PROBABILISTIC GRAPHICAL MODELS CPSC 532C (TOPICS IN AI) STAT 521A (TOPICS IN MULTIVARIATE ANALYSIS)

Lecture 2

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REVIEW: PROBABILISTIC INFERENCE (STATE ESTIMATION)

• Inference is about estimating hidden (query) variables *H* from observed (visible) measurements *v*, which we can do as follows:

$$P(h|v) = \frac{P(v,h)}{\sum_{h'} P(v,h')}$$

- Examples:
 - Medical diagnosis: H diseases, v = findings/ symptoms,
 - -Speech recognition: H = spoken words, v = acoustic waveform
 - Genetic pedigree analysis: H = genotype, v = phenotype

- Class web page http://www.cs.ubc.ca/~murphyk /Teaching/CS532c_Fall04/index.html
- Send email to 'majordormo@cs.ubc.ca' with the contents 'subscribe cpsc535c' to join class list. (Note: email address does not correspond to correct class number!)
- Homework due in class on Monday 20th.
- Monday's class starts at 9.30am as usual.

NAIVE INFERENCE

- \bullet Represent joint prob. distribution P(C,S,R,W) as a 4D table of $2^4=32$ numbers.
- We observe the grass is wet and want to know how likely it was that the sprinkler caused this event.

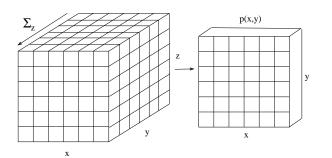
$$\begin{split} P(s=1|w=1) &= \frac{P(s=1,w=1)}{P(w=1)} \\ &= \frac{\sum_{c=0}^{1} \sum_{r=0}^{1} P(s=1,w=1,R=r,C=c)}{\sum_{c,r,s} P(S=s,w=1,R=r,C=c)} \\ &\xrightarrow[(\text{Creally})]{} \end{split}$$



• Query/hidden vars = $\{S\}$, visible vars = $\{W\}$, nuisance vars = $\{C, R\}$.

• It is easy to marginalize a joint probability distribution when it is represented as a table

 \bullet e.g., $P(X,Y) = \sum_z P(X,Y,Z)$



- Problems with representing joint as a big table
 - $-\operatorname{Representation:}$ big table of numbers is hard to understand.
 - -Inference: computing a marginal $P(X_i)$ takes $O(2^N)$ time.
 - -Learning: there are ${\cal O}(2^N)$ free parameters to estimate.
- Graphical models solve all 3 problems by providing a structured representation for joint probability distributions.
- Graphs encode conditional independence properties and represent families of probability distributions that satisfy these properties.
- Today we will study the relationship between graphs and independence properties.

INDEPENDENCE PROPERTIES OF DISTRIBUTIONS

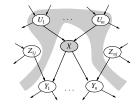
• Defn: let I(P) be the set of independence properties of the form $X \perp Y | Z$ that hold in distribution P.

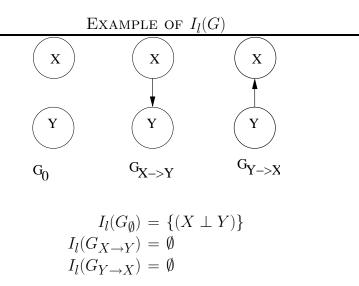
(LOCAL) INDEPENDENCE PROPERTIES OF DAGS

 \bullet Defn: let $I_l(G)$ be the set of local independence properties encoded by DAG G , namely:

 $\{X_i \perp \mathsf{NonDescendants}(X_i) | \mathsf{Parents}(X_i) \}$

- i.e., a node is conditionally independent of its non-descendants given its parents.
- Ancestors $(X_i) \subseteq NonDescendants(X_i)$





- Defn: A DAG G is an I-map (independence-map) of P if $I_l(G) \subseteq I(P)$.
- From previous example,

$$\begin{split} I_l(G_{\emptyset}) &= \{(X \perp Y)\}\\ I_l(G_{X \to Y}) &= \emptyset\\ I_l(G_{Y \to X}) &= \emptyset\\ I(P) &= \{(X \perp Y)\} \end{split}$$

• Hence all three graphs are I-maps of *P*.

