Graphical models for causal inference

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Example is compelling for causal reasons



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Causal interpretations are tied to the notion of *conditioning by intervention*

$$P(X = x \mid Y \leftarrow y) = P\{X = x \mid do(Y = y)\} = p(x \mid \mid y), \quad (1)$$

which in general is quite different from conventional conditioning or *conditioning by observation* which is

$$P(X = x | Y = y) = P\{X = x | is(Y = y)\} = p(x | y) = p(x, y)/p(y).$$

A causal interpretation of a Bayesian network involves giving (1) a simple form.

[Also distinguish p(x | y) from $P\{X = x | see(Y = y)\}$. Observation/sampling bias.]

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We say that a BN is *causal w.r.t. atomic interventions at* $B \subseteq V$ if it holds for any $A \subseteq B$ that

$$p(x \mid\mid x_A^*) = \prod_{v \in V \setminus A} p(x_v \mid x_{\mathsf{pa}(v)}) \bigg|_{x_A = x_A^*}$$

For $A = \emptyset$ we obtain standard factorisation.

Note that conditional distributions $p(x_v | x_{pa(v)})$ are stable under interventions which do not involve x_v . Such assumption must be justified in any given context.

Contrast the formula for intervention conditioning with that for observation conditioning:

$$p(x || x_A^*) = \prod_{v \in V \setminus A} p(x_v | x_{\mathsf{pa}(v)}) \bigg|_{x_A = x_A^*}$$
$$= \frac{\prod_{v \in V} p(x_v | x_{\mathsf{pa}(v)})}{\prod_{v \in A} p(x_v | x_{\mathsf{pa}(v)})} \bigg|_{x_A = x_A^*}$$

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whereas

$$p(x \mid x_A^*) = \frac{\prod_{v \in V} p(x_v \mid x_{pa(v)})}{p(x_A)}\Big|_{x_A = x_A^*}$$

An example



$$\begin{array}{rcl} p(x \,||\, x_5^*) &=& p(x_1) p(x_2 \,|\, x_1) p(x_3 \,|\, x_1) p(x_4 \,|\, x_2) \\ & \times & p(x_6 \,|\, x_3, x_5^*) p(x_7 \,|\, x_4, x_5^*, x_6) \end{array}$$

whereas

$$\begin{array}{rcl} p(x \mid x_5^*) & \propto & p(x_1)p(x_2 \mid x_1)p(x_3 \mid x_1)p(x_4 \mid x_2) \\ & \times & p(x_5^* \mid x_2, x_3)p(x_6 \mid x_3, x_5^*)p(x_7 \mid x_4, x_5^*, x_6) \end{array}$$

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DAG \mathcal{D} can also represent structural equation system:

$$X_{\nu} \leftarrow g_{\nu}(x_{\mathsf{pa}(\nu)}, U_{\nu}), \nu \in V, \tag{2}$$

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where g_v are fixed functions and U_v are independent random disturbances.

Intervention in structural equation system can be made by *replacement*, i.e. so that $X_v \leftarrow x_v^*$ is replacing the corresponding line in 'program' (2).

Corresponds to g_v and U_v being unaffected by the intervention if intervention is not made on node v. Hence the equation is *structural*.



For the network shown, we get

$$\begin{array}{rcl} X_1 & \leftarrow & \alpha_1 + U_1 \\ X_2 & \leftarrow & \alpha_2 + \beta_{21} x_1 + U_2 \\ X_3 & \leftarrow & \alpha_3 + \beta_{31} x_1 + U_3 \\ X_4 & \leftarrow & \alpha_4 + \beta_{42} x_2 + U_4 \\ X_5 & \leftarrow & \alpha_5 + \beta_{52} x_2 + \beta_{53} x_3 + U_5 \\ X_6 & \leftarrow & \alpha_6 + \beta_{63} x_3 + \beta_{65} x_5 + U_6 \\ X_7 & \leftarrow & \alpha_7 + \beta_{74} x_4 + \beta_{75} x_5 + \beta_{76} x_6 + U_7. \end{array}$$

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After intervention by replacement, the system changes to

$$\begin{array}{rcl} X_1 & \leftarrow & \alpha_1 + U_1 \\ X_2 & \leftarrow & \alpha_2 + \beta_{21} x_1 + U_2 \\ X_3 & \leftarrow & \alpha_3 + \beta_{31} x_1 + U_3 \\ X_4 & \leftarrow & x_4^* \\ X_5 & \leftarrow & \alpha_5 + \beta_{52} x_2 + \beta_{53} x_3 + U_5 \\ X_6 & \leftarrow & \alpha_6 + \beta_{63} x_3 + \beta_{65} x_5 + U_6 \\ X_7 & \leftarrow & \alpha_7 + \beta_{74} x_4^* + \beta_{75} x_5 + \beta_{76} x_6 + U_7. \end{array}$$

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Intervention by replacement in structural equation system implies \mathcal{D} causal for distribution of $X_v, v \in V$.

