Expectation maximization, FA/PCA continued

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Figure 1 is the graphical model that motivates the following discussion of factor analysis (FA).

Figure 1:



In FA, the basic idea is to choose z from some distribution in q dimensions and project it onto a p-dimensional space and then choose x given the projection. To begin with, we define the variable distributions.

$$\langle z, x \rangle$$
 is a joint Gaussian
 $x \sim \mathcal{N}(0, \Lambda \Lambda^T + \Psi)$
 $z | x \sim \mathcal{N}(\Lambda^T (\Lambda \Lambda^T + \Psi)^{-1} X, (I + \Lambda^T \Psi \Lambda^T)^{-1})$

We want to get the MLE of Λ and Ψ , given that we have data, $\mathcal{D} = \{x_n\}_{n=1}^N$.

Notice that $x = \Lambda z + \epsilon$. If we know z, then this is a linear regression. But z is a hidden variable. So we are going to use the expectation maximization (EM) algorithm. Generally speaking, this is a way of solving maximization problems in the face of hidden variables. The EM algorithm for factor analysis is an iterative algorithm with 2 steps:

- 1. the E-step:
 - compute $p(z_n|x_n, \Lambda^{(t)}, \Psi^{(t)})$
 - the posterior p(z|x) is defined above
- 2. The M-step:
 - $\Lambda^{(t=1)} = (\sum_{n} \mathbb{E}[z_n z_n^T | x_n, \Lambda^{(t)}, \Psi^{(t)}])^{-1} (\sum_{n} \mathbb{E}[z_n | x_n, \Lambda^{(t)}, \Psi^{(t)}]^T x_n)$
 - $\Psi^{(t+1)} = \text{See book}$

EM is a way of finding approximate MLE's in latent variable models. We will be thinking about these in the rest of the course. Latent variable models posit hidden structure in observed data: clustering, subspace, trees, sequences etc.

One way to think about EM: in the E-step, we will fill in the hidden variables. In the M-step, we fit parameters to match the filled in variables (akin to taking the MLE estimate in a fully observed model). So, fill in z and then estimate the parameters. This gets us around having to integrate out the latent variables.

EM general setting

- $x_n = \{1....N\}$ observed data
- z_n hidden structure
- θ are the parameters we are interested in fitting.
- There is no particular graphical model.

What if z were observed? We could find the parameters by taking the max of the loglikelihood.

$$\hat{\theta} = \arg \max_{\theta} \log p(x, z | \theta)$$
$$= \arg \max_{\theta} \log p(x | z, \theta_x) + \log p(z | \theta_z)$$

This function is called the complete log-likelihood. But z is hidden, so we are really after

$$\hat{\theta} = \arg \max_{\theta} \log p(x|\theta)$$
$$= \arg \max_{\theta} \log \sum_{z} p(x, z|\theta)$$

Where the hidden variable has been factored out.

Note Jenson's inequality:

We will have a lower bound $\log p(x)$ on Jenson's inequality. If $\lambda \in (0,1)$ and φ is convex then:

$$\lambda \varphi(x) + (1 - \lambda)\varphi(y) \ge \varphi(\lambda(x) + (1 - \lambda)(y))$$

Which generalizes to expectations –

$$\mathbf{E}[\varphi(x)] \ge \varphi(\mathbf{E}[x])$$

And if φ is concave –

 $\mathbf{E}[\varphi(x)] \le \varphi(\mathbf{E}[x])$

Now back to EM:

$$\log p(x|\theta) = \log \sum_{z} p(x, z|\theta)$$
$$= \log \sum_{z} p(x, z|\theta) \frac{q(z)}{q(z)}$$
$$= \log E_q \left[\frac{p(x, z|\theta)}{q(z)} \right]$$
$$\geq E_q \left[\log \frac{p(x, z|\theta)}{q(z)} \right]$$
$$= E_q \log p(x, z|\theta) - E_q \log q(z)$$
$$\equiv \mathcal{Q}(\theta; q)$$

This is the EM objective function. The EM algorithm will optimize the objective function. The EM is a coordinate ascent on Q:

$$E: q^{(t+1)} = \arg \max_{q} \mathcal{Q}(\theta^{(t)}, q)$$
$$M: \theta^{(t+1)} = \arg \max_{\theta} \mathcal{Q}(\theta, q^{(t+1)})$$

Holding θ fixed, the optimal q(z) is $p(z|x, \theta^{(t)})$.

$$\begin{split} &= \sum_{z} p(z|x) \log p(x,z) - \sum_{z} p(z|x) \log p(z|x) \\ &= \sum_{z} p(z|x) \log p(z|x) + \sum_{z} p(z|x) \log p(z|x) \log p(x) - \sum_{z} p(z|x) \log p(z|x) \\ &= \sum_{z} p(z|x) \log p(x) \\ &= \log p(x) \end{split}$$

M-step:

$$\theta^{(t+1)} = \arg \max_{\theta} \mathbf{E}_q \log p(x, z|\theta)$$
$$= \arg \max_{\theta} \mathbf{E}_q \log p(z|\theta) + \mathbf{E}_q \log p(x|z, \theta)$$

Which is the expected complete log-likelihood.

Mixture modeling

- E-step: estimate p(cluster|datapoint)
- $\bullet\,$ M-step: reweight the data by p($|\mathbf{x})$ and do MLE.