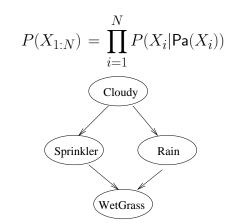
FROM I-MAP TO FACTORIZATION

- Defn: P factorizes according to G if P can be written as $P(X_1,\ldots,X_N) = \prod P(X_i|\mathsf{Pa}_G(X_i))$
- Thm 3.2.6: If G is an I-map of P, then P factorizes according to G.
 Proof:

$$\begin{split} P(X_{1:N}) &= P(X_1)P(X_2|X_1)P(X_3|X_1,X_2)\dots \text{ chain rule} \\ &= \prod_{i=1}^N P(X_i|X_{1:i-1}) \\ &= \prod_{i=1}^N P(X_i|\operatorname{Pa}(X_i),\operatorname{Ancestors}(X_i)\setminus\operatorname{Pa}(X_i)) \\ &= \prod_{i=1}^N P(X_i|\operatorname{Pa}(X_i)) \text{ since } G \text{ is I-map of } P \end{split}$$

BAYES NETS PROVIDE COMPACT REPRESENTATION OF JOINT PROBABILITY DISTRIBUTIONS

- Thm: If G is an I-map of P, then P factorizes according to G.
- Corollary: If G is an I-map of P, then we can represent P using G and a set of conditional probability distributions (CPDs), P(X_i|Pa(X_i)), one per node.
- Defn: A Bayesian network (aka belief network) representing distribution P is an I-map of P and a set of CPDs.
- For binary random variables, the Bayes net takes $O(N2^K)$ parameters ($K = \max$. num. parents), whereas full joint takes $O(2^N)$ parameters.
- Factored representation is easier to understand, easier to learn and supports more efficient inference (see later lectures).



- Thm 3.2.8: If P factorizes according to G, then G is an I-map of P.
- \bullet Proof: we must show $X \perp W | U$

$$\begin{split} P(X,W|U) &= \frac{P(X,W,U)}{P(U)} \\ &= \frac{\sum_Y P(X,W,U,Y)}{P(U)} \\ &= \frac{P(W)P(U|W)P(X|U)\sum_Y P(Y|X,W)}{P(U)} \\ &= \frac{P(W,U)}{P(U)}P(X|U)\sum_Y P(Y|X,W) \\ &= P(W|U)P(X|U) \end{split}$$

P(C, S, R, W) = P(C)P(S|C)P(R|C)P(W|S, R)

MINIMAL I-MAPS

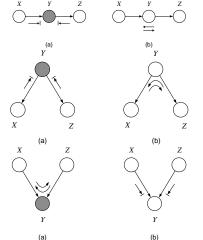
- \bullet Let G be a fully connected DAG. Then $I_l(G)= \emptyset \subseteq I(P)$ for any P.
- Hence the complete graph is an I-map for any distribution.
- Defn: A DAG G is a minimal I-map for P if it is an I-map for P, and if the removal of even a single edge from G renders it not an I-map.
- Construction: pick a node ordering, then let the parents of node X_i be the minimal subset of $U \subseteq \{X_1, \ldots, X_{i-1}\}$ s.t. $X_i \perp \{X_1, \ldots, X_i - 1\} \setminus U | U$.
- Defn (revised): A Bayesian network (aka belief network) representing distribution P is a *minimal* I-map of P and a set of CPDs.

