

STA 414/2104: Machine Learning

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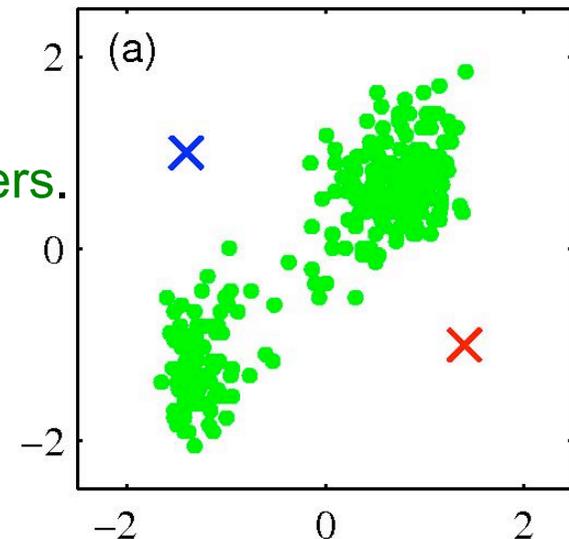
Lecture 7

Mixture Models

- We will look at the mixture models, including **Gaussian mixture** models.
- The key idea is to introduce **latent variables**, which allows complicated distributions to be formed from simpler distributions.
- We will see that mixture models can be interpreted in terms of having **discrete latent variables** (in a directed graphical model).
- Later in class, we will also look at the continuous latent variables.

K-Means Clustering

- Let us first look at the following problem: **Identify clusters**, or groups, of data points in a multidimensional space.
- We observe the dataset $\{\mathbf{x}_1, \dots, \mathbf{x}_N\}$ consisting of N D -dimensional observations
- We would like to **partition the data into K clusters**, where K is given.
- We next introduce D -dimensional vectors, **prototypes**, $\mu_k, k = 1, \dots, K$.
- We can think of μ_k as representing cluster centers.
- Our goal:
 - Find an **assignment of data points to clusters**.
 - Sum of squared distances of each data point to its closest prototype is **at the minimum**.



K-Means Clustering

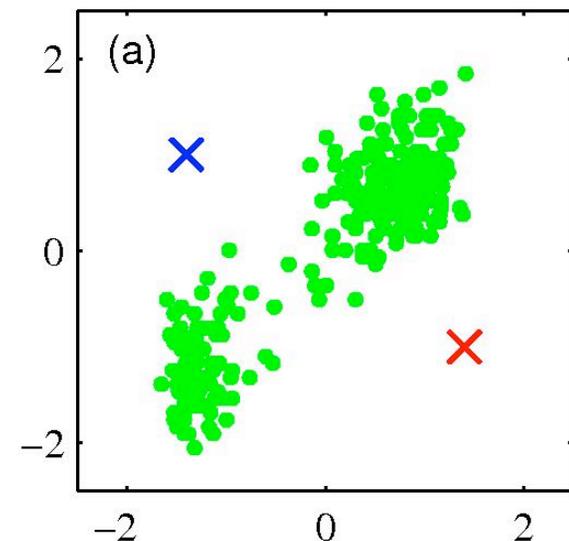
- For each data point \mathbf{x}_n we introduce a binary vector \mathbf{r}_n of length K (1-of- K encoding), which indicates which of the K clusters the data point \mathbf{x}_n is assigned to.

- Define objective (distortion measure):

$$J = \sum_{n=1}^N \sum_{k=1}^K r_{nk} \|\mathbf{x}_n - \boldsymbol{\mu}_k\|^2.$$

- It represents the **sum of squares of the distances** of each data point to its assigned prototype $\boldsymbol{\mu}_k$.

- Our goal is to find the values of r_{nk} and the cluster centers $\boldsymbol{\mu}_k$ so as to minimize the objective J .



Iterative Algorithm

- Define iterative procedure to minimize:

$$J = \sum_{n=1}^N \sum_{k=1}^K r_{nk} \|\mathbf{x}_n - \boldsymbol{\mu}_k\|^2.$$

- Given $\boldsymbol{\mu}_k$, minimize J with respect to r_{nk} (**E-step**):

$$r_{nk} = \begin{cases} 1 & \text{if } k = \arg \min_j \|\mathbf{x}_n - \boldsymbol{\mu}_j\|^2 \\ 0 & \text{otherwise} \end{cases}$$

Hard assignments of points to clusters.



which simply says **assign n^{th} data point \mathbf{x}_n to its closest cluster center.**

- Given r_{nk} , minimize J with respect to $\boldsymbol{\mu}_k$ (**M-step**):

$$\boldsymbol{\mu}_k = \frac{\sum_n r_{nk} \mathbf{x}_n}{\sum_n r_{nk}}.$$

Number of points assigned to cluster k.

