Lecture 11: Iterative Proportional Fitting

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Undirected Models

- In directed models, each node plus its parents form a clique, and the clique potential are $p(x_i|x_{\pi_i})$.
- In undirected models, we can have arbitrary cliques $C$, with arbitrary postive potentials $\psi_C$ on each one.
- But this flexibility comes at a cost: learning in fully observed directed models is trivial, but learning in fully observed undirected models is not.
- Reason? The normalization factor (partition function) $Z$:

$$P(X) = \frac{1}{Z} \prod_{\text{cliques } c} \psi_c(x_c) \quad Z = \sum_{X} \prod_{\text{cliques } c} \psi_c(x_c)$$

Discrete Nodes

- Consider undirected models with all discrete nodes.
- Now there are just a finite number of possible joint settings of $X$, and our model is defined as:

$$P(x|\theta) = \frac{1}{Z(\theta)} \prod_c \psi_c(x_c|\theta_c) \quad Z(\theta) = \sum_x \prod_c \psi_c(x_c|\theta_c)$$

where $\theta = \{\theta_c\}$.
- For a particular dataset $D = \{x^1, x^2, \ldots, x^N\}$, we can count the number of times any joint configuration $x$ has been observed:

$$n(x) = \sum_n \delta(x, x^n)$$
- We can also count the number of times a clique configuration appears (using $x_c$ as the subvector over the variables in clique $c$):

$$n(x_c) = \sum_n \delta(x_c, x_c^n) = \sum_{x_c} n([x_c, x_c])$$

Which cliques?

- In general, an undirected model can place potentials on any subset of the cliques of the graph.
- Not all cliques need to be have potentials, and not all potentials need to be over maximal cliques.
- But in general, placing one potential on each maximal clique of the graph can represent any set of potentials on subsets of cliques.
Why? We can always set the potential on a maximal clique to the product of all the potentials on cliques which were contained in that maximal clique. We can always set the potential on cliques we wanted to leave out to unity.
In terms of the counts, the log likelihood is given by:

\[ p(D|\theta) = \prod_n \prod_x p(x|\theta)^{\delta(x, x^n)} \]

\[ \log p(D|\theta) = \sum_n \sum_x \delta(x, x^n) \log p(x|\theta) \]

\[ \ell = \sum_n n(x) \log \left( \frac{1}{Z} \prod_c \psi_c(x_c) \right) \]

\[ = \sum_c \sum_n n(x_c) \log \psi_c(x_c) - N \log Z \]

So the clique counts \( n(x_c) \) are the sufficient statistics for our undirected model.

But now there is a nasty \( \log Z \) in the likelihood.

A Game

There is a table (with all positive numbers that sum to one). I show you the row sums and the column sums.

Can you find the table? (Easy).
Can you find the table if I force some spots to be zero? (Harder).
Can you find the table if instead of row sums and column sums I tell you the sums of arbitrary subsets of the elements?

Iterative Proportional Fitting (IPF)

Let’s go back to the derivative of the likelihood:

\[ \frac{\partial \ell}{\partial \psi_c(x_c)} = \frac{n(x_c)}{\psi_c(x_c)} - N \frac{p(x_c)}{\psi_c(x_c)} \]

From this we can derive another relationship:

\[ \frac{n(x_c)}{N} \frac{q(x_c)}{\psi_c(x_c)} = \frac{p(x_c|\theta)}{\psi_c(x_c)} \]

in which \( \psi_c \) appears implicitly in the model marginal \( p_c(x_c|\theta) \).

To solve for \( \psi_c \) is hard, because it appears on both sides of this implicit nonlinear equation.

The idea of IPF is to hold \( \psi_c \) fixed on the right hand side (both in the numerator and denominator) and solve for it on the left hand side. We cycle through all cliques, then iterate:

\[ \psi_c^{(t+1)}(x_c) = \psi_c^{(t)}(x_c) \frac{q(x_c)}{p^{(t)}(x_c)} \]
Properties of IPF Updates

- The IPF updates have two amazing properties:
  1. At each iteration the model marginal $p(t+1)(x_c)$ is equal to the observed marginal $q(x_c)$.
  2. The partition function $Z$ remains constant across all IPF updates.

- To show this, calculate the marginal:
  
  $$p(t+1)(x_c) = \sum_{x_c} p(t+1)(x) = \frac{Z(t)}{Z(t+1)} q(x_c)$$

  Now sum both sides over $x_c$, yielding:
  
  $$1 \cdot Z(t+1) = Z(t) \cdot 1$$

  and thus
  
  $$p(t+1)(x_c) = q(x_c)$$

KL Divergence View

- IPF can also be seen to be coordinate ascent in the likelihood using the way of expressing likelihoods using KL divergences.
  
- First, we observe that maximizing the log likelihood is equivalent to minimizing the KL divergence (cross entropy) from the observed distribution to the model distribution:
  
  $$\max \ell \Leftrightarrow \min KL[q(x) \| p(x|\theta)] = \sum_x q(x) \log \frac{q(x)}{p(x|\theta)}$$

- Next, we use a property of KL divergence based on the conditional chain rule: $p(x) = p(x_a)p(x_b|x_a)$:
  
  $$KL[q(x_a, x_b) \| p(x_a, x_b)] = KL[q(x_a) \| p(x_a)] + \sum_{x_a} q(x_a) KL[q(x_b \| x_a) \| p(x_b \| x_a)]$$

- Putting these two together, we see that:
  
  $$KL[q(x) \| p(x|\theta)] = KL[q(x_c) \| p(x_c|\theta)] + \sum_{x_c} q(x_c) KL[q(x_c) \| p(x_c|\theta)]$$

  But changing the clique potential has no effect on the conditional distribution, so the second term in unaffected. To minimize the first term, we set the marginal to the observed marginal, just as in IPF.

- In fact, we can interpret IPF updates as retaining the “old” conditional probabilities $p(t)(x_c|x_c)$ while replacing the “old” marginal probability $p(t)(x_c)$ with the observed marginal $q(x_c)$. 

IPF minimizes KL divergence

- We saw that the IPF iterations always achieve the conditions we require at the ML parameter estimates.

- But how do we know if they will converge, and if so, if they always increase the log likelihood?

- We can show that IPF is actually coordinate ascent in the log likelihood, just like EM was.

- At any iteration, the derivative of the likelihood can be written as:
  
  $$\frac{\partial \ell}{\partial \psi_c(x_c)} = \frac{n(x_c)}{\psi_c(x_c)} - N \frac{p(t)(x_c)}{\psi_c(t)(x_c)}$$

  and if we evaluate this at $\psi_c(t+1)(x_c)$ we find that the gradient is in fact equal to zero.

IPF as Coordinate Ascent

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