Occasionally used for *justification* of CBN.

Ambiguity in choice of g_v and U_v makes this problematic.

May take *stability of conditional distributions* as a primitive rather than structural equations.

Structural equations more expressive when choice of g_v and U_v can be externally justified.

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a - treatment with AZT; *I* - intermediate response (possible lung disease); *b* - treatment with antibiotics; r - survival after a fixed period.

Predict survival if $X_a \leftarrow 1$ and $X_b \leftarrow 1$, assuming stable conditional distributions.

- **B** - **b** - - -

G-computation



$$p(1_r || 1_a, 1_b) = \sum_{x_l} p(1_r, x_l || 1_a, 1_b)$$

=
$$\sum_{x_l} p(1_r | x_l, 1_a, 1_b) p(x_l | 1_a).$$

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Augment each node $v \in A$ where intervention is contemplated with additional parent variable F_v .

 F_ν has state space $\mathcal{X}_\nu\cup\{\phi\}$ and conditional distributions in the intervention diagram are

$$p'(x_{v} | x_{\mathsf{pa}(v)}, f_{v}) = \begin{cases} p(x_{v} | x_{\mathsf{pa}(v)}) & \text{if } f_{v} = \phi \\ \delta_{x_{v}, x_{v}^{*}} & \text{if } f_{v} = x_{v}^{*}, \end{cases}$$

where δ_{xy} is Kronecker's symbol

$$\delta_{xy} = \begin{cases} 1 & \text{if } x = y \\ 0 & \text{otherwise.} \end{cases}$$

 F_v is *forcing* the value of X_v when $F_v \neq \phi$.

It now holds in the extended intervention diagram that

$$p(x) = p'(x \mid F_v = \phi, v \in A),$$

but also

$$\begin{aligned} p(x \mid\mid x_B^*) &= P(X = x \mid X_B \leftarrow x_B^*) \\ &= P'(x \mid F_v = x_v^*, v \in B, F_v = \phi, v \in B \setminus A), \end{aligned}$$

In particular it holds that if $pa(v) = \emptyset$, then $p(x | x_v^*) = p(x_v || x_v^*)$.

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Treatment variable t, response r, set of observed covariates C, unobserved variables U.

When and how can $p(X_r || x_t)$ be calculated from $p(x_t, x_r, x_c)$, the latter in principle being observable from data?

In this case we could say that C is a *identifier* for assessing the effect of T on R.

Answer can be found by analysing intervention diagram.

Simplest cases known as *back-door* and *front-door* criteria and formulae.

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 $\begin{array}{l} \mathcal{D}' \text{ denotes } \mathcal{D} \text{ augmented with } F_t. \\ \text{Assume } C \supseteq C_0, \text{ where } C_0 \text{ satisfies} \\ (\text{BD1}) \text{ Covariates in } C_0 \text{ are unaffected by an intervention:} \\ C_0 \perp_{\mathcal{D}'} F_t; \\ (\text{BD2}) \text{ Intervention only affects response through the} \end{array}$

treatment it chooses: $R \perp_{\mathcal{D}'} F_t \mid C_0 \cup \{t\}$.

Then C identifies the effect of the treatment t on R as

$$p(x_r || x_t^*) = \sum_{x_{C_0}} p(x_r | x_{C_0}, x_t^*) p(x_{C_0}).$$

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The unobserved *confounder* X_u is affecting both treatment and response.

BD2 is violated; graph to the right reveals that F_t is not

d-separated from r by t, so treatment effect is not identifiable.



When X_t is randomised, possibly depending on observed covariate c, confounding is resolved.

Now $F_t \perp_{\mathcal{D}'} r \mid \{c, t\}$ and *c* is an identifier.

Sufficient covariate



Alternatively, an observed covariate c can 'screen away' the confounding effect on the treatment. Also here, $F_t \perp_{\mathcal{D}'} r \mid \{c, t\}$ and c is an identifier.

Instrumental variable



i is an instrumental variable as it affects t and it is uncorrelated with the confounders.

Graph to the right shows $r \perp_{D'} F_i | \{i, t\}$ so the effect of the instrument can be identified.

However, r is not d-separated from F_t by t so the *effect of the treatment itself is not*.

Note that in the linear case, the effect of t on r can be found as the ratio of effects of i on r and the effect of i on t, both of which are identified.

In the linear case, many more effects can be identified. But linearity and additivity of errors are very strong assumptions.

Bounds are available in the general case