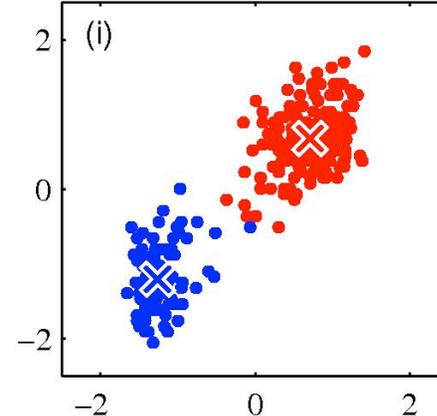
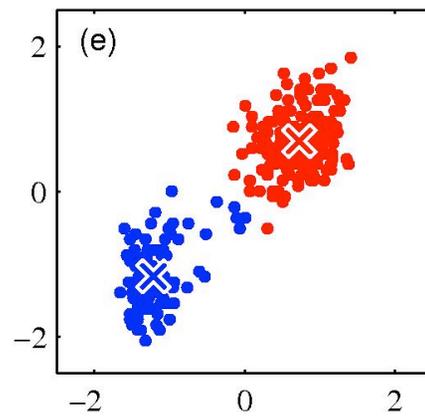
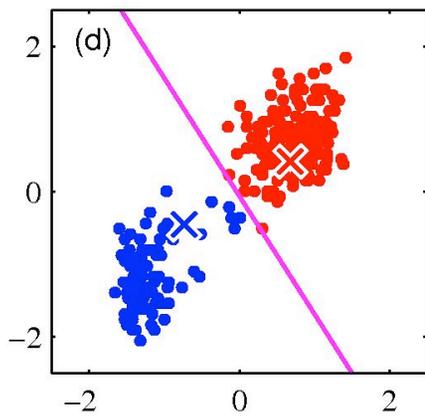
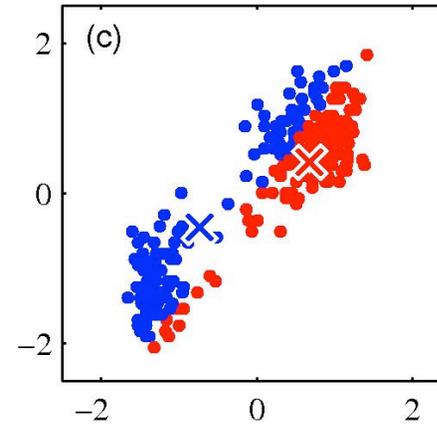
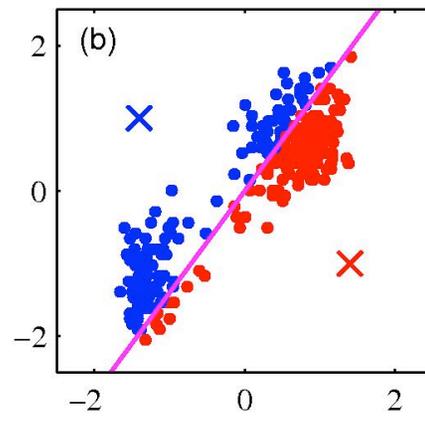
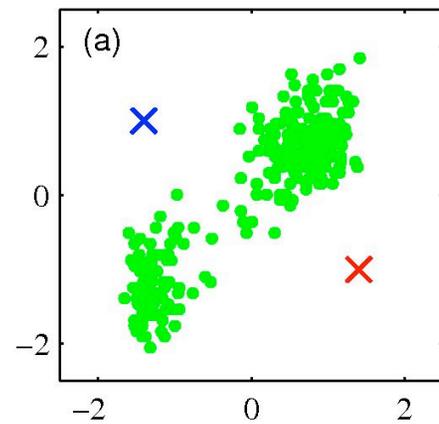


Set $\boldsymbol{\mu}_k$ equal to the **mean of all the data points assigned to cluster k.**

- Guaranteed convergence to local minimum (**not global minimum**).

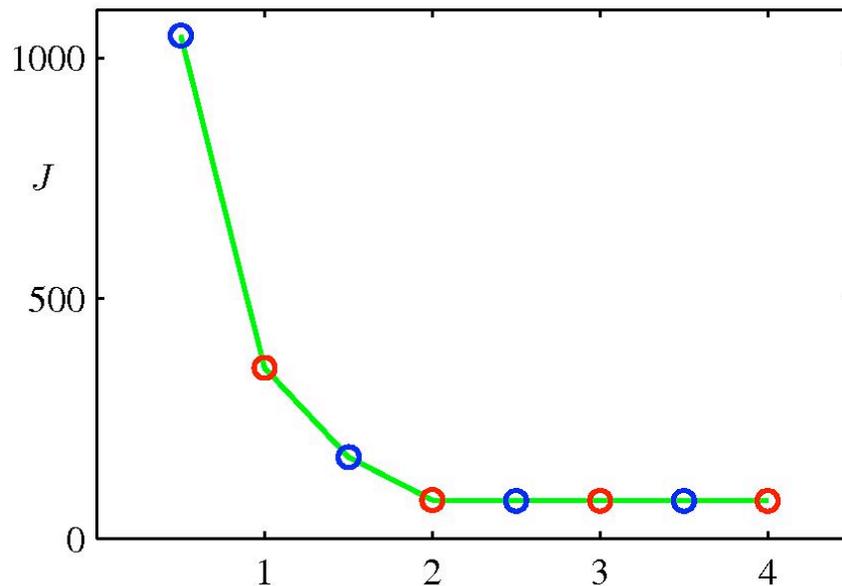
Example

- Example of using K-means (K=2) on Old Faithful dataset.



Convergence

- Plot of the cost function after each E-step (blue points) and M-step (red points)



The algorithm has converged after 3 iterations.

- K-means can be generalized by introducing a **more general dissimilarity measure**:

$$J = \sum_{n=1}^N \sum_{k=1}^K r_{nk} K(\mathbf{x}_n, \boldsymbol{\mu}_k).$$

Image Segmentation

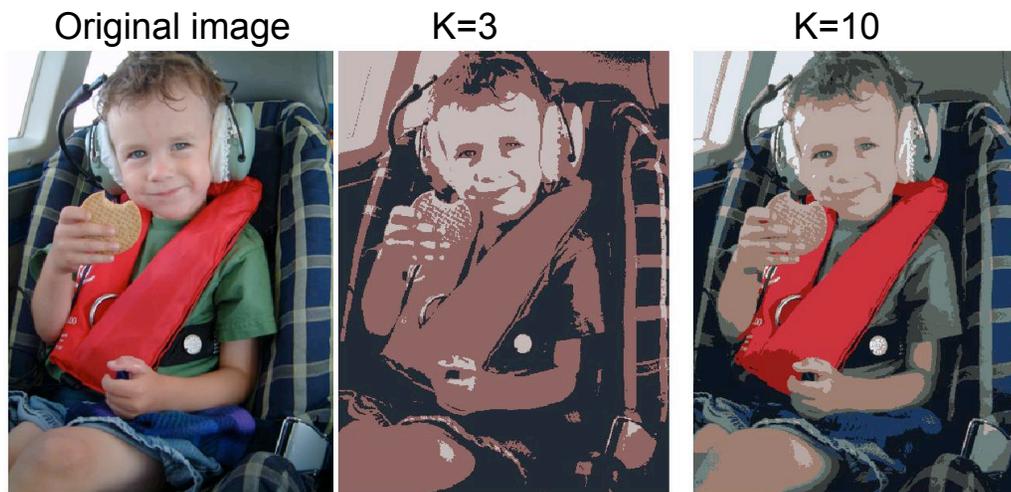
- Another application of K-means algorithm.
- **Partition an image into regions** corresponding, for example, to object parts.
- Each pixel in an image is a point in 3-D space, **corresponding to R,G,B channels.**



- For a given value of K , the algorithm represent an image using K colors.
- Another application is image compression.

