ELEC 548 Mixture Models and Expectation Maximization

A. Gaussian Mixtures

In general, a *mixture model* can be defined for arbitrary probability distributions. However as is almost always the case, the Gaussian distribution makes everything work nicely. Mixture models are a key foundation for machine learning based on probabilistic generative models.

We will use a *Mixture of Guassians* as a probabilistic generative model for clustering. A good reference for this topic is Chapter 9 of Bishop's *Pattern Recognition and Machine Learning*. The first step to generate a data point is to start with the cluster identity.

Let $z \in \{1, \dots, K\}$ be a discrete random variable.

We can define the probability of drawing a point from an individual mixture component (i.e., cluster) as

$$Pr(z=k) = \pi_k, \text{ where } k=1,\dots,K. \tag{1}$$

Note that for $\Pr(z)$ to be a valid probability distribution, $0 \le \pi_k \le 1$ and $\sum_{k=1}^K \pi_k = 1$.

Aside: We have begun our discussion of mixture models implicitly assuming that the number of cluters, K, is specified. Non-parametric models, for which K would not be prespefied, have been an area of active development in the last decade. A good introduction to this topic is Chapter 25 of Murphy *Machine Learning: A Probabilistic Perspective*.

Once we have our *prior probability*, $\Pr(z)$, we can define the distribution of the data point from the selected mixture component

$$Pr(\mathbf{x} \mid z = k) = \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}) \tag{2}$$

Combining the prior probability of the cluster and the distribution of data points within the cluster we get the marginal probability of an observed data point:

$$\Pr(\mathbf{x}) = \sum_{z} \Pr(\mathbf{x} \mid z) \Pr(z) = \sum_{k=1}^{K} \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}) \boldsymbol{\pi}_{k}. \tag{3}$$

A peak ahead: We will use the $Pr(\mathbf{x})$ density with training data to do **maximum likelihood parameter estimation**. Then, with optimized parameters we can use the *a posteriori* density, $Pr(z \mid \mathbf{x})$, to assign data points to clusters either with a ""hard decision" (choosing the class which maximizes the density) or with a "soft decision" (keeping track of how likely the datapoint is to have come from each cluster).

A.1 Maximum likelihood parameter estimation

 $\textbf{Goal:} \quad \text{Fit } \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k, \boldsymbol{\pi}_k \text{ to training data } \mathbf{x}_1, \dots, \mathbf{x}_N.$

We will use the notational shorthand $\{\mathbf{x}\}$ for the training data $\mathbf{x}_1, \dots \mathbf{x}_N$, and the shorthand θ for all the parameters of the model $\{\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k, \pi_k\}_{k=1,\dots,K}$. Then, we can write down the likelihood of our training data

$$\begin{split} \Pr(\{\mathbf{x}\} \mid \boldsymbol{\theta}) &= \prod_{n=1}^{N} \Pr(\mathbf{x}_n) \\ &= \prod_{n=1}^{N} \sum_{k=1}^{K} \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \boldsymbol{\pi}_k \\ &\underbrace{\log \Pr(\{\mathbf{x}\} \mid \boldsymbol{\theta})}_{\text{call this } \mathcal{L}} &= \sum_{n=1}^{N} \log \left[\sum_{k=1}^{K} \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \boldsymbol{\pi}_k \right]. \end{split}$$

(i). Find μ_k :

$$\frac{\partial \mathcal{L}}{\partial \boldsymbol{\mu}_k} = \sum_{n=1}^{N} \frac{1}{\sum_{j=1}^{K} \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j) \cdot \boldsymbol{\pi}_j} \cdot \boldsymbol{\pi}_k \cdot \left(\frac{\partial}{\partial \boldsymbol{\mu}_k} \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right) \tag{4}$$

Looking at our handy table of matrix derivatives, we find that for a symmetric **A**, $\frac{\partial}{\partial \mathbf{x}} \left(\mathbf{x}^\mathsf{T} \mathbf{A} \mathbf{x} \right) = 2 \mathbf{A} \mathbf{x}$. Then, applying the chain rule, we have

$$\begin{split} \frac{\partial}{\partial \boldsymbol{\mu}} \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) &= \frac{\partial}{\partial \boldsymbol{\mu}} \left(\frac{1}{(2\pi)^{\frac{D}{2}} \left| \boldsymbol{\Sigma} \right|^{\frac{1}{2}}} e^{-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})} \right) \\ &= \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) \cdot \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu}). \end{split}$$

