Graphical Models For Complex Health Data (P8124)

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Directed Graphical Models

Directed Acyclic Graphs

A directed acyclic graph (DAG)is a graph $\mathcal{G} = (V, E)$ is such that E contains only directed edges (\rightarrow) and has no sequence of directed edges from X_i to X_i $(\forall i)$.

Definitions:

$$\begin{array}{l} \mathsf{Pa}(X_i,\mathcal{G}) \equiv \{X_j: X_j \to X_i \text{ in } \mathcal{G}\} \\ \mathsf{Ch}(X_i,\mathcal{G}) \equiv \{X_j: X_j \leftarrow X_i \text{ in } \mathcal{G}\} \\ \mathsf{An}(X_i,\mathcal{G}) \equiv \{X_j: X_j \to \ldots \to X_i \text{ in } \mathcal{G}\} \\ \mathsf{De}(X_i,\mathcal{G}) \equiv \{X_j: X_j \leftarrow \ldots \leftarrow X_i \text{ in } \mathcal{G}\} \\ \end{array}$$
 (descendants of X_i)

Factorization Property

A distribution p(x) satisfies the factorization property wrt DAG $\mathcal G$ if

$$p(x) = \prod_{i=1}^{p} p(x_i|\operatorname{Pa}(X_i,\mathcal{G}))$$

where $Pa(X_i, \mathcal{G})$ are the parents of X_i in \mathcal{G} .

Definition: Bayesian network model

A pair $(\mathcal{G}, \mathcal{P})$ where \mathcal{G} is a DAG and \mathcal{P} is a set of distributions that factorize wrt \mathcal{G} .

By the chain rule of probability, any distribution

$$p(x) = p(x_1, ..., x_p)$$

$$= p(x_p | x_{p-1}, ..., x_1) p(x_{p-1}, ..., x_1)$$

$$= p(x_p | x_{p-1}, ..., x_1) p(x_{p-1} | x_{p-2}, ..., x_1) p(x_{p-2}, ..., x_1)$$

$$= \prod_{i=1}^{p} p(x_i | x_{i-1}, ..., x_1)$$

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$$= \prod_{i=1}^{p} p(x_i | x_{i-1}, ..., x_1)$$

Compare this to the DAG factorization:

$$p(x) = \prod_{i=1}^{p} p(x_i | Pa(X_i, \mathcal{G}))$$

What does the factorization property "get you"?



Figure: A complete DAG

$$p(x) = \prod_{i=1}^{p} p(x_i | Pa(X_i, \mathcal{G}))$$

$$= p(x_1)p(x_2|x_1)p(x_3|x_2, x_1)p(x_4|x_3, x_2, x_1)$$



Figure: A complete DAG

$$p(x) = \prod_{i=1}^{p} p(x_i| \operatorname{Pa}(X_i, \mathcal{G}))$$

= $p(x_1)p(x_2|x_1)p(x_3|x_2, x_1)p(x_4|x_3, x_2, x_1)$

For a complete DAG the factorization property doesn't say anything! Just equivalent to chain rule.

⇒ "A complete DAG imposes no restrictions on the data distribution."

$$X_1 \longrightarrow X_2 \longrightarrow X_3 \longrightarrow X_4$$

Figure: A chain

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A DAG with missing edges encodes conditional independencies.

Which conditional independencies?

Compare:

$$p(x) = p(x_1)p(x_2|x_1)p(x_3|x_2,x_1)p(x_4|x_3,x_2,x_1)$$

$$= p(x_1)p(x_2|x_1)p(x_3|x_2)p(x_4|x_3)$$

$$p(x_4|x_3,x_2,x_1) = p(x_4|x_3) \text{ or } X_4 \perp X_2, X_1|X_3$$
and
$$p(x_3|x_2,x_1) = p(x_3|x_2) \text{ or } X_3 \perp X_1|X_2$$

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Compare:

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and
$$p(x_3|x_2,x_1) = p(x_3|x_2)$$
 or $X_3 \perp X_1|X_2$

something like... $X_i \perp$ "ancestors" | parents ??