Image Compression

- For each data point, we store only the **identity k of the assigned cluster**.
- We also **store the values of the cluster centers μ_k** .
- Provided $K \ll N$, we require significantly less data.



- The original image has $240 \times 180 = 43,200$ pixels.
- Each pixel contains $\{R,G,B\}$ values, each of which requires 8 bits.

- Requires $43,200 \times 24 = 1,036,800$ bits to transmit directly.
- With K-means, we need to transmit K **code-book vectors μ_k** -- $24K$ bits.
- For each pixel we need to transmit **$\log_2 K$ bits** (as there are K vectors).
- **Compressed image** requires 43,248 (K=2), 86,472 (K=3), and 173,040 (K=10) bits, which amounts to compression ratios of 4.2%, 8.3%, and 16.7%.

Mixture of Gaussians

- We will look at mixture of Gaussians in terms of **discrete latent variables**.
- The Gaussian mixture can be written as a linear **superposition of Gaussians**:

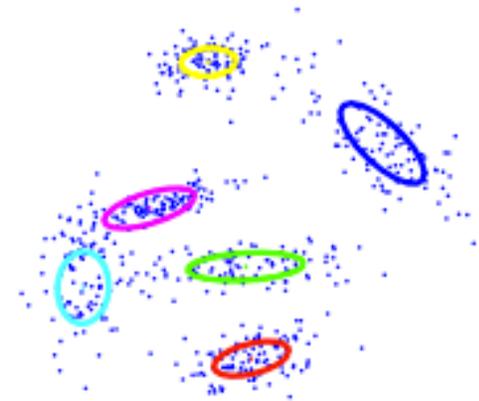
$$p(\mathbf{x}) = \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

- Introduce K-dimensional **binary random variable** \mathbf{z} having a 1-of-K representation:

$$z_k \in \{0, 1\}, \quad \sum_k z_k = 1.$$

- We will specify the distribution over \mathbf{z} in terms of mixing coefficients:

$$p(z_k = 1) = \pi_k, \quad 0 \leq \pi_k \leq 1, \quad \sum_k \pi_k = 1.$$



Mixture of Gaussians

- Because \mathbf{z} uses **1-of-K encoding**, we have:

$$p(\mathbf{z}) = \prod_{k=1}^K \pi_k^{z_k}.$$

- We can now specify the conditional distribution:

$$p(\mathbf{x}|z_k = 1) = \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k), \text{ or } p(\mathbf{x}|\mathbf{z}) = \prod_{k=1}^K \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)^{z_k}.$$

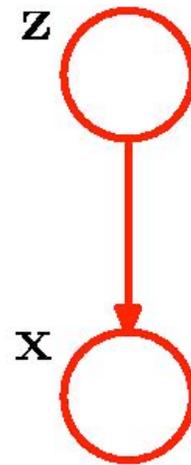
- We have therefore specified the joint distribution:

$$p(\mathbf{x}, \mathbf{z}) = p(\mathbf{x}|\mathbf{z})p(\mathbf{z}).$$

- The **marginal distribution** over \mathbf{x} is given by:

$$p(\mathbf{x}) = \sum_{\mathbf{z}} p(\mathbf{z})p(\mathbf{x}|\mathbf{z}) = \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

- The marginal distribution over \mathbf{x} is given by a **Gaussian mixture**.



Mixture of Gaussians

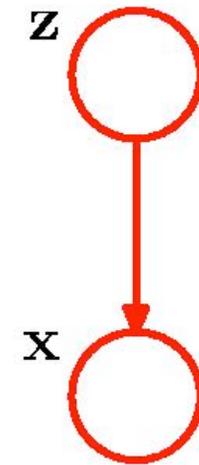
- The marginal distribution:

$$p(\mathbf{x}) = \sum_{\mathbf{z}} p(\mathbf{z})p(\mathbf{x}|\mathbf{z}) = \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

- If we have several observations $\mathbf{x}_1, \dots, \mathbf{x}_N$, it follows that for **every observed data point** \mathbf{x}_n , there is a corresponding **latent variable** \mathbf{z}_n .
- Let us look at the conditional $p(\mathbf{z}|\mathbf{x})$, responsibilities, which we will need for doing inference:

$$\gamma(z_k) = p(z_k = 1|\mathbf{x}) = \frac{p(z_k = 1)p(\mathbf{x}|z_k = 1)}{\sum_{j=1}^K p(z_j = 1)p(\mathbf{x}|z_j = 1)} = \frac{\pi_k N(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j N(\mathbf{x}|\boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}.$$

