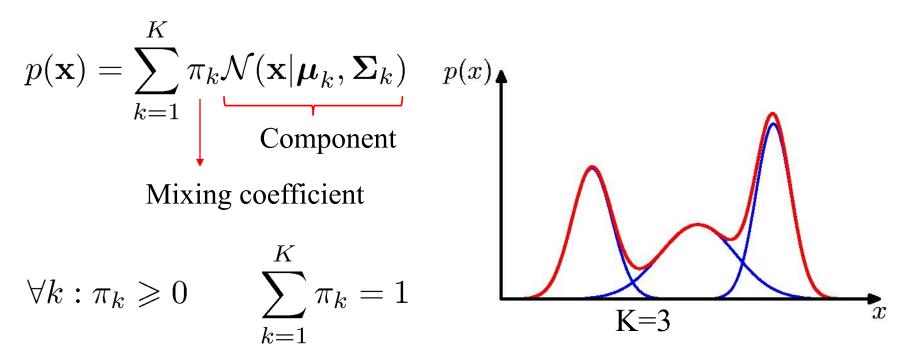
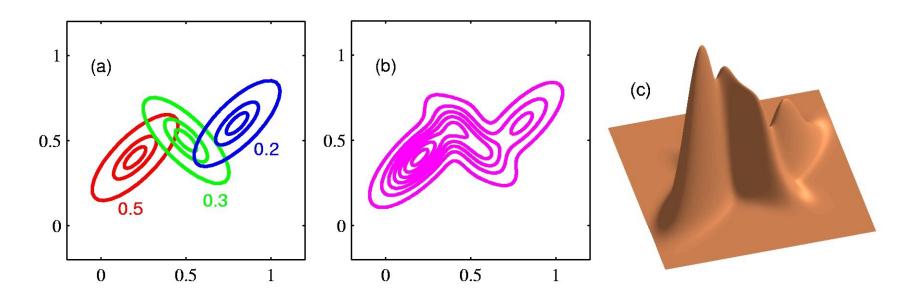
• We combine simple models into a complex model by defining a superposition of K Gaussian densities of the form:



- Note that each Gaussian component has its own mean and covariance. The parameters  $\pi_k$  are called mixing coefficients.
- More generally, mixture models can comprise linear combinations of other distributions.

• Illustration of a mixture of 3 Gaussians in a 2-dimensional space:



(a) Contours of constant density of each of the mixture components, along with the mixing coefficients

(b) Contours of marginal probability density  $p(\mathbf{x}) = \sum_{k=1}^{N} \pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$ 

(c) A surface plot of the distribution p(x).

- We will look at mixture of Gaussians in terms of discrete latent variables.
- The Gaussian mixture:

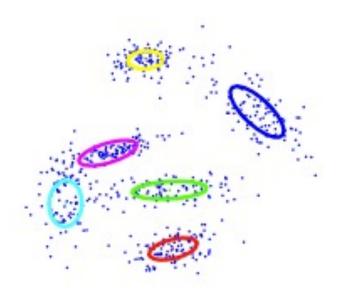
$$p(\mathbf{x}) = \sum_{k=1}^{K} \pi_k \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_K).$$

•K-dimensional binary random variable **z** having a 1-of-K representation is the latent variable:

$$\mathbf{z} = [0, 0, ..., 1, 0]^T$$
  $z_k \in \{0, 1\}, \sum_k z_k = 1.$ 

• We will specify the distribution over **z** in terms of mixing coefficients:

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

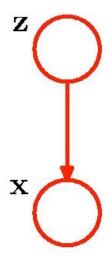


• Because **z** uses 1-of-K encoding, we have:

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

• We can now specify the conditional distribution:

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



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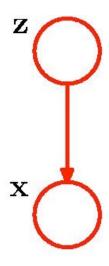
• We have therefore specified the joint distribution:

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

• The marginal distribution over **x** is given by:

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

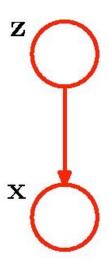
• The marginal distribution over **x** is given by a Gaussian mixture.



• 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  $x_1,...,x_N$ , it follows that for every observed data point  $x_n$ , there is a corresponding latent variable  $z_n$ .