 $p(x_4|x_3,x_2,x_1) = p(x_4|x_3)$ or $X_4 \perp X_2, X_1|X_3$

Local Markov property

Define:

$$Nd(X_i, \mathcal{G}) \equiv \{X_j : X_j \notin De(X_i, \mathcal{G})\}$$
 (non-descendants of X_i)

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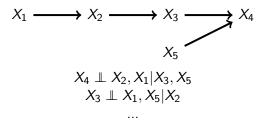
$$Nd(X_i, \mathcal{G}) \equiv \{X_j : X_j \notin De(X_i, \mathcal{G})\}$$
 (non-descendants of X_i)

A distribution p(x) satisfies the *local Markov property* wrt DAG \mathcal{G} if

$$X_i \perp \operatorname{Nd}^*(X_i, \mathcal{G}) | \operatorname{Pa}(X_i, \mathcal{G})$$

where
$$\operatorname{\sf Nd}^*(X_i,\mathcal{G}) \equiv \operatorname{\sf Nd}(X_i,\mathcal{G}) \setminus \operatorname{\sf Pa}(X_i,\mathcal{G})$$

Local Markov property



Global Markov property

Let A, B, C be disjoint subsets of X. A distribution p(x) satisfies the global Markov property wrt DAG \mathcal{G} if

$$A \perp_d B | C \implies A \perp \!\!\! \perp B | C.$$

Global Markov property

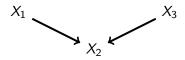
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"A graphical criterion \implies conditional independence in the distribution"

Colliders

Given a path π in a graph \mathcal{G} , a non-endpoint vertex X_j on π is called a collider if the two edges incident to X_j are both into X_j , i.e., have arrowheads at X_j .¹



¹A v-structure is a triple $\langle X_i, X_j, X_k \rangle$ such that X_j is a collider and X_i and X_k are not adjacent. A collider which is part of a v-structure (i.e., a collider with non-adjacent parents) is also called an unshielded collider.

d-separation

A path π in $\mathcal G$ between distinct vertices X_i and X_k is called a *d-connecting path* conditional on vertex set $\mathcal C$ ($\mathcal C \subseteq X \setminus \{X_i, X_k\}$) if:

- a) every collider on π is in C or is in $An(C,\mathcal{G})$ and
- b) every non-collider on π is not in C.

 X_i and X_k are *d-separated* conditional on C if there is no d-connecting path conditional on C between X_i and X_k .

 $A \perp_d B \mid C$ if $X_i \perp_d X_k \mid C$ for all $X_i \in A$ and $X_k \in B$.

d-separation (equivalent def)

A vertex X_j is **active** on a path relative to C just in case either

- i) X_j is a collider, and X_j or any of its descendents is in C, or
- ii) X_j is a noncollider and is not in C.

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A path π is **active** relative to C just in case every vertex on π is active relative to C.

 X_i and X_k are d-separated given C just in case there is no path between X_i and X_k that is **active** relative to C.

⇒ "Active can be thought of as carrying association"

d-separation examples

[on the board]

Equivalence of Markov properties in BNs

Theorem. Let $\mathcal G$ be a DAG. For any probability distribution which has a density wrt product measure, the factorization, local Markov, and Global Markov properities (wrt $\mathcal G$) are equivalent:

Factorization ← Local Markov ← Global Markov

Prove: global \implies local

Need to show that $X_i \perp_d \operatorname{Nd}(X_i) \setminus \operatorname{Pa}(X_i) | \operatorname{Pa}(X_i)$. Then global property $\implies X_i \perp \operatorname{Nd}(X_i) \setminus \operatorname{Pa}(X_i) | \operatorname{Pa}(X_i) | \operatorname{local}$.

Any path from X_i to $Nd(X_i)$ must go through either $Ch(X_i)$ or $Pa(X_i)$. If the path goes through $Ch(X_i)$ then there must be a collider on that path, but it is not in the conditioning set $Pa(X_i)$ (nor an ancestor of $Pa(X_i)$ by acyclicity), so that path must be not active. If instead the path goes through $Pa(X_i)$, the parents would be non-colliders on the path, and since they are in the conditionining set, they are also not active. So all paths are not active, therefore the paths cannot be d-connecting.

