

- Basic review of Gaussians
- Mixture models
- EM for mixture models

### **Recall from last time: Expectation Maximization (EM)**

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- A general purpose method for learning from incomplete data
- Main idea:
  - If we had complete data (sufficient statistics) we could easily maximize the likelihood
  - So in the case of missing values, we will "fantasize" what they should be, based on the current parameter setting
  - This means we compute expected sufficient statistics
  - Then we improve the parameter setting, based on these statistics



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#### **Quick refresher: Gaussian distributions**

• The univariate Gaussian (Normal distribution) with mean  $\mu$  and variance  $\sigma$ :

$$p(x;\mu,\Sigma) = \frac{1}{\sqrt{(2\pi\sigma^2)}} \exp\left(-\frac{(x-\mu)^2}{2\sigma^2}\right)$$

• Suppose we have data with n attributes (so now a data instance  $\vec{x}$  is an n-dimensional vector)

The <u>multivariate</u> Gaussian distribution with **mean vector**  $\vec{\mu}$  (*n*-dimensional vector) and **covariance matrix**  $\Sigma$  ( $n \times n$  matrix, symmetric, positive semi-definite) is:

$$p(\vec{x}) = \frac{1}{(2\pi)^{n/2} |\Sigma|^{1/2}} \exp\left(\frac{1}{2} (\vec{x} - \vec{\mu})^T \Sigma^{-1} (\vec{x} - \vec{\mu})\right)$$

where  $|\Sigma|$  denotes the determinant of  $\Sigma$ 

### **Mixture of Gaussians**

- Suppose we have a set of data points,  $\langle ec{x_1}, \ldots ec{x_m} 
  angle$
- Gaussian assumption: Suppose there are k Gaussian distributions  $\mathcal{N}(\vec{\mu_j}, \Sigma_j), j = 1 \dots k$  which generate data.
- Every data point is generated by first selecting one of these distributions, with probability p<sub>j</sub>, j = 1, ... k, and then generating a point from the distribution.
- Can we estimate  $p_j, \vec{\mu_j}, \Sigma_j, j = 1 \dots k$  from the data?
- We'll do the 1-dimensional case first...

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### Maximum likelihood to the rescue!

- Write down the log likelihood of the data given a set of parameters p<sub>i</sub>, μ<sub>i</sub>, σ<sub>i</sub>, i = 1,...k:
  - $\log L = \log p(\langle x_1, \dots, x_n \rangle | p_1, \dots, p_k, \mu_1, \dots, \mu_k, \sigma_1 \dots, \sigma_k)$   $= \log \prod_{i=1}^m p(x_i | p_1, \dots, p_k, \mu_1, \dots, \mu_k, \sigma_1 \dots, \sigma_k) \text{ (assuming data is iid)}$   $\sum_{i=1}^m \left(\sum_{i=1}^k p(x_i | p_1, \dots, p_k, \mu_1, \dots, \mu_k, \sigma_1 \dots, \sigma_k) \right)$

$$=\sum_{i=1}^m \log\left(\sum_{j=1}^\kappa p(x_i|\mu_j,\sigma_j)p_j
ight)$$
 (given how the data is drawn)

• Now compute derivative wrt  $p_j$ ,  $\mu_j$ ,  $\sigma_j$ , set to 0 and solve...

#### This cannot be solved analytically!



# A simple problem!

- Suppose we knew the latent variables (i.e., we knew which Gaussian generated which point)
- Let  $k_i$  be the Gaussian that generated point *i*. Then we have:

$$\log L = \sum_{i=1}^{m} \log \left( p(x_i | \mu_{k_i}, \sigma_{k_i}, p_{k_i}) \right)$$
$$= \sum_{i=1}^{m} \left( \log p(\mathbf{x}_i | \mu_{k_i}, \sigma_{k_i}) + \log p_{k_i} \right)$$
$$= \sum_{i=1}^{m} \left( -\frac{(x_i - \mu_{k_i})^2}{2\sigma_{k_i}^2} - \frac{1}{2} \log \left( 2\pi \sigma_{k_i}^2 \right) + \log p_{k_i} \right)$$

• Now it is easy to maximize the likelihood!

