# Mixture of Gaussians Expectation Maximization (EM)

# Part 2

Most of the slides are due to Christopher Bishop BCS Summer School, Exeter, 2003. The rest of the slides are based on lecture notes by A. Ng

## Limitations of K-means

- Hard assignments of data points to clusters small shift of a data point can flip it to a different cluster
- Not clear how to choose the value of K
- Solution: replace 'hard' clustering of K-means with 'soft' probabilistic assignments
- Represents the probability distribution of the data as a *Gaussian mixture model*

### **Gaussian Mixtures**

• Linear super-position of Gaussians

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

• Normalization and positivity require

$$\sum_{k=1}^{K} \pi_k = 1 \qquad 0 \leqslant \pi_k \leqslant 1$$

• Can interpret the mixing coefficients as prior probabilities

$$p(\mathbf{x}) = \sum_{k=1}^{K} p(k) p(\mathbf{x} \mid k)$$

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## Maximum Likelihood for the GMM

• The log likelihood function takes the form

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

- Note: sum over components appears inside the log
- There is no closed form solution for maximum likelihood
- How to maximize the log likelihood
  - solved by expectation-maximization (EM) algorithm

# EM Algorithm – Informal Derivation

- The solutions are not closed form since they are coupled
- Suggests an iterative scheme for solving them:
  - Make initial guesses for the parameters
  - Alternate between the following two stages:
    - 1. E-step: evaluate responsibilities
    - 2. M-step: update parameters using ML results

#### Jensen's inequality:

Let *f* be a convex function  $(f''(x) \ge 0 \text{ for all } x \in \Re)$  and *X* be a and random variable. Then  $E[f(X)] \ge f(E[X])$ .



f is convex X =a with probability 0.5 X =b with probability 0.5

E[X] is given by the midpoint between a and b. E[f(X)] is the midpoint between f(a) and f(b).  $E[f(X)] \ge f(EX)$ . Further, if f is a strictly convex function (f''(x) > 0), then E[f(X)] = f(E[X]) holds true if and only if E[X] = X with probability 1 (i.e., if X is a constant).

Jensen's inequality also holds for concave functions  $(f''(x) \le 0)$ , but with the direction of all the inequalities reversed ( $E[f(X)] \le f(E[X])$ , etc.).

# **Problem Definition**

- Suppose we have an estimation problem in which we have a training set {*x*<sub>1</sub>,.., *x<sub>m</sub>*} of iid samples.
- We wish to fit the parameters of a model  $p(x, z; \theta)$  to the data.
- We want to maximize the likelihood

$$l(\theta) = \sum_{i=1}^{m} \log p(x,\theta) = \sum_{i=1}^{m} \log \sum_{z} p(x,z;\theta)$$

- Doing it explicitly may be hard, since z's are the nonobserved.
- If *z*'s were observed, then (often) maximum likelihood estimation would be easy.

# EM at glance

- Our strategy will be to repeatedly
  - construct a lower-bound on  $l(\theta)$  (E-step),
  - optimize that lower-bound (M-step).



# EM algorithm derivation

#### EM algorithm derivation (cont.)

$$l(\theta) \ge \sum_{i=1}^{m} \sum_{z_i} Q_i(z_i) \log \frac{p(x_i, z_i; \theta)}{Q_i(z_i)}$$

We want a lower bound to be equal to *l* at the previous  $\theta$ 



## EM algorithm derivation (cont.)

To ensure that, we should choose  $Q_i(z)$  such that inequality in our derivation above would hold with equality. We require that:

$$\frac{p(x_i, z_i; \theta)}{Q_i(z_i)} = const \qquad \Longrightarrow \qquad Q_i(z_i) \propto p(x_i, z_i; \theta)$$

Since we know that  $\sum_{z} Q_i(z_i) = 1$ , then  $Q_i(z_i) = \frac{p(x_i, z_i; \theta)}{\sum_{z} p(x_i, z_i; \theta)} = \frac{p(x_i, z_i; \theta)}{p(x_i; \theta)} = p(z_i | x_i; \theta)$  Repeat until convergence {

- E-step: For each *i* set  $Q_i(z_i) \coloneqq p(z_i | x_i; \theta)$ 

- M-step:  

$$\theta \coloneqq \arg \max_{\theta} \sum_{i=1}^{m} \sum_{z_i} Q_i(z_i) \log \frac{p(x_i, z_i; \theta)}{Q_i(z_i)}$$
Lower bound on  $l(\theta)$ 