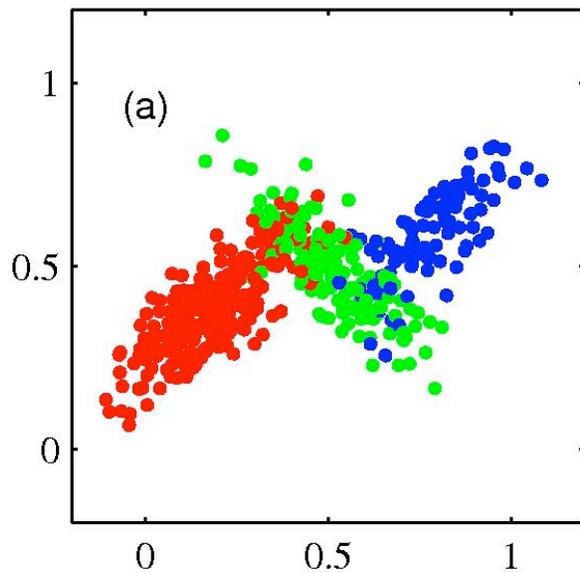
responsibility that component k takes for explaining the data x



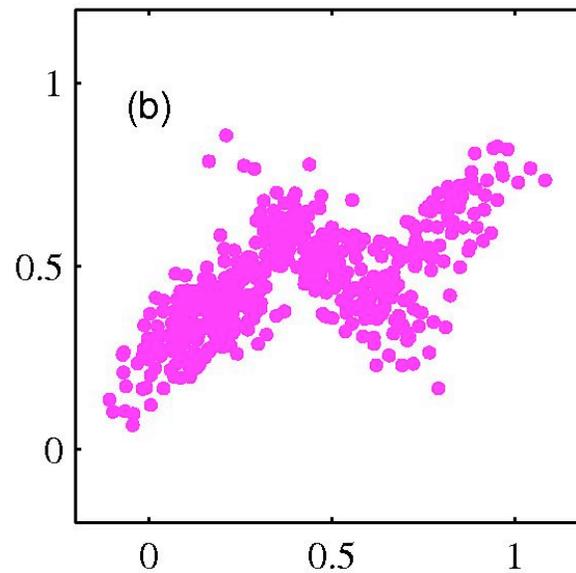
- We will view π_k as **prior probability** that $z_k=1$, and $\gamma(z_k)$ is the **corresponding posterior** once we have observed the data.

Example

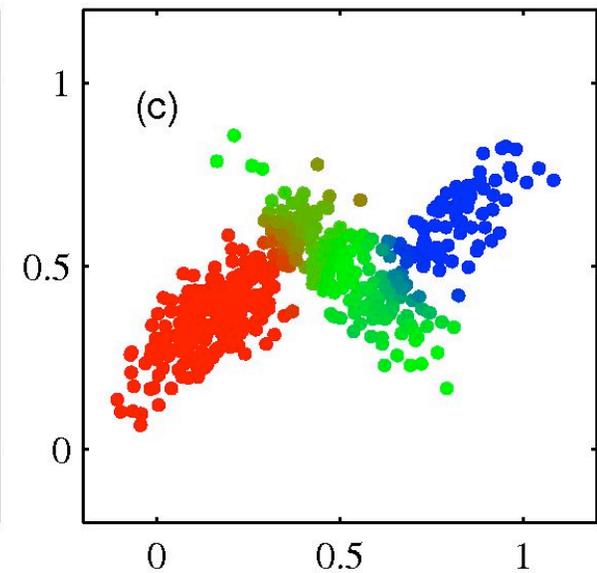
- 500 points drawn from a mixture of 3 Gaussians.



Samples from the **joint distribution** $p(\mathbf{x}, \mathbf{z})$.



Samples from the **marginal distribution** $p(\mathbf{x})$.



Same samples where colors represent the value of responsibilities.

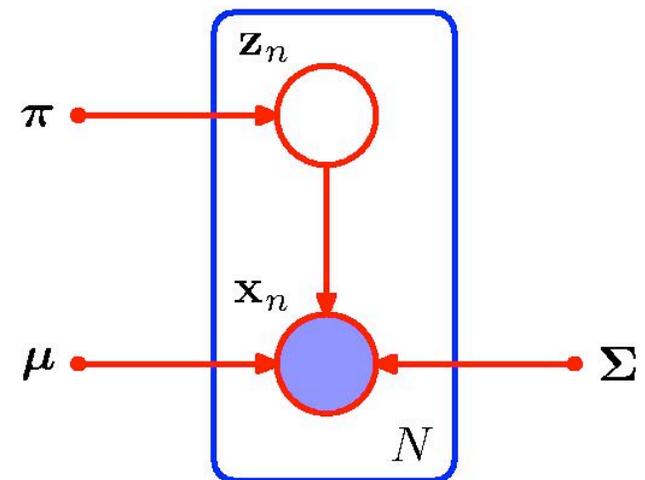
Maximum Likelihood

- Suppose we observe a dataset $\{\mathbf{x}_1, \dots, \mathbf{x}_N\}$, and we model the data using mixture of Gaussians.
- We represent the dataset as an N by D matrix \mathbf{X} .
- The corresponding **latent variables** will be represented and an N by K matrix \mathbf{Z} .

- The log-likelihood takes form:

$$\ln p(\mathbf{X} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \sum_{n=1}^N \ln \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

↑
Model parameters



Graphical model for a Gaussian mixture model for a set of i.i.d. data point $\{\mathbf{x}_n\}$, and corresponding latent variables $\{\mathbf{z}_n\}$.