The reverse direction (global \iff local) is trickier: proof by induction on graph size.

I-map

Let $\mathcal P$ be a set of distributions (model) over X. We define $\mathcal I(\mathcal P)$ to be the set of independence assertions of the form $(A \perp\!\!\!\perp B \mid C)$ that hold in $\mathcal P$. (A,B,C) are disjoint subsets of X.)

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Note: \mathcal{P} may have independencies that are not reflected in $\mathcal{I}(\mathcal{G})$!

Example: deterministic relationships among variables

Consider:

$$X_1 \longrightarrow X_2 \longrightarrow X_3$$

where $X_3 = 2 \times X_2$.

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$$X_1 \longrightarrow X_2 \longrightarrow X_3$$

where $X_3 = 2 \times X_2$.

 $X_3 \perp X_1 | X_2$ (by Markov property) but also $X_2 \perp X_1 | X_3$ (by determinism).

Example: exact "cancellation" or "balancing"

Consider:

$$X_1 \longrightarrow X_2 \longrightarrow X_3$$

where

$$X_1 = \epsilon_1$$

$$X_2 = \alpha X_1 + \epsilon_2$$

$$X_3 = \beta X_2 - \alpha \beta X_1 + \epsilon_3$$

$$\epsilon_1, \epsilon_2, \epsilon_3 \sim N(0, 1), \quad \alpha, \beta > 0$$

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No independencies follow from Markov property, but $X_3 \perp X_1$ (by exact cancellation).

It is common to rule out "extra" or "non-graphical" conditional independencies, *by assumption*. This restricts the allowed set of distributions, and may not always be appropriate.

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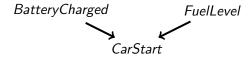
$$A \perp \!\!\! \perp B \mid C \implies A \perp_d B \mid C.$$

In conjunction with the global Markov property this means we're assuming $A \perp\!\!\!\perp B \mid C \iff A \perp_d B \mid C$.

... is not always an appropriate assumption to make, but greatly simplifies things ("all conditional independence information is in the graph") and plays an important role in structure learning (as we will discuss later).

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Also note that if a distribution satisfies the Markov and faithfulness assumptions wrt \mathcal{G} we sometimes say that \mathcal{G} is a *perfect map* for that distribution.

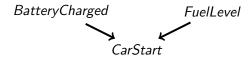


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By d-separation: $BatteryCharged \perp FuelLevel$.

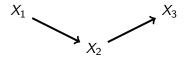


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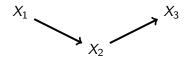
By d-separation: BatteryCharged \bot FuelLevel.

By faithfulness assumption: $BatteryCharged \not\perp FuelLevel \mid CarStart$.

Conditioning on *CarStart* makes *FuelLevel* informative about *BatteryCharged*.

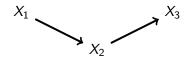


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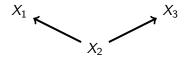


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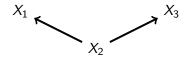
By d-separation: $X_1 \not\perp X_3$.

By d-separation: $X_1 \perp X_3 | X_2$.

Conditioning on X_2 makes X_1 non-informative about X_3 .

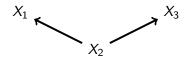


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Markov equivalence

Markov equivalence

Definition. Two DAGs \mathcal{G}_1 and \mathcal{G}_2 are called Markov equivalent if $\mathcal{I}(\mathcal{G}_1)=\mathcal{I}(\mathcal{G}_2)$.

Theorem. Two DAGs \mathcal{G}_1 and \mathcal{G}_2 are Markov equivalent iff they share the same adjacencies and unshielded colliders.