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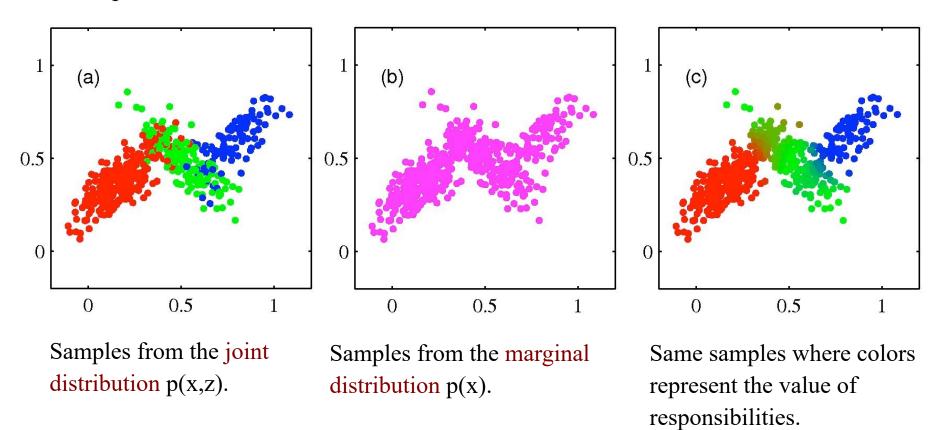
- If we have several observations  $x_1,...,x_N$ , it follows that for every observed data point  $x_n$ , there is a corresponding latent variable  $z_n$ .
- Let us look at the conditional  $p(\mathbf{z} \mid \mathbf{x})$ , "responsibilities", which we will need for doing inference:

$$\gamma(z_k) = p(z_k = 1 | \mathbf{x}) = \frac{p(z_k = 1)p(\mathbf{x} | z_k = 1)}{\sum_{j=1}^{K} p(z_j = 1)p(\mathbf{x} | z_j = 1)} = \frac{responsibility that}{responsibility that component k takes for explaining the data x} = \frac{\pi_k N(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^{K} \pi_j N(\mathbf{x} | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}.$$

• We will view  $\pi_k$  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.



- Suppose we observe a dataset  $\{x_1,...,x_N\}$ , and we model the data using mixture of Gaussians.
- We represent the dataset as an N by D matrix **X**.
- The corresponding latent variables will be represented as an N by K matrix **Z**.
- The log-likelihood takes form:

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

Model parameters

Graphical model for a Gaussian mixture model for a set of i.i.d. data point  $\{x_n\}$ , and corresponding latent variables  $\{z_n\}$ .

• The log-likelihood takes form:

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

• Differentiating with respect to  $\mu_k$  and setting to zero:

$$0 = \sum_{n} \frac{\pi_{k} \mathcal{N}(\mathbf{x}_{n} | \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k})}{\sum_{j} \pi_{j} \mathcal{N}(\mathbf{x}_{n} | \boldsymbol{\mu}_{j}, \boldsymbol{\Sigma}_{j})} \sum_{k=1}^{n-1} (\mathbf{x}_{n} - \boldsymbol{\mu}_{k}). \qquad \boldsymbol{\pi}$$

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

$$\boldsymbol{\mu}_{k} = \frac{1}{N_{k}} \sum_{n} \gamma(z_{nk}) \mathbf{x}_{n}, \quad N_{k} = \sum_{n} \gamma(z_{nk}).$$

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

• The log-likelihood takes form:

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

• Differentiating with respect to  $\Sigma_k$  and setting to zero:

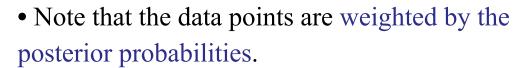
$$\Sigma_k = \frac{1}{N_k} \sum_{n=1}^N \gamma(z_{nk}) (\mathbf{x}_n - \boldsymbol{\mu}_k) (\mathbf{x}_n - \boldsymbol{\mu}_k)^T. \quad \boldsymbol{\pi} \leftarrow \sum_{k=1}^{\infty} \sum_{n=1}^{\infty} \gamma(z_{nk}) (\mathbf{x}_n - \boldsymbol{\mu}_k) (\mathbf{x}_n - \boldsymbol{\mu}_k)^T.$$

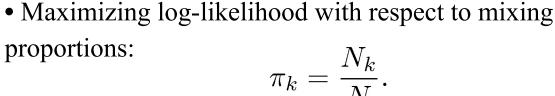
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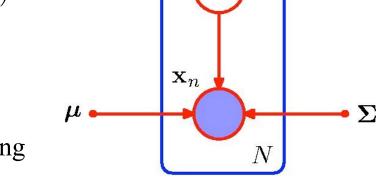
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• Differentiating with respect to  $\Sigma_k$  and setting to zero:

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







• Mixing proportion for the k<sup>th</sup> component is given by the average responsibility which that component takes for explaining the data.