Plugging this into (4), we have

$$\begin{split} \frac{\partial \mathcal{L}}{\partial \pmb{\mu}_k} &= \sum_{n=1}^N \underbrace{\frac{\mathcal{N}(\mathbf{x}_n \mid \pmb{\mu}_k, \pmb{\Sigma}_k) \cdot \pi_k}{\sum_{j=1}^K \mathcal{N}(\mathbf{x}_n \mid \pmb{\mu}_j, \pmb{\Sigma}_j) \cdot \pi_j}}_{\text{call this } \gamma_{nk}} \cdot \pmb{\Sigma}^{-1}(\mathbf{x} - \pmb{\mu}) \\ &= \pmb{\Sigma}_k^{-1} \sum_{n=1}^N \gamma_{nk}(\mathbf{x} - \pmb{\mu}) \\ &= 0 \end{split}$$

Defining $N_k = \sum_{n=1}^N \gamma_{nk}$, we the result

$$\mu_k = \frac{1}{N_k} \sum_{n=1}^{N} \gamma_{nk} \mathbf{x}_n$$
 (5)

Notice that this result is very similar to the cluster update for K-means, with γ_{nk} replacing r_{nk} as tge "responsibility" that cluster k takes in explaining the observation \mathbf{x}_n . Notice that $0 \leq \gamma_{nk} \leq 1$ and $\sum_{k=1}^K \gamma_{nk} = 1$. Let's think about this by looking at what γ_{nk} is:

$$\begin{split} \gamma_{n\,k} &= \Pr(z_n = k \mid \mathbf{x}_n) \\ &= \frac{\Pr(\mathbf{x}_n \mid z_n = k) \Pr(z_n = k)}{\Pr(\mathbf{x}_n)} \\ &= \frac{\mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \cdot \boldsymbol{\pi}_k}{\sum_{j=1}^K \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j) \cdot \boldsymbol{\pi}_j} \end{split}$$

So, while π_k is the <u>prior</u> probability that $z_n=k$, γ_{nk} is the <u>posterior</u> probability that $z_n=k$ after we have observed \mathbf{x}_n .

Unlike in the case of K-means, where training data points are assigned to a particular cluster and N_k is the number of points in that cluster, for the mixture model, we can think of N_k as the "effective number" of points in cluster k.

Thus, in equation (5), μ_k is a weighted mean of the training data, where the weights are given by the responsibilities, γ_{nk} . A small value of γ_{nk} means that cluster k bears little responsibility for data point \mathbf{x}_n and a large value means that it is very repsonsible for \mathbf{x}_n . (5) is very similar to the cluster update in K-means, exect that rather than hard assignments to each cluster – $r_{nk}=0$ or 1 – each data point has a soft assignment based on γ_{nk} .

(ii). Find Σ_k :

Looking at our handy table of matrix derivatives, we find that for a symmetric $\frac{\partial}{\partial \mathbf{X}}\operatorname{Tr}\left(\mathbf{X}^{-1}\mathbf{A}\right)=-\mathbf{X}^{-1}\mathbf{A}\mathbf{X}^{-1}$. Then, applying product and chain rules, we have

$$\begin{split} \frac{\partial}{\partial \boldsymbol{\Sigma}} \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) &= \frac{\partial}{\partial \boldsymbol{\Sigma}} \left(\frac{1}{(2\pi)^{\frac{D}{2}} |\boldsymbol{\Sigma}|^{\frac{1}{2}}} e^{-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})} \right) \\ &= \frac{1}{(2\pi)^{\frac{D}{2}}} \left(e^{-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})} \frac{\partial}{\partial \boldsymbol{\Sigma}} \frac{1}{|\boldsymbol{\Sigma}|^{\frac{1}{2}}} + \frac{1}{|\boldsymbol{\Sigma}|^{\frac{1}{2}}} \frac{\partial}{\partial \boldsymbol{\Sigma}} e^{-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})} \right) \\ &= \frac{1}{(2\pi)^{\frac{D}{2}}} e^{-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})} \left[\left(-\frac{1}{2} \frac{|\boldsymbol{\Sigma}| \, \boldsymbol{\Sigma}^{-1}}{|\boldsymbol{\Sigma}|^{\frac{3}{2}}} \right) \right. \\ &\left. + \frac{1}{|\boldsymbol{\Sigma}|^{\frac{1}{2}}} \left(-\frac{1}{2} \frac{\partial}{\partial \boldsymbol{\Sigma}} \operatorname{Tr} \left(\boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu}) (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \right) \right) \right] \\ &= -\frac{1}{2} \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) \left(\boldsymbol{\Sigma}^{-1} - \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu}) (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \boldsymbol{\Sigma}^{-1} \right) \end{split}$$