Markov equivalence

$$X_i o X_j o X_k$$
 $X_i \leftarrow X_j \leftarrow X_k$
 $X_i \leftarrow X_j o X_k$
 $X_i \leftarrow X_j o X_k$
 $X_i \to X_j \leftarrow X_k$
b)

Figure: a) Three Markov equivalent DAG models. b) The CPDAG representation of (a). c) A DAG that is not Markov equivalent to the graphs in (a).

Implications of Markov equivalence

Markov equivalence is *almost* like "observational equivalence" – the data cannot distinguish between Markov equivalent graphs (unless we use more than conditional independence information).

So, using only conditional independence information, we cannot learn the "correct" structure within an equivalence class. Equivalence classes are the natural "units" for structure learning. (More on this later!)

Markov blankets

 $\mathsf{Mb}(X_i,\mathcal{G})$ is called the Markov blanket of X_i . It is a set of vertices that "screens off" all other vertices in \mathcal{G} , i.e., $X_i \perp X \setminus \{\mathsf{Mb}(X_i,\mathcal{G}),X_i\}|\mathsf{Mb}(X_i,\mathcal{G}).$

In a DAG, $\mathsf{Mb}(X_i,\mathcal{G}) \equiv \mathsf{Pa}(X_i) \cup \mathsf{Ch}(X_i) \cup \mathsf{Pa}(\mathsf{Ch}(X_i))$.

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In a DAG,
$$\mathsf{Mb}(X_i,\mathcal{G}) \equiv \mathsf{Pa}(X_i) \cup \mathsf{Ch}(X_i) \cup \mathsf{Pa}(\mathsf{Ch}(X_i))$$
.

If you want only to predict X_i , $Mb(X_i, \mathcal{G})$ is sufficient. [example on board]



Yes, in a boring sense: just use the complete DAG.



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The complete DAG is an I-map of any model, since $\mathcal{I}(\mathcal{G})$ is empty.

No!

No! Even leaving issues such as determinism and exact cancellation aside, there are sets of conditional independence facts that correspond to no DAG over the *observed* variables.

Consider:

$$X_1 \perp \!\!\! \perp X_3 | (X_2, X_4)$$
 and $X_2 \perp \!\!\! \perp X_4 | (X_1, X_3)$ (and all vars are pairwise dependent)

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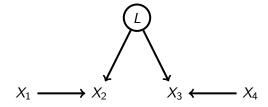
 \Rightarrow there is no faithful BN to describe these independencies (may use a MRF)

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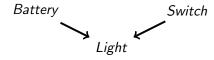
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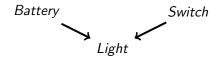
⇒ Might have to posit *unobserved* variables to explain observed independencies

Context-specific independence



Consider a battery-powered light bulb, with a switch. Charge in the battery will cause the bulb to light up provided the switch is on, but not otherwise. The dependence of *Light* and *Battery* arises entirely through the condition *Switch* = 'on'. When *Switch* = 'off', *Light* and *Battery* are independent.

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These independencies are correctly represented by the simple DAG, but it is not fully informative. The independence is "context-specific," i.e., only appears in the context of a certain variable setting. Representing this system with a single DAG isn't wrong, but there are alternative, richer representations that may be more useful. Likewise for "interactions."

Context-specific independence

$$Switch = "on"$$
 $Switch = "off"$ $Battery \longrightarrow Light$ $Battery$ $Light$

We could represent this situation using a set of Bayesian Networks, a.k.a. a multinet. One BN for every "context" setting.

Special structure: Hidden Markov Models (HMMs)

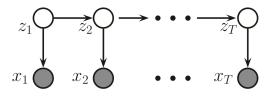


Figure 10.4 A first-order HMM.

(from Murphy (2012))

Special structure: Dynamic Bayesian Networks (DBNs)

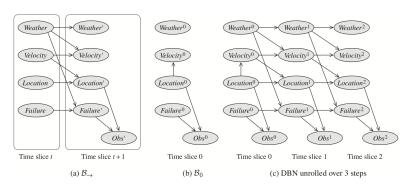


Figure 6.1 A highly simplified DBN for monitoring a vehicle: (a) the 2-TBN; (b) the time 0 network; (c) resulting unrolled DBN over three time slices.