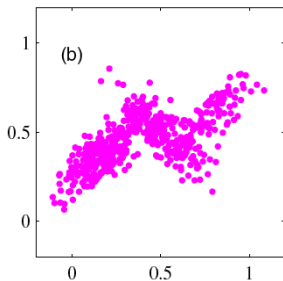
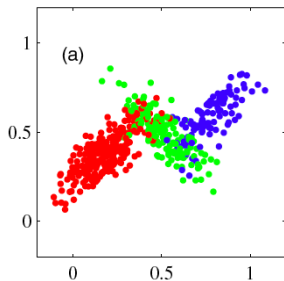
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- ▶ We can derive closed form updates for all parameters

Visualizing a Mixture of Gaussians



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- Conditional probability (using Bayes rule) of \mathbf{z} given \mathbf{x}

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- γ_k can be viewed as the **responsibility**

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- Set derivatives to 0:

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M-Step: Estimate Parameters

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- We used:

$$\mathcal{N}(\mathbf{x}|\mu, \Sigma) = \frac{1}{\sqrt{(2\pi)^d |\Sigma|}} \exp \left(-\frac{1}{2} (\mathbf{x} - \mu)^T \Sigma^{-1} (\mathbf{x} - \mu) \right)$$

and:

$$\frac{\partial (\mathbf{x}^T A \mathbf{x})}{\partial \mathbf{x}} = \mathbf{x}^T (A + A^T)$$

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- We just take the center-of gravity of the data that the Gaussian is responsible for
- Just like in K-means, except the data is weighted by the posterior probability of the Gaussian.
- Guaranteed to lie in the convex hull of the data (Could be big initial jump)

M-Step (variance, mixing coefficients)

- We can get similarly expression for the variance

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- Note that this is not a closed form solution of the parameters, as they depend on the responsibilities $\gamma_k^{(n)}$, which are complex functions of the parameters
- But we have a simple iterative scheme to optimize

EM Algorithm for GMM

- **Initialize** the means μ_k , covariances Σ_k and mixing coefficients π_k
- Iterate until convergence:
 - ▶ **E-step**: Evaluate the responsibilities given current parameters

$$\gamma_k^{(n)} = p(z^{(n)}|\mathbf{x}) = \frac{\pi_k \mathcal{N}(\mathbf{x}^{(n)}|\mu_k, \Sigma_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(\mathbf{x}^{(n)}|\mu_j, \Sigma_j)}$$

- ▶ **M-step**: Re-estimate the parameters given current responsibilities

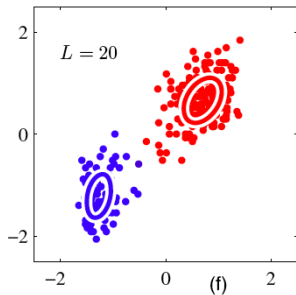
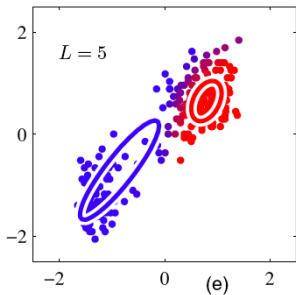
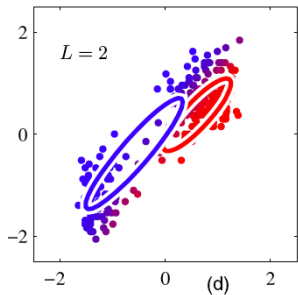
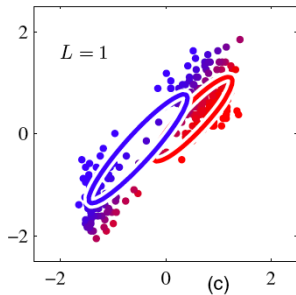
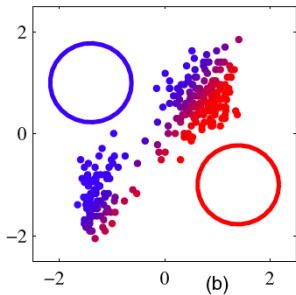
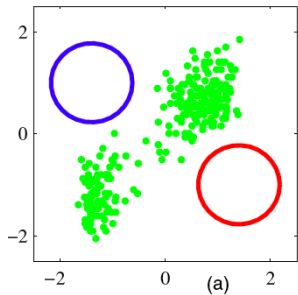
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- In K-means, weights are 0 or 1