# Maximum likelihood estimation: Univariate Gaussian

- Working with  $k_i$  (the component which generated instance *i*) is awkward.
- To make things easy, we'll define an indicator variable,  $\delta_{ij}$ , which is equal to 1 if and only if  $k_i = j$ , 0 otherwise.
- With this notation, the likelihood becomes:

$$\log L = \sum_{i=1}^{m} \sum_{j=1}^{k} \delta_{ij} \left( -\frac{(x_i - \mu_j)^2}{2\sigma_j^2} - \frac{1}{2} \log (2\pi\sigma_j^2) + \log p_j \right)$$

Note that we can reverse the sums at will

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# Maximum likelihood estimation: Univariate Gaussian (2)

• To compute the means  $\mu_j$ , we have:

$$\frac{\partial \log L}{\partial \mu_j} = \sum_{i=1}^m \delta_{ij} \left( -\frac{2(x_i - \mu_j)(-1)}{2\sigma_j^2} \right) = 0$$

Solving this, we get:

$$\mu_j = \frac{\sum_{i=1}^m \delta_{ij} x_i}{\sum_{i=1}^m \delta_{ij}}$$

In other words, to estimate the mean of the jth component, we compute the sample mean of the instances generated by it

• Similarly, we get for the standard deviation:

$$\sigma_j^2 = \frac{\sum_{i=1}^m \delta_{ij} (x_i - \mu_j)^2}{\sum_{i=1}^m \delta_{ij}}$$

# Maximum likelihood estimation: Univariate Gaussian (3)

- To find  $p_j$ , we also have to take into account that  $\sum_j p_j = 1$ .
- Assume without loss of generality (wlog) that we replace  $p_k = 1 \sum_{j=1}^{k-1} p_j$ :

$$\frac{\partial \log L}{\partial p_j} = \sum_{i=1}^m \left( \delta_{ij} \frac{1}{p_j} + \delta_{ik} \frac{1}{1 - \sum_{j=1}^{k-1} p_j} (-1) \right) = 0, \forall j = 1, \dots k - 1$$

• By manipulating this equation, we get:

$$\frac{\sum_{i=1}^{m} \delta_{ij}}{\sum_{i=1}^{m} \delta_{ik}} = \frac{p_j}{p_k}, \forall j = 1, \dots k-1$$

• Now we take sums on both sides:

$$\sum_{j=1}^{k-1} \frac{\sum_{i=1}^{m} \delta_{ij}}{\sum_{i=1}^{m} \delta_{ik}} = \sum_{j=1}^{k-1} \frac{p_j}{p_k}$$

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### Maximum likelihood estimation: Univariate Gaussian (4)

- We know that  $\sum_{i=1}^{m} \sum_{j=1}^{k} \delta_{ij} = m$  (because each instance is generated by exactly one Gaussian component)
- We also know that  $\sum_{j=1}^{k-1} p_j = 1 p_k$
- Plugging everything back in the equation, we get:

$$\frac{m - \sum_{i=1}^{m} \delta_{ik}}{\sum_{i=1}^{m} \delta_{ik}} = \frac{1 - p_k}{p_k}$$

• Solving, we get that the probability  $p_k$  is given by the empirical fraction of the points coming from the *k*th distribution:

$$p_k = \frac{\sum_{i=1}^m \delta_{ik}}{m}$$

And because we did this wlog, it holds for any  $p_j$ .

# Maximum likelihood estimation: Multivariate Gaussian

- It is just a straightforward generalization of the univariate case
- The probability  $p_j$  is just the number of points coming from the jth distribution

$$p_j = \frac{\sum_{i=1}^m \delta_{ij}}{m}$$

• The mean of the distribution is given by the empirical mean of the points coming from it:

$$\vec{\mu_j} = \frac{\sum_{i=1}^m \delta_{ij} \vec{x_i}}{\sum_{i=1}^m \delta_{ij}}$$

• The covariance matrix is the covariance of the points from the distribution

$$\Sigma_{j} = \frac{\sum_{i=1}^{m} \delta_{ij} \left( \vec{x_{i}} - \vec{\mu_{j}} \right) \left( \vec{x_{i}} - \vec{\mu_{j}} \right)^{T}}{\sum_{i=1}^{m} \delta_{ij}}$$

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### **EM for Mixture of Gaussians**

- We start with an initial guess for the parameters  $p_j$ ,  $\vec{\mu_j}$ ,  $\Sigma_j$
- We will iterate an e-step, in which we "complete" the data, with an M-step, in which we re-compute the parameters
- In the "hard EM" version, completing the data means that each data point is assumed to be generated by *exactly one Gaussian*
- This is very related to *k-means clustering* (which you may know)
- In the "soft EM" version (also usually known as EM), we assume that each data point could have been generated from *any component*
- In this case, each point will contribute to the mean and variance estimate of each component.