Thus, we have

$$\begin{split} \frac{\partial \mathcal{L}}{\partial \boldsymbol{\Sigma}_k} &= \sum_{n=1}^N \frac{1}{\sum_{j=1}^K \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j) \cdot \boldsymbol{\pi}_j} \cdot \boldsymbol{\pi}_k \cdot \left(\frac{\partial}{\partial \boldsymbol{\Sigma}_k} \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right) \\ &= -\frac{1}{2} \sum_{n=1}^N \gamma_{nk} \left(\boldsymbol{\Sigma}^{-1} - \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu}) (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \boldsymbol{\Sigma}^{-1} \right) \end{split}$$

= 0

Rearranging,

$$\begin{split} \boldsymbol{\Sigma}^{-1} \sum_{n=1}^{N} \gamma_{n\,k} &= \sum_{n=1}^{N} \gamma_{n\,k} (\mathbf{x} - \boldsymbol{\mu}) (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \boldsymbol{\Sigma}^{-1} \\ \boldsymbol{\Sigma}^{-1} N_k &= \boldsymbol{\Sigma}^{-1} \left(\sum_{n=1}^{N} \gamma_{n\,k} (\mathbf{x} - \boldsymbol{\mu}) (\mathbf{x} - \boldsymbol{\mu})^\mathsf{T} \right) \boldsymbol{\Sigma}^{-1} \end{split}$$

Front and back multiplying by the covariance matrix, Σ , we have our result

$$\left[\quad \boldsymbol{\Sigma}_k = \frac{1}{N_k} \sum_{n=1}^N \gamma_{nk} (\mathbf{x}_n - \boldsymbol{\mu}) (\mathbf{x}_n - \boldsymbol{\mu})^\mathsf{T} \quad \right]$$
 (6)

As with the means, this is simply a weighted sample covariance of the training data.

(iii). Find π_k :

Finding a solution for π_k , the prior probilities of mixture component k is slightly more complicated, because of the constraint that the prior probabilities must sum to 1. Rather than a set-the-derivative-equal-to-zero maximization, this is a *constrained* maximization. This can be done with the Lagrange multiplier technique. Instead of maximiz \mathcal{L} , we maximize

$$\mathcal{L}' = \mathcal{L} + \lambda \left(\sum_{k=1}^{K} \pi_k \right),$$

where λ is the Lagrange multiplier who's value we'll discover by enforcing the constraint on the solution.

$$\begin{split} \frac{\partial \mathcal{L}'}{\partial \pi_k} &= \frac{\partial \mathcal{L}}{\partial \pi_k} + \lambda \\ &= \sum_{n=1}^N \frac{1}{\sum_{j=1}^K \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j) \cdot \pi_j} \cdot \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) + \lambda \\ &= \sum_{n=1}^N \frac{\gamma_{nk}}{\pi_k} + \lambda = 0 \end{split}$$

Rearranging and multiplying both sides by π_k ,

$$\begin{split} \sum_{n=1}^{N} \gamma_{n\,k} &= -\lambda \cdot \pi_{k} \\ \Longrightarrow & \pi_{k} = \frac{-N_{k}}{\lambda} \end{split}$$

Now, we can enforce the constraint that $\sum_{k=1}^K \pi_k = 1.$

$$-\frac{\sum_{k=1}^{K} N_k}{\lambda} = 1$$

$$\implies \lambda = -N$$

Thus,

$$\pi_k = \frac{N_k}{N} \tag{7}$$

In other words, the prior probability of each mixture component is the effective fraction of training data for which the component is given responsibility. If we were doing hard assignment, this would just be the number of points assigned to the component.

Wait a second! The maximum likelihood parameters given by (5), (6), and (7) are not actually in closed form! The resposibilities, γnk , appear in the right hand sides of the equations, but depend on the values of the parameters. This suggests an iterative solution – using initial guesses of the parameters to estimate the responsibilities, then recalculating the parameters. This turns out to be an instance of the **Expectation Maximization (EM) Algorithm**.

The EM algorithm is a powerful and general method for optimizing the parameters for models with **latent variables**. A latent variable is a random variable defined as part of generative model that represents something that is not actually observed. In the clustering mixture model, the $\{z_n\}$ are latent variables – unlike the $\{x_n\}$, they are never directly observed.