Maximum Likelihood

- The log-likelihood:

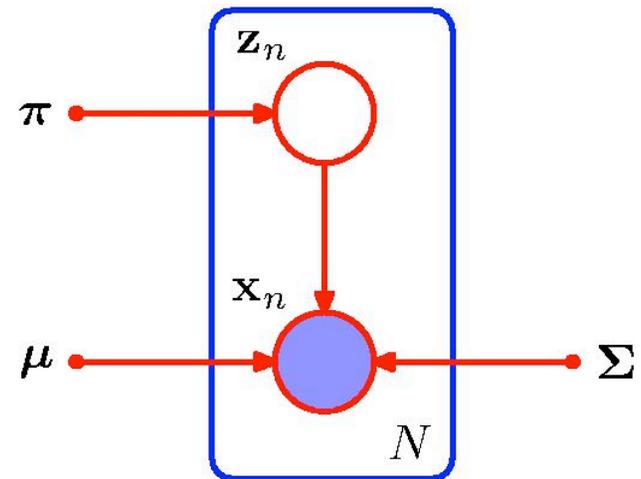
$$\ln p(\mathbf{X}|\boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \sum_{n=1}^N \ln \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

- Differentiating with respect to $\boldsymbol{\mu}_k$ and setting to zero:

$$0 = \sum_n \frac{\pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_j \pi_j \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)} \boldsymbol{\Sigma}_K^{-1} (\mathbf{x}_n - \boldsymbol{\mu}_k).$$

$\gamma(z_{nk})$ Soft assignment

$$\boldsymbol{\mu}_k = \frac{1}{N_k} \sum_n \gamma(z_{nk}) \mathbf{x}_n, \quad N_k = \sum_n \gamma(z_{nk}).$$



- We can interpret N_k as **effective number of points** assigned to cluster k .
- The mean $\boldsymbol{\mu}_k$ is given by the mean of all the data points **weighted by the posterior** $\gamma(z_{nk})$ that component k was responsible for generating \mathbf{x}_n .

Maximum Likelihood

- The log-likelihood:

$$\ln p(\mathbf{X}|\boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \sum_{n=1}^N \ln \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

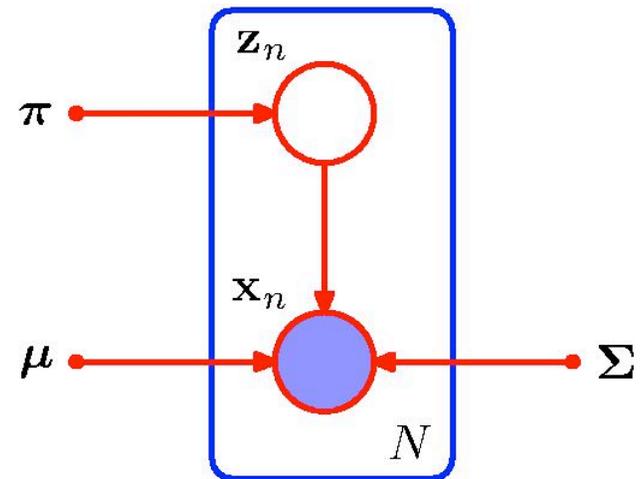
- Differentiating with respect to $\boldsymbol{\Sigma}_k$ and setting to zero:

$$\boldsymbol{\Sigma}_k = \frac{1}{N_k} \sum_{n=1}^N \gamma(z_{nk}) (\mathbf{x}_n - \boldsymbol{\mu}_k)(\mathbf{x}_n - \boldsymbol{\mu}_k)^T.$$

- Note that the data points are **weighted by the posterior probabilities**.
- Maximizing log-likelihood with respect to mixing proportions:

$$\pi_k = \frac{N_k}{N}.$$

- Mixing proportion for the k^{th} component is given by the **average responsibility which that component takes for explaining the data**.



Maximum Likelihood

- The log-likelihood:

$$\ln p(\mathbf{X}|\boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \sum_{n=1}^N \ln \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

- Note that the maximum likelihood **does not have a closed form solution**.
- Parameter updates **depend on responsibilities**

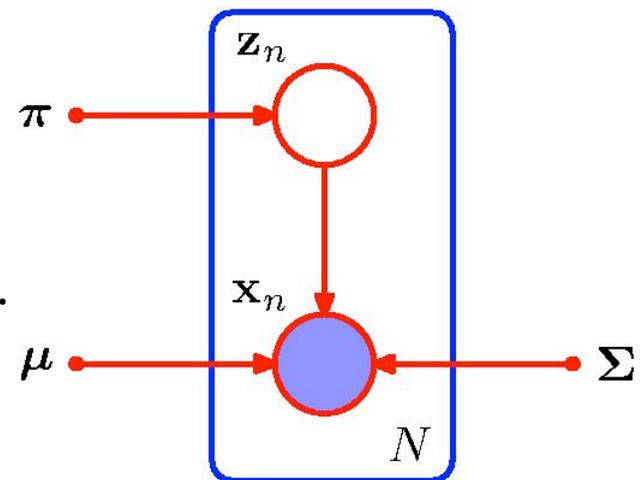
$\gamma(z_{nk})$, which themselves depend on those parameters:

$$\gamma(z_{nk}) = p(z_{nk} = 1 | \mathbf{x}) = \frac{\pi_k N(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j N(\mathbf{x}_n | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}.$$

- Iterative Solution:

E-step: Update responsibilities $\gamma(z_{nk})$.

M-step: Update model parameters $\pi_k, \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k$, for $k=1, \dots, K$.



EM algorithm

- Initialize the means μ_k , covariances Σ_k , and mixing proportions π_k .
- **E-step: Evaluate responsibilities** using current parameter values:

$$\gamma(z_{nk}) = p(z_{nk} = 1 | \mathbf{x}) = \frac{\pi_k N(\mathbf{x}_n | \mu_k, \Sigma_k)}{\sum_{j=1}^K \pi_j N(\mathbf{x}_n | \mu_j, \Sigma_j)}.$$