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- In the E-step we compute $p(\mathbf{Z}|\mathbf{X}, \Theta^{old})$
- In the M-step we maximize w.r.t Θ

$$Q(\Theta, \Theta^{old}) = \sum_z p(\mathbf{Z}|\mathbf{X}, \Theta^{old}) \ln p(\mathbf{X}, \mathbf{Z}|\Theta)$$

General EM Algorithm

1. Initialize Θ^{old}
2. E-step: Evaluate $p(\mathbf{Z}|\mathbf{X}, \Theta^{old})$
3. M-step:

$$\Theta^{new} = \arg \max_{\Theta} Q(\Theta, \Theta^{old})$$

where

$$Q(\Theta, \Theta^{old}) = \sum_z p(\mathbf{Z}|\mathbf{X}, \Theta^{old}) \ln p(\mathbf{X}, \mathbf{Z}|\Theta)$$

4. Evaluate log likelihood and check for convergence (or the parameters). If not converged, $\Theta^{old} = \Theta$, Go to step 2

- Beyond this slide, read if you are interested in more details

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- A good way to show that this is OK is to show that there is a single function that is improved by both the E-step and the M-step.
 - ▶ The function we need is called **Free Energy**.

Why EM converges

- Free energy F is a cost function that is reduced by both the E-step and the M-step.

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- The **entropy** term encourages "soft" assignments. It would be happiest spreading the assignment probabilities for each datapoint equally between all the Gaussians.

Free Energy

- Our goal is to maximize

$$p(\mathbf{X}|\Theta) = \sum_{\mathbf{z}} p(\mathbf{X}, \mathbf{z}|\Theta)$$

- Typically optimizing $p(\mathbf{X}|\Theta)$ is difficult, but $p(\mathbf{X}, \mathbf{Z}|\Theta)$ is easy
- Let $q(\mathbf{Z})$ be a distribution over the latent variables. For any distribution $q(\mathbf{Z})$ we have

$$\ln p(\mathbf{X}|\Theta) = \mathcal{L}(q, \Theta) + KL(q||p(\mathbf{Z}|\mathbf{X}, \Theta))$$

where

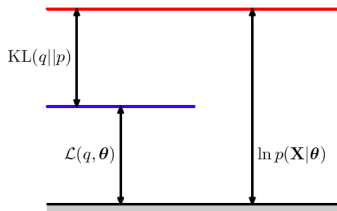
$$\mathcal{L}(q, \Theta) = \sum_{\mathbf{z}} q(\mathbf{Z}) \ln \left\{ \frac{p(\mathbf{X}, \mathbf{Z}|\Theta)}{q(\mathbf{Z})} \right\}$$

$$KL(q||p) = - \sum_{\mathbf{z}} q(\mathbf{Z}) \ln \left\{ \frac{p(\mathbf{Z}|\mathbf{X}, \Theta)}{q(\mathbf{Z})} \right\}$$

More on Free Energy

- Since the KL-divergence is always positive and have value 0 only if $q(Z) = p(\mathbf{Z}|\mathbf{X}, \Theta)$
- Thus $\mathcal{L}(q, \Theta)$ is a lower bound on the likelihood

$$\mathcal{L}(q, \Theta) \leq \ln p(\mathbf{X}|\Theta)$$



E-step and M-step

$$\ln p(\mathbf{X}|\Theta) = \mathcal{L}(q, \Theta) + KL(q||p(\mathbf{Z}|\mathbf{X}, \Theta))$$

