The EM algorithm for Bayesian networks

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Entropy

The *entropy* of a discrete probability distribution P is

$$\operatorname{Ent}(P) = -\sum_{x} p(x) \log p(x).$$

Entropy is a measure of *spread* of the distribution and it is always positive.

The entropy is never larger than the entropy of the uniform distribution:

Let $P_u(x) = 1/|\mathcal{X}|$, then it holds that

 $0 \leq \operatorname{Ent}(P) \leq \operatorname{Ent}(P_u) = \log |\mathcal{X}|.$

Proof on next overhead.

Uniform distribution has maximal entropy

The information inequality

$$\sum_{x} p(x) \log p(x) \ge \sum_{x} p(x) \log q(x)$$

yields

$$\operatorname{Ent}(P) = -\sum_{x} p(x) \log p(x)$$
$$\leq -\sum_{x} p(x) \log \frac{1}{|\mathcal{X}|}$$
$$= -\sum_{x} \frac{1}{|\mathcal{X}|} \log \frac{1}{|\mathcal{X}|} = \operatorname{Ent}(P_u).$$

Kullback-Leibler divergence

The *KL divergence* between P and Q is

$$KL(P:Q) = \sum_{x} p(x) \log \frac{p(x)}{q(x)}.$$

Also known as *relative entropy* of Q with respect to P. Information inequality says that $KL(P:Q) \ge 0$ and KL(P:Q) = 0 if and only if P = Q, so KL divergence defines an (asymmetric) distance measure between probability distributions.

Incomplete observations

Bayesian network with conditional probability distributions only partially known:

$$p(x) = \prod_{v \in V} p(x_v \,|\, x_{\operatorname{pa}(v)}, \theta)$$

where $\theta \in \Theta \subseteq \mathcal{R}^k$ is unknown parameter.

Instead of *complete data* (x^1, \ldots, x^n) , only *incomplete data* $(x_{A_1}^1, \ldots, x_{A_n}^n)$ available, where $A_i \subseteq V$.

Example: paternity cases. Unknown parameters: gene frequencies, probability of paternity, possibly mutation rates.

EM algorithm

Complete data x, incomplete data (observed) y = g(x). Complete data log-likelihood:

 $l(\theta) = \log L(x \,|\, \theta) = \log p(x \,|\, \theta).$

The marginal log-likelihood is

$$l_y(\theta) = \log L(\theta \mid y) = \log p(y \mid \theta).$$

Wish to maximize l_y in θ but l_y is unpleasant:

$$l_y(\theta) = \log \sum_{x:g(x)=y} p(x \mid \theta).$$

However, we assume that we know how to maximize l. How can this be exploited?

E-step and M-step

We let θ^* be arbitrary but fixed.

The **E-step** calculates *expected* complete data log-likelihood $q(\theta | \theta^*)$:

$$q(\theta \mid \theta^*) = \mathbf{E}_{\theta^*} \{ l(\theta) \mid y \} = \sum_{x:g(x)=y} p(x \mid y, \theta^*) \log p(x \mid \theta).$$

The **M-step** maximizes $q(\cdot | \theta^*)$ for fixed θ^* : The algorithm alternates between an E-step and an M-step. *After an E-step and subsequent M-step, the likelihood*

function has never decreased, as we shall now show.

EM algorithm

Since $p(x \mid y, \theta) = \chi_{q(x)}(y)p(x \mid \theta)/p(y \mid \theta)$ we have $q(\theta \,|\, \theta^*) \quad = \quad \sum p(x \,|\, y, \theta^*) \log\{p(y \,|\, \theta) p(x \,|\, y, \theta)\}$ $= \log p(y \mid \theta) + \sum p(x \mid y, \theta^*) \log p(x \mid y, \theta)$ $= l_y(\theta) - \sum p(x \mid y, \theta^*) \log p(x \mid y, \theta^*)$ $-\sum p(x \mid y, \theta^*) \log \frac{p(x \mid y, \theta^*)}{p(x \mid y, \theta)}$ $= \overline{l_u(\theta)} - \operatorname{Ent} P^y_{\theta^*} - \overline{KL}(P^y_{\theta^*} : \overline{P^y_{\theta}}).$