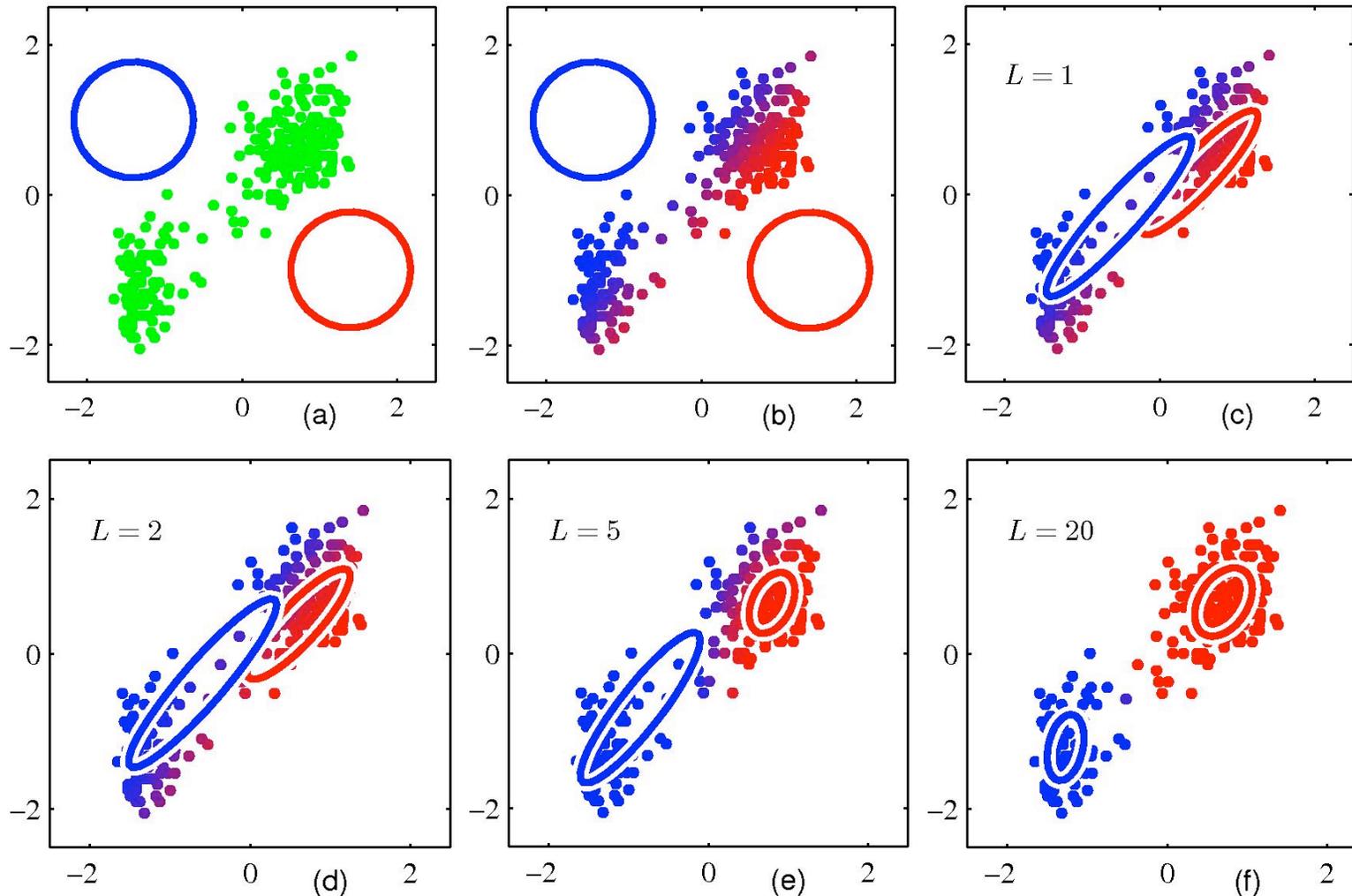
- **M-step: Re-estimate model parameters** using the current responsibilities:

$$\begin{aligned}\mu_k^{new} &= \frac{1}{N_k} \sum_n \gamma(z_{nk}) \mathbf{x}_n, & N_k &= \sum_n \gamma(z_{nk}), \\ \Sigma_k^{new} &= \frac{1}{N_k} \sum_{n=1}^N \gamma(y_{nk}) (\mathbf{x}_n - \mu_k)(\mathbf{x}_n - \mu_k)^T, \\ \pi_k^{new} &= \frac{N_k}{N}.\end{aligned}$$

- Evaluate the log-likelihood and **check for convergence**.

Mixture of Gaussians: Example

- Illustration of the EM algorithm (much slower convergence compared to K-means)



An Alternative View of EM

- The goal of EM is to **find maximum likelihood solutions** for models with latent variables.
- We represent the **observed dataset** as an N by D matrix \mathbf{X} .
- **Latent variables** will be represented and an N by K matrix \mathbf{Z} .
- The set of all **model parameters** is denoted by θ .
- The log-likelihood takes form:

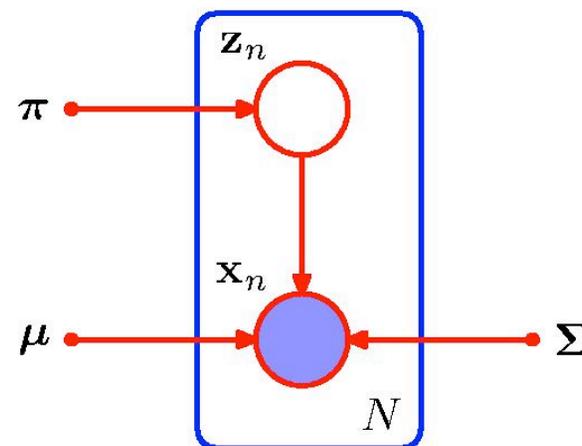
$$\ln p(\mathbf{X}|\theta) = \ln \left[\sum_{\mathbf{Z}} p(\mathbf{X}, \mathbf{Z}|\theta) \right].$$

- Note: even if the joint distribution belongs to exponential family, the marginal typically does not!

- We will call:

$\{\mathbf{X}, \mathbf{Z}\}$ as **complete** dataset.

$\{\mathbf{X}\}$ as **incomplete** dataset.