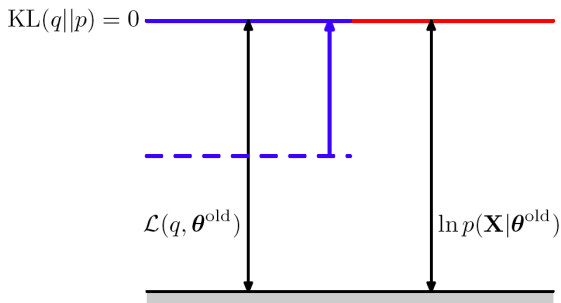
- In the E-step we maximize w.r.t $q(\mathbf{Z})$ the lower bound $\mathcal{L}(q, \Theta)$
- Since $\ln p(\mathbf{X}|\theta)$ does not depend on $q(\mathbf{Z})$, the maximum \mathcal{L} is obtained when the KL is 0
- This is achieved when $q(\mathbf{Z}) = p(\mathbf{Z}|\mathbf{X}, \Theta)$
- The lower bound \mathcal{L} is then

$$\begin{aligned}\mathcal{L}(q, \Theta) &= \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \Theta^{old}) \ln p(\mathbf{X}, \mathbf{Z}|\Theta) - \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \Theta^{old}) \ln p(\mathbf{Z}|\mathbf{X}, \Theta^{old}) \\ &= Q(\Theta, \Theta^{old}) + \text{const}\end{aligned}$$

with the content the entropy of the q distribution, which is independent of Θ

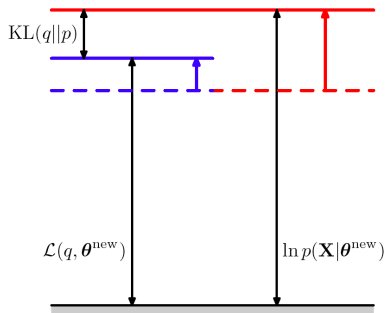
- In the M-step the quantity to be maximized is the expectation of the complete data log-likelihood
- Note that Θ is only inside the logarithm and optimizing the complete data likelihood is easier

Visualization of E-step



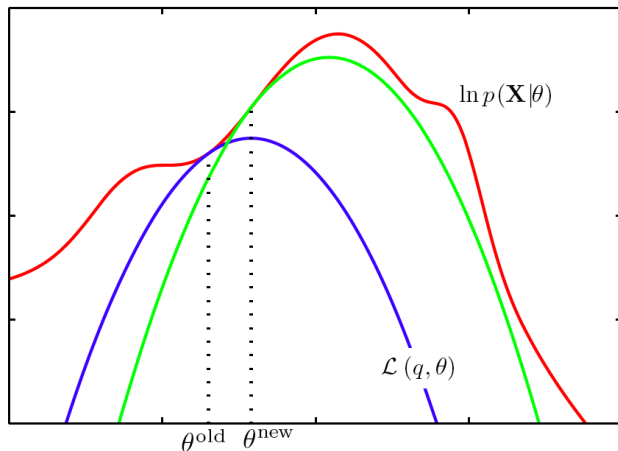
- The q distribution equal to the posterior distribution for the current parameter values Θ^{old} , causing the lower bound to move up to the same value as the log likelihood function, with the KL divergence vanishing.

Visualization of M-step



- The distribution $q(\mathbf{Z})$ is held fixed and the lower bound $\mathcal{L}(q, \Theta)$ is maximized with respect to the parameter vector Θ to give a revised value Θ^{new} . Because the KL divergence is nonnegative, this causes the log likelihood $\ln p(\mathbf{X}|\Theta)$ to increase by at least as much as the lower bound does.

Visualization of the EM Algorithm



- The EM algorithm involves alternately computing a lower bound on the log likelihood for the current parameter values and then maximizing this bound to obtain the new parameter values. See the text for a full discussion.

Summary: EM is coordinate descent in Free Energy

$$\begin{aligned}\mathcal{L}(q, \Theta) &= \sum_{\mathbf{z}} p(\mathbf{z}|\mathbf{X}, \Theta^{old}) \ln p(\mathbf{X}, \mathbf{z}|\Theta) - \sum_{\mathbf{z}} p(\mathbf{z}|\mathbf{X}, \Theta^{old}) \ln p(\mathbf{z}|\mathbf{X}, \Theta^{old}) \\ &= Q(\Theta, \Theta^{old}) + \text{const} \\ &= \text{expected energy} - \text{entropy}\end{aligned}$$

- The **E-step** minimizes F by finding the best distribution over hidden configurations for each data point.
- The **M-step** holds the distribution fixed and minimizes F by changing the parameters that determine the energy of a configuration.