### Hard EM for Mixture of Gaussians

- 1. Guess an initial parameter setting  $p_j, \vec{\mu_j}, \Sigma_j, j = 1 \dots k$
- 2. Repeat until convergence:
  - (a) *E-step:* For each i = 1, ..., m and each j = 1, ..., k:

 $k_i = rg\max_j p(ec{x_i}$  drawn from distribution  $j|p_j, ec{\mu_j}, \Sigma_j)$ 

where  $p(ec{x_i}$  drawn from distribution  $j|p_j, ec{\mu_j}, \Sigma_j) \propto p_j p(ec{x_i}|ec{\mu_j}, \Sigma_j)$ 

(b) <u>*M-step:*</u> Update the parameters of the model to maximize the likelihood of the data

$$p_{j} = \frac{1}{m} \sum_{i=1}^{m} \delta_{ij} \qquad \vec{\mu_{j}} = \frac{\sum_{i=1}^{m} \delta_{ij} \vec{x_{i}}}{\sum_{i=1}^{m} \delta_{ij}}$$
$$\Sigma_{j} = \frac{\sum_{i=1}^{m} \delta_{ij} (\vec{x_{i}} - \vec{\mu_{j}}) (\vec{x_{i}} - \vec{\mu_{j}})^{T}}{\sum_{i=1}^{m} \delta_{ij}}$$

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#### Soft EM for Mixture of Gaussians

- 1. Guess an initial parameter setting  $p_j, \vec{\mu_j}, \Sigma_j, j = 1, \dots k$
- 2. Repeat until convergence:
  - (a) *E-step:* For each i = 1, ..., m and each j = 1, ..., k:

 $w_j(i) = p(\vec{x_i} ext{ drawn from distribution } j | p_j, \vec{\mu_j}, \Sigma_j)$ 

where  $p(ec{x_i}$  drawn from distribution  $j|p_j, ec{\mu_j}, \Sigma_j) \propto p_j p(ec{x_i}|ec{\mu_j}, \Sigma_j)$ 

(b) <u>*M-step:*</u> Update the parameters of the model to maximize the likelihood of the data

$$p_{j} = \frac{1}{m} \sum_{i=1}^{m} w_{j}(i) \qquad \vec{\mu_{j}} = \frac{\sum_{i=1}^{m} w_{j}(i) \vec{x_{i}}}{\sum_{i=1}^{m} w_{j}(i)}$$
$$\Sigma_{j} = \frac{\sum_{i=1}^{m} w_{j}(i) (\vec{x_{i}} - \vec{\mu_{j}}) (\vec{x_{i}} - \vec{\mu_{j}})^{T}}{\sum_{i=1}^{m} w_{j}(i)}$$

# EM in general

- Whenever we are trying to model data drawn probabilistically, and we have missing values in the data, EM is an option
- We need some structured or parametric form of the distribution (we saw Bayes nets and mixtures of Gaussians as examples)
- We starts with a guess for the parameters of the distribution
- You can think of the E-step as trying to "complete" the data, by filling in the missing values
- The M-step will compute new parameters, given the completed data

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# Theoretical properties of EM

• Each iteration improves the likelihood:

$$L(\theta_{i+1}|D) \ge L(\theta_i|D)$$

 If the parameters do not change in one iteration, θ<sub>i+1</sub> = θ<sub>i</sub>, then the gradient of the log-likelihood function is 0 at θ<sub>i</sub>:

$$\frac{\partial L(\theta|D)}{\partial \theta}(\theta_i) = 0$$

This means that  $\theta_i$  is a min, max or saddle point

- In practice, convergence only occurs at local maxima
- See textbook for a detailed description of general EM and its properties

# EM pros and cons

- Much easier to implement than gradient descent; no parameter tuning is necessary, and no projection of the parameters (we compute them directly normalized)
- Converges much faster than vanilla gradient descent
- Not very sensitive to the starting point
- Speed comparison with fancy gradient descent is unclear

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# Summary

- Difficulty of inference with missing data is that it generates a complex likelihood function
- We have a non-linear optimization problem, and we can use two solutions:
  - Gradient descent: always works for non-linear optimization
  - EM: targeted towards optimizing likelihood
- Both are only guaranteed to converge to local maxima, so we need restarts
- Both use inference to compute expected sufficient statistics of the data
- Hence, the inference step is a bottleneck for both