A.2 EM Algorithm for Gaussian Mixture Model

- **1. Initialize** parameters $\mu_k, \Sigma_k, \pi_k \forall k \in 1, ..., K$. (Also need to decide on K!)
- 2. E-step: Evaluate responsibilities given current parameter values

$$\gamma_{nk} = \frac{\mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}$$

3. M-step: Re-estimate parameters using the current values of the responsibilities

$$\begin{split} & \boldsymbol{\mu}_k^{\text{new}} = \frac{1}{N_k} \sum_{n=1}^N \gamma_{nk} \mathbf{x}_n \\ & \boldsymbol{\Sigma}_k^{\text{new}} = \frac{1}{N_k} \sum_{n=1}^N \gamma_{nk} (\mathbf{x}_n - \boldsymbol{\mu}^{\text{new}}) (\mathbf{x}_n - \boldsymbol{\mu}^{\text{new}})^{\text{T}} \\ & \boldsymbol{\pi}_k^{\text{new}} = \frac{N_k}{N} \end{split}$$

where
$$N_k = \sum_{n=1}^N \gamma_{nk}$$
.

4. Evaluate the log likelihood of training data:

$$\log \Pr(\{\mathbf{x}_n\} \mid \boldsymbol{\theta}) = \sum_{n=1}^N \log \left[\sum_{k=1}^K \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \cdot \boldsymbol{\pi}_k \right]$$

in order to see whether it has converged. (Alternatively, one can track the convergence of the parameters after step 3.) If the convergence criteria are not satisfied go to step 2.

After each iteration of the EM algorithm the log likelihood of the training data, log $Pr(\{\mathbf{x}_n\} \mid \theta)$, increases (meaning that the parameter estimates are improving).

Mixtures of Exponential Family Distributions: We have calculated the M-step parameter updates for clustering using a mixture of Gaussians model. The solution ends up being weighted versions of the typical maximum-likelihood parameter estimates for μ_k and Σ_k . Here's a conjecture

Conjecture. For a mixture model defined using an distribution from the exponential family, the parameter updates in the M-step of the EM algorithm will take the form of weighted averages of the sample estimates of these parameters.

Here's the outline of a proof.

Proof. If we consider a distribution in the very general exponential family, we can define

$$\Pr(\mathbf{x} \mid z = k) = f(\mathbf{x} \mid \boldsymbol{\eta}_k) = h(\mathbf{x}) \exp\left(\boldsymbol{\eta}_k^{\mathsf{T}} T(\mathbf{x}) - A(\boldsymbol{\eta}_k)\right) \tag{8}$$

where η_k are the so-called "natural parameters" of the distribution $f(\mathbf{x})$ for component k. The typical parameters that we think of may not be the natural ones, but can be simply derived from them. The functions h(), T(), and A() are determined by the type of distribution. If we have some data $\{\mathbf{x}_n\}$, for a general exponential family distribution, the maximum likelihood estimates for the natural parameters are the solution to the equation

$$\frac{\partial}{\partial \pmb{\eta}} A(\pmb{\eta}) = \frac{\sum_{n=1}^{N} T(\mathbf{x}_n)}{N}.$$

So now let's consider our mixture model.

$$\frac{\partial \mathcal{L}}{\partial \boldsymbol{\eta}_k} = \sum_{n=1}^N \frac{1}{\sum_{j=1}^K f(\mathbf{x} \mid \boldsymbol{\eta}_j)} \cdot \boldsymbol{\pi}_k \cdot \left(\frac{\partial}{\partial \boldsymbol{\eta}_k} f(\mathbf{x} \mid \boldsymbol{\eta}_k) \right) \tag{9}$$

From (8), we can see that

$$\frac{\partial}{\partial \boldsymbol{\eta}} f(\mathbf{x} \mid \boldsymbol{\eta}) = f(\mathbf{x}) \left(T(\mathbf{x}) - \frac{\partial}{\partial \boldsymbol{\eta}} A(\boldsymbol{\eta}) \right)$$

Plugging this in, we have

$$\frac{\partial \mathcal{L}}{\partial \boldsymbol{\eta}_k} = \sum_{n=1}^N \gamma_{nk} \left(T(\mathbf{x}) - \frac{\partial}{\partial \boldsymbol{\eta}_k} A(\boldsymbol{\eta}_k) \right) = 0 \tag{10}$$

$$\implies \frac{\partial}{\partial \boldsymbol{\eta}_k} A(\boldsymbol{\eta}_k) = \frac{1}{N_k} \sum_{n=1}^N \gamma_{nk} T(\mathbf{x}). \tag{11}$$

Notice that this has the same form as the general maximum likelihood parameter estimate, but weighted by the responsibilities γ_{nk} .