# EM for MoG revisited

- For  $1 \le i \le N$ ,  $1 \le j \le K$ , define hidden variables  $Z_{ij}$  $Z_{ij} = \begin{cases} 1 \text{ if sample i was generated by component } \mathbf{k} \\ 0 \text{ otherwise} \end{cases}$
- $z_{ij}$  are indicator random variables, they indicate which Gaussian component generated sample  $x_i$
- Let z<sub>i</sub> = {z<sub>i1</sub>,..., z<sub>iK</sub>} indicator r.v. correspond to sample x<sub>i</sub>.
   We say that z<sub>i</sub> = k, when its k'st coordinate is 1 and the rest are 0.
- Conditioned on  $Z_i$ , distribution of  $X_i$  is Gaussian

$$p(\mathbf{x}_i \mid z_i = k) \sim N(\mu_k, \Sigma_k)$$

E-step:

$$Q_i(z_i = k) = p(z_i = k | x_i; \mu, \Sigma, \pi)$$
$$= \frac{\pi_k N(x_i | \mu_k, \Sigma_k)}{\sum_j \pi_j N(x_i | \mu_j, \Sigma_j)} = \gamma_k(x_i)$$

#### EM for MoG revisited

$$\begin{array}{ll} \text{M-step:} & \max_{\mu,\Sigma,\pi} \boxed{\sum_{i=1}^{m} \sum_{z_i} Q_i(z_i) \log \frac{p(x_i, z_i; \theta)}{Q_i(z_i)}} \\ & = \sum_{i=1}^{N} \sum_{k=1}^{K} \gamma_k(x_i) \log \frac{\pi_k N(x_i \mid \mu_k, \Sigma_k)}{\gamma_k(x_i)} \\ & = \sum_{i=1}^{N} \sum_{k=1}^{K} \gamma_k(x_i) \log \frac{\pi_k N(x_i \mid \mu_k, \Sigma_k)}{\gamma_k(x_i)} \\ & \nabla_{\mu}(\ldots) \stackrel{\text{set}}{=} 0 \implies \mu_k = \frac{\sum_{i=1}^{N} \gamma_k(x_i) x_i}{\sum_{i=1}^{N} \gamma_k(x_i)} \\ & \text{Similarly,} \\ & \pi_k = \frac{1}{N} \sum_{i=1}^{N} \gamma_k(x_i), \qquad \Sigma_k = \frac{\sum_{i=1}^{N} \gamma_k(x_i) (x_i - \mu_k) (x_i - \mu_k)^T}{\sum_{i=1}^{N} \gamma_k(x_i)} \end{array}$$

# K-means Algorithm

- Goal: represent a data set in terms of K clusters each of which is summarized by a prototype  $\mu_k$
- Initialize prototypes, then iterate between two phases:
  - E-step: assign each data point to nearest prototype
  - M-step: update prototypes to be the cluster means

#### Responsibilities

• *Responsibilities* assign data points to clusters

$$r_{nk} \in \{\mathsf{0},\mathsf{1}\}$$

such that

$$\sum_{k} r_{nk} = 1$$

• Example: 5 data points and 3 clusters

$$(r_{nk}) = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 0 & 1 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \\ 1 & 0 & 0 \end{pmatrix}$$

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# Minimizing the Cost Function

- E-step: minimize J w.r.t.  $r_{nk}$ 
  - assigns each data point to nearest prototype
- M-step: minimize J w.r.t  $\mu_k$

- gives

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

- each prototype set to the mean of points in that cluster

• Convergence guaranteed since there is a finite number of possible settings for the responsibilities

- Example from R. Gutierrez-Osuna
- Training set of 900 examples forming an annulus
- Mixture model with m = 30 Gaussian components of unknown mean and variance is used
- Training:
  - Initialization:
    - means to 30 random examples
    - covaraince matrices initialized to be diagonal, with large variances on the diagonal (compared to the training data variance)
  - During EM training, components with small mixing coefficients were trimmed
    - This is a trick to get in a more compact model, with fewer than 30 Gaussian components

# **EM Example**



from R. Gutierrez-Osuna

# **EM Motion Segmentation Example**

Three frames from the MPEG "flower garden" sequence



Figure from "Representing Images with layers,", by J. Wang and E.H. Adelson, IEEE Transactions on Image Processing, 1994, c 1994, IEEE