• The log-likelihood takes form:

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

- 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:

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

 $\mathbf{z}_n$ 

• The log-likelihood:

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• Iterative Solution:

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

M-step: Update model parameters:  $\mu_k \sum_k \pi_k$ , for k = 1,...,K.

## EM algorithm

- ullet Initialize the means  $\mu_k$  , covariances  $\Sigma_k$  , and mixing proportions  $\pi_k$
- E-step: Evaluate responsibilities using current parameter values:

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

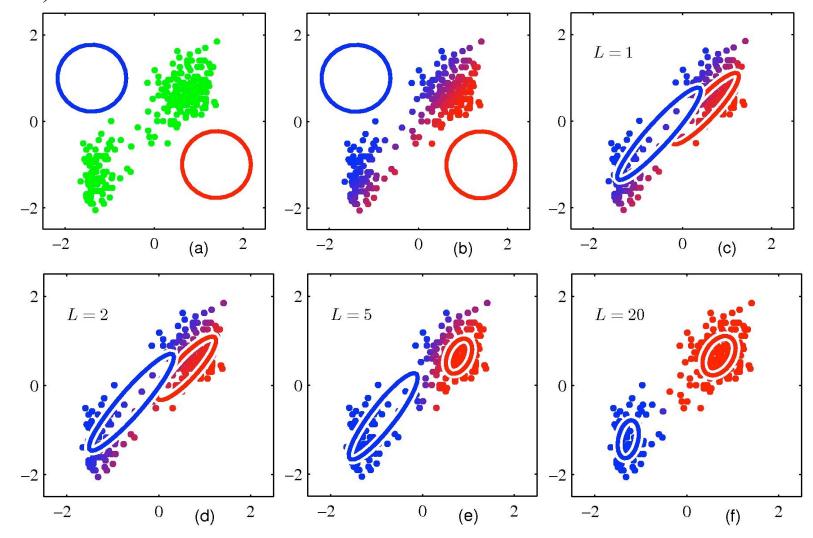
• M-step: Re-estimate model parameters using the current responsibilities:

$$oldsymbol{\mu}_k^{new} = rac{1}{N_k} \sum_n \gamma(z_{nk}) \mathbf{x}_n, \quad N_k = \sum_n \gamma(z_{nk}),$$
 $oldsymbol{\Sigma}_k^{new} = rac{1}{N_k} \sum_{n=1}^N \gamma(y_{nk}) (\mathbf{x}_n - oldsymbol{\mu}_k) (\mathbf{x}_n - oldsymbol{\mu}_k)^T,$ 
 $oldsymbol{\pi}_k^{new} = rac{N_k}{N}.$ 
 $oldsymbol{\mu}_k^{new}$ 

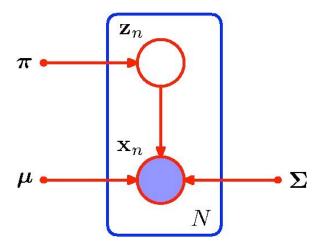
• Evaluate the log-likelihood and check for convergence.

### Mixture of Gaussians: Example

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

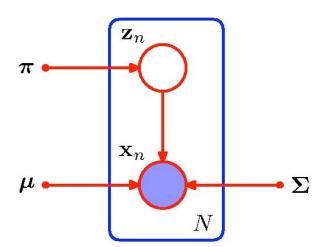


- The goal of EM is to find maximum likelihood solutions for models with latent variables.
- We represent the observed dataset as an N by D matrix X.
- Latent variables will be represented and an N by K matrix **Z**.
- The set of all model parameters is denoted by  $\theta$ .



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- We represent the observed dataset as an N by D matrix X.
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- The set of all model parameters is denoted by  $\theta$ .
- The log-likelihood takes form:

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



- We will call:
  - $\{X, Z\}$  as complete dataset.
    - $\{X\}$  as incomplete dataset.