An Alternative View of EM

- In practice, we are **not given a complete dataset** $\{\mathbf{X}, \mathbf{Z}\}$, but only incomplete dataset $\{\mathbf{X}\}$.
- Our knowledge about the latent variables is given only by **the posterior distribution** $p(\mathbf{Z}|\mathbf{X}, \theta)$.
- Because we cannot use the complete data log-likelihood, we can consider **expected complete-data log-likelihood**:

$$Q(\theta, \theta^{old}) = \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \theta^{old}) \ln p(\mathbf{X}, \mathbf{Z}|\theta).$$

- In the E-step, we use the current parameters θ^{old} to compute **the posterior over the latent variables** $p(\mathbf{Z}|\mathbf{X}, \theta^{old})$.
- We use this posterior to compute expected complete log-likelihood.
- In the M-step, we find the revised parameter estimate θ^{new} by **maximizing the expected complete log-likelihood**:

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

← Tractable

The General EM algorithm

- Given a joint distribution $p(\mathbf{Z}, \mathbf{X}|\theta)$ over observed and latent variables governed by parameters θ , the goal is to maximize the likelihood function $p(\mathbf{X}|\theta)$ with respect to θ .
- Initialize parameters θ^{old} .
- **E-step**: Compute posterior over latent variables: $p(\mathbf{Z}|\mathbf{X}, \theta^{old})$.
- **M-step**: Find the new estimate of parameters θ^{new} :

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

where

$$Q(\theta, \theta^{old}) = \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \theta^{old}) \ln p(\mathbf{X}, \mathbf{Z}|\theta).$$

- Check for convergence of either log-likelihood or the parameter values.

Otherwise:

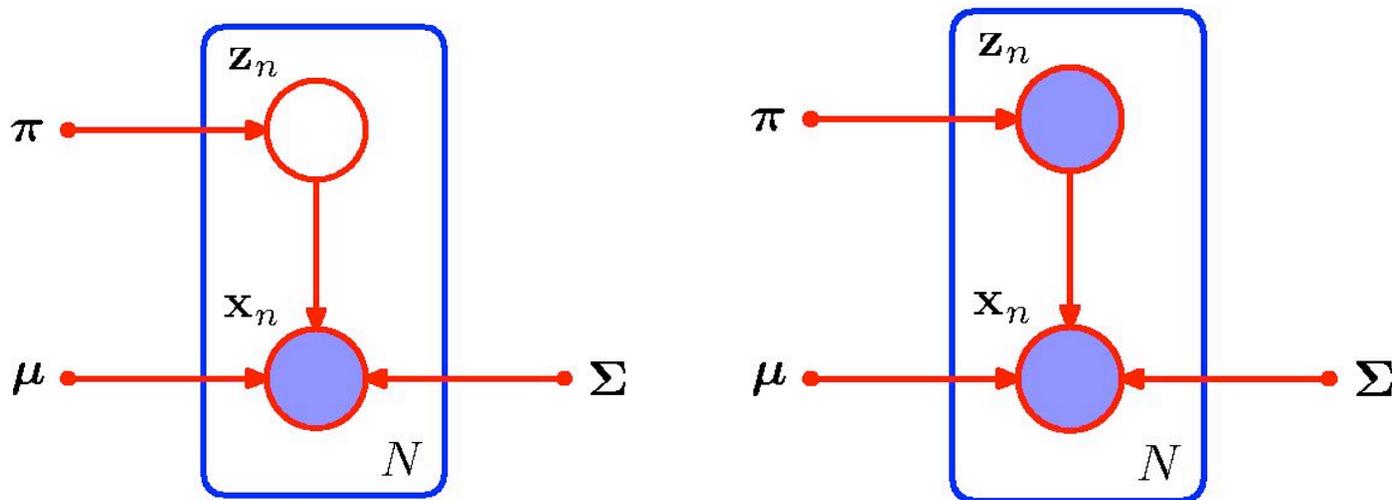
$$\theta^{new} \leftarrow \theta^{old}, \quad \text{and iterate.}$$

Gaussian Mixtures Revisited

- We now consider the application of the latent variable view of EM the case of **Gaussian mixture model**.

- Recall:

$$\ln p(\mathbf{X}|\boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \sum_{n=1}^N \ln \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$



$\{\mathbf{X}\}$ -- incomplete dataset. $\{\mathbf{X}, \mathbf{Z}\}$ -- complete dataset.

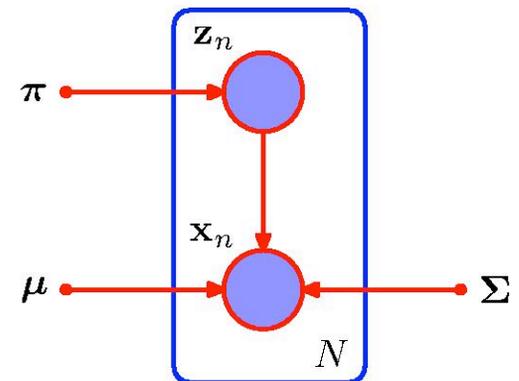
Maximizing Complete Data

- Consider the problem of maximizing the likelihood for the complete data:

$$p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \prod_{n=1}^N \prod_{k=1}^K \left[\pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right]^{z_{nk}}.$$

$$\ln p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \sum_{k=1}^K \left[\sum_{n=1}^N z_{nk} \ln \pi_k + z_{nk} \ln \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right].$$

Sum of K independent contributions, one for each mixture component.



$\{\mathbf{X}, \mathbf{Z}\}$

-- complete dataset.

- Maximizing with respect to **mixing proportions** yields:

$$\pi_k = \frac{1}{N} \sum_{n=1}^N z_{nk}.$$

- And similarly for the means and covariances.