GLOBAL MARKOV PROPERTIES OF DAGS

- By chaining together local independencies, we can infer more global independencies.
- Defn: X is d-separated (directed-separated) from Y given Z if along every undirected path between X and Y there is a node w s.t. either
 - -W has converging arrows ($\rightarrow w \leftarrow$) and neither W nor its descendants are in z; or
 - -W does not have converging arrows and $W \in Z$.
- \bullet Defn: $I(G) = \mathsf{all}$ independence properties that correspond to d-separation:

$$I(G) = \{(X \perp Y | Z) : d - sep_G(X; Y | Z)\}$$

A is d-separated from B given C if we cannot send a ball from any node in A to any node in B according to the rules below, where shaded nodes are in C.



Completeness of d-separation - v1

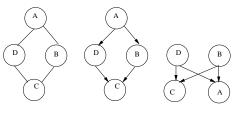
- Defn (Completeness) v1: For any distribution P that factorizes over G, if $(X \perp Y | Z) \in I(P)$, then $dsep_G(X; Y | Z)$.
- Contrapositive rule: $(A \Rightarrow B) \iff (\neg B \Rightarrow \neg A).$
- Defn (Completeness, contrapositive form) v1. If X and Y are not d-separated given Z, then X and Y are dependent in all distributions P that factorize over G.
- This definition of completeness is too strong since *P* may have conditional independencies that are not evident from the graph.
- \bullet eg. Let G be the graph $X \to Y,$ where P(Y|X) is $\begin{array}{c|c} A & B = 0 & B = 1 \\ \hline 0 & 0.4 & 0.6 \\ 1 & 0.4 & 0.6 \end{array}$
- G is I-map of P since $I(G) = \emptyset \subseteq I(P) = \{(X \perp Y)\}.$
- \bullet But the CPD encodes $X\perp Y$ which is not evident in the graph.

- Thm 3.3.3 (Soundness): If P factorizes according to G, then $I(G) \subseteq I(P)$.
- i.e., any independence claim made by the graph is satisfied by all distributions P that factorize according to G (no false claims of independence).
- Pf: see later (when we discuss undirected graphs).

Completeness of d-separation - v2

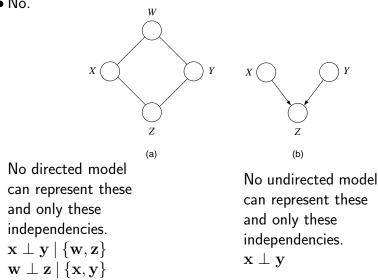
- Defn (Completeness) v2: If $(X \perp Y|Z)$ in all distributions P that factorize over G, then $dsep_G(X;Y|Z)$.
- Defn (Completeness, contrapositive form) v2: If X and Y are not d-separated given Z, then X and Y are dependent in *some* distribution P that factorizes over G.
- Thm 3.3.5: d-separation is complete.
- Proof: See Koller & Friedman p90.
- Hence d-separation captures as many of the independencies as possible (without reference to the particular CPDs) for all distributions that factorize over some DAG.

- Can we find a graph that captures all the independencies in an arbitrary distribution (and no more)?
- Defn: A DAG G is a perfect map (P-map) for a distribution P if I(P) = I(G).
- Thm: not every distribution has a perfect map.
- Pf by counterexample. Suppose we have a model where $A \perp C | \{B, D\}$, and $B \perp D | \{A, C\}$. This cannot be represented by any Bayes net.
- e.g., BN1 wrongly says $B \perp D | A$, BN2 wrongly says $B \perp D$.

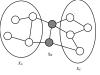


EXPRESSIVE POWER

- Can we always convert directed \leftrightarrow undirected?
- No.



- Graphs with one node per random variable and edges that connect pairs of nodes, but now the edges are undirected.
- Defn: Let H be an undirected graph. Then $sep_H(A; C|B)$ iff all paths between A and C go through some nodes in B (simple graph separation).



 \bullet Defn: the global Markov properties of a UG H are

 $I(H) = \{(X \perp Y | Z) : sep_H(X; Y | Z)\}$

- UGs can model symmetric (non-causal) interactions that directed models cannot.
- aka Markov Random Fields, Markov Networks.

CONDITIONAL PARAMETERIZATION?