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

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

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

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

## The General EM algorithm

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

where 
$$\begin{aligned} \theta^{new} &= \arg\max_{\theta} \mathcal{Q}(\theta, \theta^{old}). \\ \mathcal{Q}(\theta, \theta^{old}) &= \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \theta^{old}) \ln p(\mathbf{X}, \mathbf{Z}|\theta). \end{aligned}$$

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

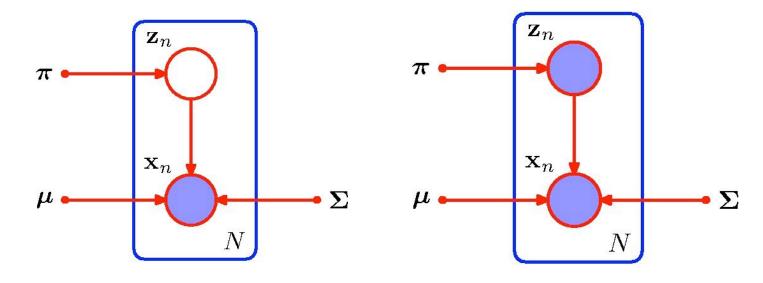
Otherwise:

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

#### Gaussian Mixtures Revisited

• We now consider the application of the latent variable view of EM to the case of Gaussian mixture model.

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



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

# Maximizing Complete Data

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

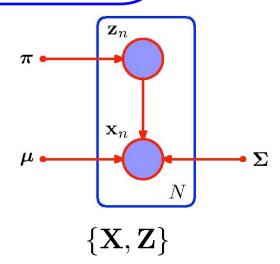
$$p(\mathbf{X}, \mathbf{Z} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \prod_{n=1}^{N} \prod_{k=1}^{K} \left[ \pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right]^{z_{nk}}.$$

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

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

• Maximizing with respect to mixing proportions yields:

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



-- complete dataset.

#### Posterior Over Latent Variables

• Remember:

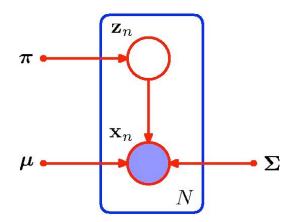
$$p(\mathbf{x}|\mathbf{z}) = \prod_{k=1}^K \mathcal{N}(\mathbf{x}|\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)^{z_k}, \quad p(\mathbf{z}) = \prod_{k=1}^K \pi_k^{z_k}.$$

• The posterior over latent variables takes form:

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

• Note that the posterior factorizes over n points, so that under the posterior distribution  $\{z_n\}$  are independent.

$$p(z_{nk} = 1 | \mathbf{X}, \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{\pi_k \mathcal{N}(x_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(x_n | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}$$



# Expected Complete Log-Likelihood

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

$$\mathbb{E}[z_{nk}] = \frac{\sum_{z_{nk}} z_{nk} [\pi_k \mathcal{N}(\mathbf{x}_n | \mu_k, \mathbf{\Sigma}_k)]^{z_{nk}}}{\sum_{z_{nj}} [\pi_j \mathcal{N}(\mathbf{x}_n | \mu_j, \mathbf{\Sigma}_j)]^{z_{nj}}}$$
Under posterior
$$= \frac{\pi_k N(\mathbf{x}_n | \boldsymbol{\mu}_k, \mathbf{\Sigma}_k)}{\sum_{j=1}^K \pi_j N(\mathbf{x}_n | \boldsymbol{\mu}_j, \mathbf{\Sigma}_j)} = \gamma(z_{nk}).$$

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- This represent the responsibility of component k for data point  $x_n$ .
- 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:

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

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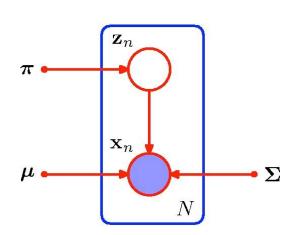
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• Maximizing with respect to model parameters we obtain:

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

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

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



## Relationship to K-Means

ullet Consider a Gaussian mixture model in which covariances are shared and are given by  $\epsilon I$ 

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

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

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

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

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

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

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

- Hence in the limit, maximizing the expected complete log-likelihood is equivalent to minimizing the distortion measure in the K-means algorithm.
- This is why assignment step in k-Means is called E-step, and finding the cluster centers is called the M-step.