Posterior Over Latent Variables

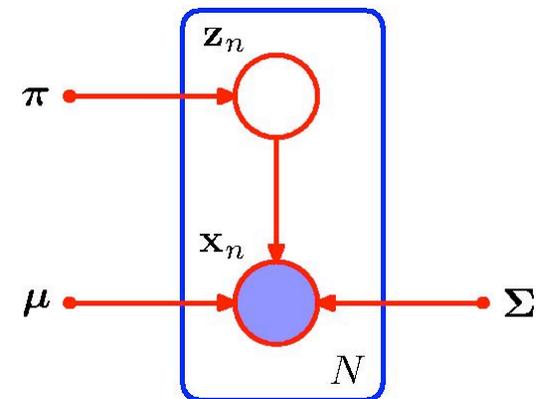
- Remember:

$$p(\mathbf{x}|\mathbf{z}) = \prod_{k=1}^K \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)^{z_k}, \quad p(\mathbf{z}) = \prod_{k=1}^K \pi_k^{z_k}.$$

- The **posterior over latent variables** takes form:

$$p(\mathbf{Z}|\mathbf{X}, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) \propto \prod_{n=1}^N \prod_{k=1}^K \left[\pi_k \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right]^{z_k}.$$

- Note that the **posterior factorizes over n points**, so that under the posterior distribution $\{\mathbf{z}_n\}$ are independent.



Expected Complete Log-Likelihood

- The expected value of indicator variable z_{nk} under the posterior distribution is:

$$\begin{aligned}\mathbb{E}[z_{nk}] &= \frac{\sum_{\mathbf{z}_n} z_{nk} \prod_j [\pi_j \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)]^{z_{nj}}}{\sum_{\mathbf{z}_n} \prod_j [\pi_j \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)]^{z_{nj}}} \\ &= \frac{\pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)} = \gamma(z_{nk}).\end{aligned}$$

- This represent **the responsibility** of component k for data point \mathbf{x}_n .
- The **complete-data log-likelihood**:

$$\ln p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \sum_{n=1}^N \sum_{k=1}^K z_{nk} \left[\ln \pi_k + \ln \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right].$$

- The **expected complete data log-likelihood** is:

$$\mathbb{E}_{\mathbf{Z}} \left[\ln p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) \right] = \sum_{n=1}^N \sum_{k=1}^K \gamma(z_{nk}) \left[\ln \pi_k + \ln \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right].$$

Expected Complete Log-Likelihood

- The expected complete data log-likelihood is:

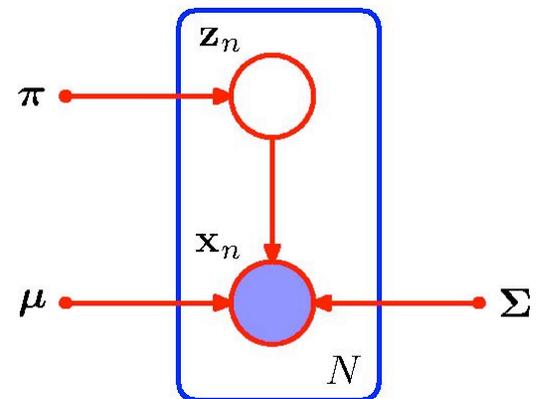
$$\mathbb{E}_{\mathbf{Z}} [\ln p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma})] = \sum_{n=1}^N \sum_{k=1}^K \gamma(z_{nk}) \left[\ln \pi_k + \ln \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right].$$

- Maximizing the respect to model parameters we obtain:

$$\boldsymbol{\mu}_k^{new} = \frac{1}{N_k} \sum_n \gamma(z_{nk}) \mathbf{x}_n, \quad N_k = \sum_n \gamma(z_{nk}),$$

$$\boldsymbol{\Sigma}_k^{new} = \frac{1}{N_k} \sum_{n=1}^N \gamma(y_{nk}) (\mathbf{x}_n - \boldsymbol{\mu}_k) (\mathbf{x}_n - \boldsymbol{\mu}_k)^T,$$

$$\pi_k^{new} = \frac{N_k}{N}.$$



Relationship to K-Means

- Consider a Gaussian mixture model in which **covariances are shared** and are given by ϵI .

$$p(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) = \frac{1}{(2\pi\epsilon)^{D/2}} \exp\left[-\frac{1}{2\epsilon}\|\mathbf{x} - \boldsymbol{\mu}_k\|^2\right].$$

- Consider EM algorithm for a mixture of K Gaussians, in which **we treat ϵ as a fixed constant**. The **posterior responsibilities** take form:

$$\gamma(z_{nk}) = \frac{\pi_k \exp(-\|\mathbf{x}_n - \boldsymbol{\mu}_k\|^2/2\epsilon)}{\sum_{j=1}^K \pi_j \exp(-\|\mathbf{x}_n - \boldsymbol{\mu}_j\|^2/2\epsilon)}.$$

- Consider the limit $\epsilon \rightarrow 0$.
- In the denominator, the term for which $\|\mathbf{x}_n - \boldsymbol{\mu}_j\|^2$ is smallest will go to zero **most slowly**. Hence $\gamma(z_{nk}) \rightarrow r_{nk}$, where

$$r_{nk} = \begin{cases} 1 & \text{if } k = \arg \min_j \|\mathbf{x}_n - \boldsymbol{\mu}_j\|^2 \\ 0 & \text{otherwise} \end{cases}$$

Relationship to K-Means

- Consider EM algorithm for a mixture of K Gaussians, in which we treat ϵ as a fixed constant. The posterior responsibilities take form:

$$\gamma(z_{nk}) = \frac{\pi_k \exp(-\|\mathbf{x}_n - \boldsymbol{\mu}_k\|^2/2\epsilon)}{\sum_{j=1}^K \pi_j \exp(-\|\mathbf{x}_n - \boldsymbol{\mu}_j\|^2/2\epsilon)}.$$

- Finally, in the limit $\epsilon \rightarrow 0$, the expected complete log-likelihood becomes:

$$\mathbb{E}_{\mathbf{Z}} [\ln p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma})] \rightarrow -\frac{1}{2} \sum_{n=1}^N \sum_{k=1}^K r_{nk} \|\mathbf{x}_n - \boldsymbol{\mu}_k\|^2 + \text{const.}$$

- Hence in the limit, maximizing the expected complete log-likelihood is equivalent to minimizing the distortion measure J for the K-means algorithm.