- In directed models, we started with $p(\mathbf{X}) = \prod_i p(\mathbf{x}_i | \mathbf{x}_{\pi_i})$ and we derived the d-separation semantics from that.
- Undirected models: have the semantics, need parametrization.
- What about this "conditional parameterization"?

$$p(\mathbf{X}) = \prod_{i} p(\mathbf{x}_{i} | \mathbf{x}_{\text{neighbours}(i)})$$

• Good: product of local functions.

Good: each one has a simple conditional interpretation.

Bad: local functions cannot be arbitrary, but must agree properly in order to define a valid distribution.

• OK, what about this "marginal parameterization"?

$$p(\mathbf{X}) = \prod_{i} p(\mathbf{x}_{i}, \mathbf{x}_{\text{neighbours}(i)})$$

• Good: product of local functions.

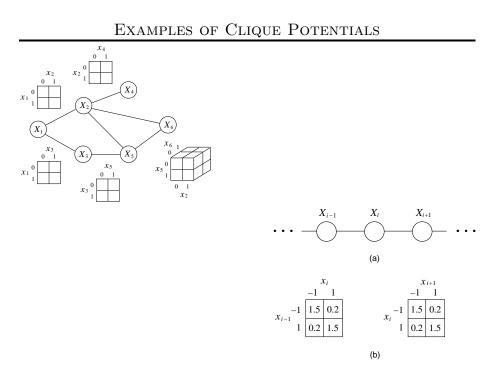
Good: each one has a simple marginal interpretation.

Bad: only very few pathalogical marginals on overalpping nodes can be multiplied to give a valid joint.

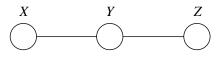
- Whatever factorization we pick, we know that only connected nodes can be arguments of a single local function.
- A *clique* is a fully connected subset of nodes.
- Thus, consider using a product of clique potentials:

$$\mathsf{P}(\mathbf{X}) = \frac{1}{Z} \prod_{\text{cliques } c} \psi_c(\mathbf{x}_c) \qquad \qquad Z = \sum_{\mathbf{X}} \prod_{\text{cliques } c} \psi_c(\mathbf{x}_c)$$

- Each clique potential $\psi_c(\mathbf{x}_c) > 0$ is an arbitrary positive function of its arguments.
- The normalization term Z is called the partition function (a function of the parameters ψ) and ensures $\sum_{\mathbf{x}} \mathsf{P}(\mathbf{x}) = 1$.
- Without loss of generality we can restrict ourselves to *maximal cliques.* (Why?)
- \bullet A distribution P that is representable by a UG H in this way is called a Gibbs distribution over H.



INTERPRETATION OF CLIQUE POTENTIALS



 \bullet The model implies $\mathbf{x} \perp \mathbf{z} \mid \mathbf{y}$

$$p(\mathbf{x}, \mathbf{y}, \mathbf{z}) = p(\mathbf{y})p(\mathbf{x}|\mathbf{y})p(\mathbf{z}|\mathbf{y})$$

• We can write this as:

$$\begin{split} p(\mathbf{x}, \mathbf{y}, \mathbf{z}) &= p(\mathbf{x}, \mathbf{y}) p(\mathbf{z} | \mathbf{y}) = \psi_{\mathbf{x} \mathbf{y}}(\mathbf{x}, \mathbf{y}) \psi_{\mathbf{y} \mathbf{z}}(\mathbf{y}, \mathbf{z}) \\ p(\mathbf{x}, \mathbf{y}, \mathbf{z}) &= p(\mathbf{x} | \mathbf{y}) p(\mathbf{z}, \mathbf{y}) = \psi_{\mathbf{x} \mathbf{y}}(\mathbf{x}, \mathbf{y}) \psi_{\mathbf{y} \mathbf{z}}(\mathbf{y}, \mathbf{z}) \end{split}$$

cannot have all potentials be marginals cannot have all potentials be conditionals

• The positive clique potentials can only be thought of as general "compatibility", "goodness" or "happiness" functions over their variables, but not as probability distributions.