Expected and complete data likelihood



$$l_{y}(\theta) - \operatorname{Ent}(P_{\theta^{*}}^{g}) = q(\theta \mid \theta^{*}) + KL(P_{\theta^{*}}^{g} : P_{\theta}^{g})$$
$$\nabla l_{y}(\theta^{*}) = \nabla q(\theta^{*} \mid \theta^{*})$$

Likelihood monotonicity of EM algorithm

Let $\theta_0 = \theta^*$ and $\theta_{n+1} = \arg \max_{\theta} q(\theta \,|\, \theta_n).$ Then

$$l_{y}(\theta_{n+1}) = q(\theta_{n+1} | \theta_{n}) + \operatorname{Ent}(P_{\theta_{n}}^{y}) + KL(P_{\theta_{n+1}}^{y} : P_{\theta_{n}}^{y})$$

$$\geq q(\theta_{n} | \theta_{n}) + \operatorname{Ent}(P_{\theta_{n}}^{y}) = l_{y}(\theta_{n}).$$

So likelihood never decreases. Note, this also holds if just $q(\theta_{n+1} | \theta_n) \ge q(\theta_n | \theta_n)$.

E-step for Bayesian networks

The complete data likelihood is

$$\log p(x \mid \theta) = \sum_{i=1}^{n} \log p(x^{i} \mid \theta) = \sum_{x} n(x) \log p(x \mid \theta).$$

where $n(x) = \#\{i : x^{i} = x\}$. Using factorization we get
 $\log p(x \mid \theta) = \sum_{x} \sum_{v} n(x) \log p(x_{v} \mid x_{\operatorname{pa}(v)}, \theta)$
 $= \sum_{v} \sum_{x_{v \cup \operatorname{pa}(v)}} n(x_{v \cup \operatorname{pa}(v)}) \log p(x_{v} \mid x_{\operatorname{pa}(v)}, \theta),$

with $n(x_{v \cup pa(v)}) = #\{i : x_{v \cup pa(v)}^i = x_{v \cup pa(v)}\}$. So E-step equivalent to computing

 $n^*(x_{v \cup \mathrm{pa}(v)}) = \mathbf{E}\{N(x_{v \cup \mathrm{pa}(v)}) \,|\, y, \theta^*\}.$

Computing expected counts

We now get

$$n^{*}(x_{v \cup \mathrm{pa}(v)}) = \mathbf{E}\{N(x_{v \cup \mathrm{pa}(v)}) \mid y, \theta^{*}\}$$
$$= \sum_{i} \mathbf{E}\{\chi_{x_{v \cup \mathrm{pa}(v)}}(x_{v \cup \mathrm{pa}(v)}^{i}) \mid y, \theta^{*}\}$$
$$= \sum_{i} \mathbf{E}\{\chi_{x_{v \cup \mathrm{pa}(v)}}(x_{v \cup \mathrm{pa}(v)}^{i}) \mid x_{A_{i}}^{i}, \theta^{*}\}$$
$$= \sum_{i} p(x_{v \cup \mathrm{pa}(v)} \mid x_{A_{i}}^{i}, \theta^{*}).$$

Each of the latter terms can be calculated by probability propagation as can the marginal likelihood function:

$$\log p(y \mid \theta) = \sum_{i} \log p(x_{A_i}^i \mid \theta).$$

M-step for Bayesian networks

Note the similarity between the complete data likelihood and q:

$$\log p(x \mid \theta) = \sum_{v} \sum_{x_{v \cup \operatorname{pa}(v)}} n(x_{v \cup \operatorname{pa}(v)}) \log p(x_v \mid x_{\operatorname{pa}(v)}, \theta)$$

whereasx

$$q(\theta \mid \theta^*) = \sum_{v} \sum_{x_{v \cup \mathrm{pa}(v)}} n^*(x_{v \cup \mathrm{pa}(v)}) \log p(x_v \mid x_{\mathrm{pa}(v)}, \theta).$$

So any algorithm which maximizes the complete data likelihood can be used to maximize q in the M-step.